

## Supplemental Information – MIAME format for cDNA Microarrays:

Paper: Linkage of Cardiac Gene Expression Profiles and ETS2 with Lifespan Variability in Rats.

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All methods for cDNA probe preparation, microarray hybridization and analysis were performed by Incyte Genomics as described by Yue et al. <sup>1</sup>.

1. **Experimental design:** *the set of hybridization experiments as a whole.* These experiments were designed to quantify mRNA transcript abundance in rat heart as a function of relative age and survival. Three strains of rats (Fisher 344, Wistar and Fisher x Brown Norway) were used for comparative and predictive purposes. All the animals were normal, and hybridizations were based on comparisons with 6 month old animals.

Experimentally, we used microarrays coupled with a bioinformatics tool, Prediction Analysis of Microarrays (PAM) <sup>2,3</sup>, to identify a putative set of gene transcripts that best characterize survival in Fisher 344 and Wistar inbred rat strains. This predictive gene set was employed to identify conserved *cis*-elements, which were utilized to define a set of longevity-dependent, strain independent transcription factor transcripts that could be linked to the regulation of these genes. We validated the putative survival gene set by demonstrating its ability to predict relative survival from microarray analyses (n=4) of a third longer-lived rat strain (Fisher x Brown Norway).

Experimental Relationships between the array and sample entities are as follows:

RNA samples from Fischer 344: 6 months (Cy3 labeled) vs 12, 18, 24 or 30 month (Cy5 labeled) – Hybridized against Gems 1 and 2 microarrays

Gem 0222ATLD: Fischer Left Ventricle 6 vs 12 month	Gem 1
Gem 022GARR8: Fischer Left Ventricle 6 vs 12 month	Gem 1
Gem 022QC3WE: Fischer Left Ventricle 6 vs 12 month	Gem 2
Gem 022XC3WF: Fischer Left Ventricle 6 vs 12 month	Gem 2
Gem 0226AASG: Fischer Left Ventricle 6 vs 18 month	Gem 1
Gem 022NBESF: Fischer Left Ventricle 6 vs 18 month	Gem 1
Gem 0224C3WG: Fischer Left Ventricle 6 vs 18 month	Gem 2
Gem 022BC3WH: Fischer Left Ventricle 6 vs 18 month	Gem 2
Gem 0220A026: Fischer Left Ventricle 6 vs 24 month	Gem 1
Gem 022DAASH: Fischer Left Ventricle 6 vs 24 month	Gem 1
Gem 022IC3WI: Fischer Left Ventricle 6 vs 24 month	Gem 2
Gem 022PC3WJ: Fischer Left Ventricle 6 vs 24 month	Gem 2
Gem 022SA2MZ: Fischer Left Ventricle 6 vs. 30 month	Gem 1
Gem 022WBBJL: Fischer Left Ventricle 6 vs. 30 month	Gem 16

Gem 022WC3WK: Fischer Left Ventricle 6 vs. 30 month Gem 2  
 Gem 0223C3WL: Fischer Left Ventricle 6 vs. 30 month Gem 2

Wistar Rats: 6 months (Cy3 labeled) vs 12 , 18 , 24 or 27 month (Cy5 labeled) – Hybridized against Gems 1 and 2 microarrays

Gem 0229ATLE: Wistar Left Ventricle 6 vs 12 month Gem 1  
 Gem 022GATLF: Wistar Left Ventricle 6 vs 18 month Gem 1  
 Gem 022NATLG: Wistar Left Ventricle 6 vs 24 month Gem 1  
 Gem 022ABBJN: Wistar Left Ventricle 6 vs 27 month Gem 1  
 Gem 022OC3WO: Wistar Left Ventricle 6 vs 27 month Gem 2  
 Gem 022VC3WP: Wistar Left Ventricle 6 vs 27 month Gem 2

Fisher x Brown Norway: 6 months (Cy3 labeled) vs. 36 month (Cy5 labeled) – Hybridized against Gems 1 and 2 microarrays

Gem 0224B8T2: Fischer x Norway Brown Left Ventricle 6 vs. 36 month Gem 1  
 Gem 0223BBJM Fischer x Norway Brown Left Ventricle 6 vs. 36 month Gem 1  
 Gem 022AC3WM Fischer x Norway Brown Left Ventricle 6 vs. 36 month Gem 2  
 Gem 022HC3WN Fischer x Norway Brown Left Ventricle 6 vs. 36 month Gem 2

2. **Array design:** *each array used and each element (spot, feature) on the array.*

(i) *Description of Array as a whole:* Two cDNA-based microarrays, containing PCR amplified cDNA fragments arrayed on an ino-modified glass slides, were employed in this study: Gem Type Mouse 1 ( 9024 cDNAs) and Gem Mouse Type 2 (10,176 cDNAs). Both of these arrays contained mouse cDNAs amplified by PCR and Internal control cDNAs (yeast and other), and Incyte Genomics performed all microarray manipulations, including the production and hybridization of labeled probes to these arrays.

(ii) *Description of each type of element or spot used:* Complete details for cDNA clone amplification by PCR (i.e., Synthesis of PCR products), isolation, quality control (i.e., Qualification and quantification of PCR products) and arraying on glass slides (i.e., Arraying and post-processing, Array qualification: SYTO 61 dye) can be found in <sup>1</sup>.

(iii) *A description of the specific properties of each element (DNA sequence identifier and quality-control indicators):*

**Gem Mouse1:** 9024 elements

Location	PlateRow	PlateCol	PlateID	GeneName	CloneID	CloneSource	AccessionNum
	1	A	1	2159971 Internal_Control_E	Internal_Control_E	Controls	
	2	A	3	2159971 Internal_Control_F	Internal_Control_F	Controls	
	3	A	5	2159971 Internal_Control_G	Internal_Control_G	Controls	

4 A	7	2159971	Internal_Control_H	Internal_Control_H	Controls		
5 A	9	2159971	Internal_Control_I	Internal_Control_I	Controls		
6 A	11	2159971	Internal_Control_J	Internal_Control_J	Controls		
7 C	1	2159971	Internal_Control_K	Internal_Control_K	Controls		
8 C	3	2159971	Internal_Control_L	Internal_Control_L	Controls		
9 C	5	2159971	Internal_Control_M	Internal_Control_M	Controls		
10 C	7	2159971	Internal_Control_N	Internal_Control_N	Controls		
11 C	9	2159971	Internal_Control_O	Internal_Control_O	Controls		
12 C	11	2159971	Internal_Control_P	Internal_Control_P	Controls		
13 E	1	2159971	Internal_Control_Q	Internal_Control_Q	Controls		
14 E	3	2159971	Internal_Control_R	Internal_Control_R	Controls		
15 E	5	2159971	Internal_Control_S	Internal_Control_S	Controls		
16 E	7	2159971	Internal_Control_T	Internal_Control_T	Controls		
17 E	9	2159971	Internal_Control_U	Internal_Control_U	Controls		
18 E	11	2159971	Internal_Control_V	Internal_Control_V	Controls		
19 G	1	2159971	Internal_Control_W	Internal_Control_W	Controls		
20 G	3	2159971	Internal_Control_X	Internal_Control_X	Controls		
21 G	5	2159971	Internal_Control_Y	Internal_Control_Y	Controls		
22 G	7	2159971	Internal_Control_Z	Internal_Control_Z	Controls		
23 G	9	2159971	Internal_Control_AA	Internal_Control_AA	Controls		
24 G	11	2159971	Internal_Control_AB	Internal_Control_AB	Controls		
25 A	1	021MA1E8	IMAGE EST {IMAGE:780337}	780337	IMAGE Consortium	AA415819	
26 A	3	021MA1E8	ESTs {IMAGE:804426}	804426	IMAGE Consortium	AA474746	
27 A	5	021MA1E8	ESTs {IMAGE:805250}	805250	IMAGE Consortium	AA473306	
28 A	7	021MA1E8	Mus musculus metal response element DNA-binding protein M96 mRNA, complete cds {IMAGE:945218}	945218	IMAGE Consortium	AA545607	
29 A	9	021MA1E8	IMAGE EST {IMAGE:834148}	834148	IMAGE Consortium	AA413090	
30 A	11	021MA1E8	Mus musculus putative G-protein (GP-1) mRNA, complete cds {IMAGE:760753}	760753	IMAGE Consortium	AA386616	
31 C	1	021MA1E8	IMAGE EST {IMAGE:805825}	805825	IMAGE Consortium	AA437482	
32 C	3	021MA1E8	ESTs, Highly similar to vasopressin activated calcium mobilizing receptor-like protein [H.sapiens] {IMAGE:833817}	833817	IMAGE Consortium	AA412921	
33 C	5	021MA1E8	ESTs, Highly similar to THYMIDYLATE KINASE [Homo sapiens] {IMAGE:760921}	760921	IMAGE Consortium	AA386718	
34 C	7	021MA1E8	ESTs {IMAGE:760847}	760847	IMAGE Consortium	AA386794	
35 C	9	021MA1E8	IMAGE EST {IMAGE:894406}	894406	IMAGE Consortium	AA517679	
36 C	11	021MA1E8	ESTs, Highly similar to NUCLEOPROTEIN TPR [H.sapiens] {IMAGE:932755}	932755	IMAGE Consortium	AA538426	
37 E	1	021MA1E8	Mus musculus mRNA for type II phosphatidylinositolphosphate kinase-alpha, complete cds {IMAGE:932803}	932803	IMAGE Consortium	AA538499	
38 E	3	021MA1E8	IMAGE EST {IMAGE:892346}	892346	IMAGE Consortium	AA501253	
39 E	5	021MA1E8	IMAGE EST {IMAGE:804944}	804944	IMAGE Consortium	AA397342	

40 E	7 021MA1E8	IMAGE EST {IMAGE:805843}	805843 IMAGE Consortium	AA399856
41 E	9 021MA1E8	IMAGE EST {IMAGE:834844}	834844 IMAGE Consortium	AA467182
42 E	11 021MA1E8	IMAGE EST {IMAGE:760968}	760968 IMAGE Consortium	AA387446
43 G	1 021MA1E8	IMAGE EST {IMAGE:894180}	894180 IMAGE Consortium	AA517039
44 G	3 021MA1E8	Fibroblast growth factor receptor 1 {IMAGE:933285}	933285 IMAGE Consortium	AA542013
45 G	5 021MA1E8	X-ray repair complementing defective repair in Chinese hamster {IMAGE:805093}	805093 IMAGE Consortium	AA397212
46 G	7 021MA1E8	ESTs, Weakly similar to C13F10.2 [C.elegans] {IMAGE:934431}	934431 IMAGE Consortium	AA545049
47 G	9 021MA1E8	Mus musculus antigen containing epitope to monoclonal antibody MMS-85/12 mRNA, partial cds {IMAGE:945585}	945585 IMAGE Consortium	AA546085
48 G	11 021MA1E8	ESTs {IMAGE:945711}	945711 IMAGE Consortium	AA546205
49 A	1 021TA1E9	IMAGE EST {IMAGE:775033}	775033 IMAGE Consortium	AA387356
50 A	3 021TA1E9	Mus musculus mRNA for protein L, partial cds {IMAGE:933939}	933939 IMAGE Consortium	AA545633
51 A	5 021TA1E9	IMAGE EST {IMAGE:805306}	805306 IMAGE Consortium	AA473329
52 A	7 021TA1E9	ESTs, Highly similar to TRANSCRIPTION INITIATION FACTOR IIE, BETA SUBUNIT [H.sapiens] {IMAGE:833858}	833858 IMAGE Consortium	AA412879
53 A	9 021TA1E9	ESTs, Highly similar to Cdc5-like protein [R.norvegicus] {IMAGE:833661}	833661 IMAGE Consortium	AA466838
54 A	11 021TA1E9	IMAGE EST {IMAGE:760616}	760616 IMAGE Consortium	AA386522
55 C	1 021TA1E9	IMAGE EST {IMAGE:760808}	760808 IMAGE Consortium	AA386657
56 C	3 021TA1E9	IMAGE EST {IMAGE:894135}	894135 IMAGE Consortium	AA516975
57 C	5 021TA1E9	IMAGE EST {IMAGE:761095}	761095 IMAGE Consortium	AA387662
58 C	7 021TA1E9	IMAGE EST {IMAGE:774890}	774890 IMAGE Consortium	AA387079
59 C	9 021TA1E9	IMAGE EST {IMAGE:804889}	804889 IMAGE Consortium	AA472871
60 C	11 021TA1E9	ESTs {IMAGE:805129}	805129 IMAGE Consortium	AA473234
61 E	1 021TA1E9	ESTs {IMAGE:804709}	804709 IMAGE Consortium	AA474386
62 E	3 021TA1E9	ESTs, Weakly similar to 36k hydrophobic nucleolar protein [M.musculus] {IMAGE:805549}	805549 IMAGE Consortium	AA397253
63 E	5 021TA1E9	Max binding protein {IMAGE:834147}	834147 IMAGE Consortium	AA413113
64 E	7 021TA1E9	ESTs {IMAGE:834900}	834900 IMAGE Consortium	AA467249
65 E	9 021TA1E9	ESTs, Highly similar to TRANSCRIPTION FACTOR IIIA [Xenopus laevis] {IMAGE:760965}	760965 IMAGE Consortium	AA386829
66 E	11 021TA1E9	IMAGE EST {IMAGE:932748}	932748 IMAGE Consortium	AA538419
67 G	1 021TA1E9	ESTs {IMAGE:760606}	760606 IMAGE Consortium	AA386539
68 G	3 021TA1E9	Mus musculus Ankhzn mRNA, complete cds {IMAGE:894596}	894596 IMAGE Consortium	AA517717
69 G	5 021TA1E9	ESTs {IMAGE:933135}	933135 IMAGE Consortium	AA538173
70 G	7 021TA1E9	IMAGE EST {IMAGE:774798}	774798 IMAGE Consortium	AA386741
71 G	9 021TA1E9	IMAGE EST {IMAGE:775326}	775326 IMAGE Consortium	AA387120
72 G	11 021TA1E9	ESTs {IMAGE:790071}	790071 IMAGE Consortium	AA388068
73 A	1 021DA1EA	IMAGE EST {IMAGE:779711}	779711 IMAGE Consortium	AA414632

74 A	3 021DA1EA	ESTs, Highly similar to ZINC FINGER PROTEIN 18 [Homo sapiens] {IMAGE:803658}	803658 IMAGE Consortium	AA422560
75 A	5 021DA1EA	IMAGE EST {IMAGE:804181}	804181 IMAGE Consortium	AA474405
76 A	7 021DA1EA	ESTs {IMAGE:833451}	833451 IMAGE Consortium	AA465960
77 A	9 021DA1EA	Mus musculus serine/threonine kinase (sak-a) mRNA, complete cds {IMAGE:836382}	836382 IMAGE Consortium	AA426866
78 A	11 021DA1EA	ESTs {IMAGE:793064}	793064 IMAGE Consortium	AA444244
79 C	1 021DA1EA	Mus musculus mRNA for kinesin-like protein 2beta {IMAGE:803803}	803803 IMAGE Consortium	AA422958
80 C	3 021DA1EA	Early growth response 1 {IMAGE:833284}	833284 IMAGE Consortium	AA465846
81 C	5 021DA1EA	ESTs, Weakly similar to nuclear factor YC [M.musculus] {IMAGE:836484}	836484 IMAGE Consortium	AA426942
82 C	7 021DA1EA	IMAGE EST {IMAGE:793017}	793017 IMAGE Consortium	AA437635
83 C	9 021DA1EA	IMAGE EST {IMAGE:779264}	779264 IMAGE Consortium	AA466982
84 C	11 021DA1EA	ESTs, Weakly similar to ZINC FINGER PROTEIN ZFP-26 [M.musculus] {IMAGE:793764}	793764 IMAGE Consortium	AA415708
85 E	1 021DA1EA	ESTs, Highly similar to SET PROTEIN [Homo sapiens] {IMAGE:780092}	780092 IMAGE Consortium	AA414765
86 E	3 021DA1EA	CCAAT/enhancer binding protein alpha (C/EBP), related sequence 1 {IMAGE:790966}	790966 IMAGE Consortium	AA473938
87 E	5 021DA1EA	ESTs, Weakly similar to L8004.7 gene product [S.cerevisiae] {IMAGE:835792}	835792 IMAGE Consortium	AA422581
88 E	7 021DA1EA	ESTs, Moderately similar to S1-1 protein [R.norvegicus] {IMAGE:792947}	792947 IMAGE Consortium	AA437614
89 E	9 021DA1EA	ESTs {IMAGE:779026}	779026 IMAGE Consortium	AA414072
90 E	11 021DA1EA	ESTs {IMAGE:803824}	803824 IMAGE Consortium	AA422967
91 G	1 021DA1EA	IMAGE EST {IMAGE:779624}	779624 IMAGE Consortium	AA413821
92 G	3 021DA1EA	IMAGE EST {IMAGE:803600}	803600 IMAGE Consortium	AA422752
93 G	5 021DA1EA	IMAGE EST {IMAGE:791255}	791255 IMAGE Consortium	AA416321
94 G	7 021DA1EA	Ecotropic viral integration site 5 {IMAGE:791360}	791360 IMAGE Consortium	AA415421
95 G	9 021DA1EA	IMAGE EST {IMAGE:835819}	835819 IMAGE Consortium	AA422616
96 G	11 021DA1EA	ESTs, Weakly similar to ELL [M.musculus] {IMAGE:944891}	944891 IMAGE Consortium	AA545429
97 A	1 021KA1EB	IMAGE EST {IMAGE:775514}	775514 IMAGE Consortium	AA388288
98 A	3 021KA1EB	Mus musculus mRNA for protein encoded by BCL7B gene {IMAGE:775379}	775379 IMAGE Consortium	AA387248
99 A	5 021KA1EB	ESTs {IMAGE:789701}	789701 IMAGE Consortium	AA387431
100 A	7 021KA1EB	IMAGE EST {IMAGE:774895}	774895 IMAGE Consortium	AA387095
101 A	9 021KA1EB	ESTs {IMAGE:775004}	775004 IMAGE Consortium	AA387064
102 A	11 021KA1EB	IMAGE EST {IMAGE:775208}	775208 IMAGE Consortium	AA387199
103 C	1 021KA1EB	IMAGE EST {IMAGE:790122}	790122 IMAGE Consortium	AA387971
104 C	3 021KA1EB	Maternal embryonic message 3 {IMAGE:775464}	775464 IMAGE Consortium	AA388122
105 C	5 021KA1EB	ESTs, Highly similar to ISOCITRATE DEHYDROGENASE [Bos taurus] {IMAGE:775189}	775189 IMAGE Consortium	AA387198
106 C	7 021KA1EB	ESTs, Weakly similar to Tpr gene product [H.sapiens]	789711 IMAGE Consortium	AA387491

		{IMAGE:789711}		
107 C	9	021KA1EB ESTs {IMAGE:789815}	789815 IMAGE Consortium	AA388172
108 C	11	021KA1EB IMAGE EST {IMAGE:778301}	778301 IMAGE Consortium	AA413894
109 E	1	021KA1EB ESTs, Highly similar to p105 coactivator [R.norvegicus] {IMAGE:779592}	779592 IMAGE Consortium	AA413831
110 E	3	021KA1EB ESTs {IMAGE:779755}	779755 IMAGE Consortium	AA414614
111 E	5	021KA1EB ESTs, Moderately similar to the KIAA0197 gene is expressed ubiquitously. [H.sapiens] {IMAGE:790971}	790971 IMAGE Consortium	AA415358
112 E	7	021KA1EB ESTs, Moderately similar to KIAA0368 [H.sapiens] {IMAGE:803637}	803637 IMAGE Consortium	AA422809
113 E	9	021KA1EB ESTs {IMAGE:835873}	835873 IMAGE Consortium	AA422200
114 E	11	021KA1EB ESTs {IMAGE:792954}	792954 IMAGE Consortium	AA437708
115 G	1	021KA1EB IMAGE EST {IMAGE:803949}	803949 IMAGE Consortium	AA435335
116 G	3	021KA1EB ESTs {IMAGE:791505}	791505 IMAGE Consortium	AA437433
117 G	5	021KA1EB ESTs, Highly similar to DTDP-4-DEHYDRORHAMNOSE REDUCTASE [Salmonella typhimurium] {IMAGE:836126}	836126 IMAGE Consortium	AA426893
118 G	7	021KA1EB ESTs {IMAGE:779165}	779165 IMAGE Consortium	AA467578
119 G	9	021KA1EB ESTs, Highly similar to EUKARYOTIC INITIATION FACTOR 4A-LIKE NUK-34 [Homo sapiens] {IMAGE:779315}	779315 IMAGE Consortium	AA414089
120 G	11	021KA1EB ESTs, Weakly similar to predicted using Genefinder [C.elegans] {IMAGE:793605}	793605 IMAGE Consortium	AA415799
121 A	1	021RA1EC ESTs {IMAGE:944409}	944409 IMAGE Consortium	AA545589
122 A	3	021RA1EC ESTs {IMAGE:944330}	944330 IMAGE Consortium	AA545518
123 A	5	021RA1EC Protein kinase C, beta {IMAGE:317270}	317270 IMAGE Consortium	W13213
124 A	7	021RA1EC ESTs {IMAGE:317466}	317466 IMAGE Consortium	W34061
125 A	9	021RA1EC IMAGE EST {IMAGE:314717}	314717 IMAGE Consortium	W09623
126 A	11	021RA1EC EST {IMAGE:318969}	318969 IMAGE Consortium	W34482
127 C	1	021RA1EC Hemochromatosis {IMAGE:313224}	313224 IMAGE Consortium	W11889
128 C	3	021RA1EC ESTs, Weakly similar to predicted using Genefinder [C.elegans] {IMAGE:313261}	313261 IMAGE Consortium	W11916
129 C	5	021RA1EC Mus musculus diacylglycerol acyltransferase (Dgat) mRNA, complete cds {IMAGE:313859}	313859 IMAGE Consortium	W10786
130 C	7	021RA1EC ESTs, Weakly similar to weak similarity with SINR protein [C.elegans] {IMAGE:314146}	314146 IMAGE Consortium	W09723
131 C	9	021RA1EC IMAGE EST {IMAGE:314562}	314562 IMAGE Consortium	W12782
132 C	11	021RA1EC IMAGE EST {IMAGE:315676}	315676 IMAGE Consortium	W09957
133 E	1	021RA1EC ESTs, Weakly similar to similar to peptidase family C19 [C.elegans] {IMAGE:313741}	313741 IMAGE Consortium	W10753
134 E	3	021RA1EC IMAGE EST {IMAGE:314290}	314290 IMAGE Consortium	W12679
135 E	5	021RA1EC ESTs, Weakly similar to HYPOTHETICAL 25.2 KD PROTEIN IN ACB1-KSS1 INTERGENIC REGION [S.cerevisiae] {IMAGE:317894}	317894 IMAGE Consortium	W12956
136 E	7	021RA1EC IMAGE EST {IMAGE:318815}	318815 IMAGE Consortium	W34362
137 E	9	021RA1EC ESTs, Weakly similar to vacuolar proton-ATPase subunit	329703 IMAGE Consortium	W11739

		M9.2 [H.sapiens] {IMAGE:329703}		
138 E	11 021RA1EC	ESTs {IMAGE:329863}	329863 IMAGE Consortium	W14142
139 G	1 021RA1EC	IMAGE EST {IMAGE:315773}	315773 IMAGE Consortium	W09924
140 G	3 021RA1EC	IMAGE EST {IMAGE:315913}	315913 IMAGE Consortium	W08983
141 G	5 021RA1EC	IMAGE EST {IMAGE:316467}	316467 IMAGE Consortium	W11186
142 G	7 021RA1EC	ESTs {IMAGE:316852}	316852 IMAGE Consortium	W11347
143 G	9 021RA1EC	ESTs, Moderately similar to CYTOCHROME P450 IVF3 [Homo sapiens] {IMAGE:330995}	330995 IMAGE Consortium	W14009
144 G	11 021RA1EC	IMAGE EST {IMAGE:317396}	317396 IMAGE Consortium	W34044
145 A	1 021YA1ED	ESTs {IMAGE:330237}	330237 IMAGE Consortium	W14738
146 A	3 021YA1ED	ESTs {IMAGE:316914}	316914 IMAGE Consortium	W11926
147 A	5 021YA1ED	ESTs, Highly similar to SEC14L [H.sapiens] {IMAGE:313396}	313396 IMAGE Consortium	W10134
148 A	7 021YA1ED	M.musculus spasmodic polyprotein (SP) mRNA {IMAGE:314078}	314078 IMAGE Consortium	W09829
149 A	9 021YA1ED	IMAGE EST {IMAGE:315025}	315025 IMAGE Consortium	W12569
150 A	11 021YA1ED	ATPase, Ca <sup>++</sup> transporting, cardiac muscle, fast twitch 1 {IMAGE:318735}	318735 IMAGE Consortium	W34420
151 C	1 021YA1ED	IMAGE EST {IMAGE:314127}	314127 IMAGE Consortium	W09719
152 C	3 021YA1ED	ESTs {IMAGE:314625}	314625 IMAGE Consortium	W10903
153 C	5 021YA1ED	IMAGE EST {IMAGE:315536}	315536 IMAGE Consortium	W09604
154 C	7 021YA1ED	ESTs {IMAGE:316083}	316083 IMAGE Consortium	W08930
155 C	9 021YA1ED	IMAGE EST {IMAGE:316143}	316143 IMAGE Consortium	W12536
156 C	11 021YA1ED	IMAGE EST {IMAGE:316423}	316423 IMAGE Consortium	W11159
157 E	1 021YA1ED	ESTs {IMAGE:316107}	316107 IMAGE Consortium	W12259
158 E	3 021YA1ED	Cytochrome P450, 2b13, phenobarbital inducible, type c {IMAGE:316288}	316288 IMAGE Consortium	W12874
159 E	5 021YA1ED	Mus musculus Balb/c zinc finger protein PZF (Pzf) mRNA, complete cds {IMAGE:316427}	316427 IMAGE Consortium	W11161
160 E	7 021YA1ED	IMAGE EST {IMAGE:317436}	317436 IMAGE Consortium	W13495
161 E	9 021YA1ED	IMAGE EST {IMAGE:317678}	317678 IMAGE Consortium	W11218
162 E	11 021YA1ED	Microsomal triglyceride transfer protein {IMAGE:317806}	317806 IMAGE Consortium	W13053
163 G	1 021YA1ED	IMAGE EST {IMAGE:314044}	314044 IMAGE Consortium	W09811
164 G	3 021YA1ED	SRY-box containing gene 4 {IMAGE:315058}	315058 IMAGE Consortium	W14398
165 G	5 021YA1ED	ESTs {IMAGE:316182}	316182 IMAGE Consortium	W12523
166 G	7 021YA1ED	ESTs, Weakly similar to predicted using Genefinder [C.elegans] {IMAGE:316850}	316850 IMAGE Consortium	W11346
167 G	9 021YA1ED	IMAGE EST {IMAGE:316835}	316835 IMAGE Consortium	W11342
168 G	11 021YA1ED	Prostaglandin E receptor EP1 subtype {IMAGE:331219}	331219 IMAGE Consortium	W14312
169 A	1 0215A1EE	DIFF6 PROTEIN {IMAGE:334138}	334138 IMAGE Consortium	W16222
170 A	3 0215A1EE	ESTs, Highly similar to ZINC FINGER PROTEIN HF.12 [Homo sapiens] {IMAGE:334378}	334378 IMAGE Consortium	W36520
171 A	5 0215A1EE	Mouse s-adenosylmethionine synthetase mRNA {IMAGE:348007}	348007 IMAGE Consortium	W29782

172 A	7 0215A1EE	IMAGE EST {IMAGE:335167}	335167 IMAGE Consortium	W17570
173 A	9 0215A1EE	ESTs, Weakly similar to TRANSCRIPTIONAL ACTIVATOR GCN5 [Saccharomyces cerevisiae] {IMAGE:335158}	335158 IMAGE Consortium	W36963
174 A	11 0215A1EE	ESTs, Weakly similar to weakly similar to ANK repeat region of Fowlpox virus BamHI-orf7 protein [C.elegans] {IMAGE:332328}	332328 IMAGE Consortium	W07874
175 C	1 0215A1EE	ESTs {IMAGE:331549}	331549 IMAGE Consortium	W13929
176 C	3 0215A1EE	IMAGE EST {IMAGE:335302}	335302 IMAGE Consortium	W36450
177 C	5 0215A1EE	ESTs, Weakly similar to scaffold protein Pbp1 homolog [M.musculus] {IMAGE:336296}	336296 IMAGE Consortium	W36584
178 C	7 0215A1EE	IMAGE EST {IMAGE:336375}	336375 IMAGE Consortium	W36659
179 C	9 0215A1EE	ESTs {IMAGE:333346}	333346 IMAGE Consortium	W16259
180 C	11 0215A1EE	ESTs, Highly similar to GLYCOGEN PHOSPHORYLASE, MUSCLE FORM [Oryctolagus cuniculus] {IMAGE:334236}	334236 IMAGE Consortium	W16286
181 E	1 0215A1EE	IMAGE EST {IMAGE:336783}	336783 IMAGE Consortium	W20810
182 E	3 0215A1EE	CLARA CELL PHOSPHOLIPID-BINDING PROTEIN PRECURSOR {IMAGE:337134}	337134 IMAGE Consortium	W36838
183 E	5 0215A1EE	ESTs {IMAGE:337217}	337217 IMAGE Consortium	W20887
184 E	7 0215A1EE	IMAGE EST {IMAGE:334335}	334335 IMAGE Consortium	W16251
185 E	9 0215A1EE	ESTs, Weakly similar to similar to DNA directed RNA polymerase II [C.elegans] {IMAGE:334501}	334501 IMAGE Consortium	AA052404
186 E	11 0215A1EE	ESTs, Highly similar to FIBRINOGEN GAMMA-A CHAIN PRECURSOR [Homo sapiens] {IMAGE:335055}	335055 IMAGE Consortium	W17873
187 G	1 0215A1EE	Mus musculus mRNA for NIPSNAP2 protein {IMAGE:333579}	333579 IMAGE Consortium	W15931
188 G	3 0215A1EE	ESTs, Weakly similar to HYPOTHETICAL 11.0 KD PROTEIN IN FAA3-BET1 INTERGENIC REGION [Saccharomyces cerevisiae] {IMAGE:334008}	334008 IMAGE Consortium	W16162
189 G	5 0215A1EE	ESTs {IMAGE:651903}	651903 IMAGE Consortium	AA212155
190 G	7 0215A1EE	ESTs {IMAGE:331599}	331599 IMAGE Consortium	W08056
191 G	9 0215A1EE	Apurinic/aprimidinic endonuclease {IMAGE:331901}	331901 IMAGE Consortium	W08299
192 G	11 0215A1EE	IMAGE EST {IMAGE:331794}	331794 IMAGE Consortium	W08098
193 A	1 021CA1EF	ESTs, Weakly similar to LL5 protein [R.norvegicus] {IMAGE:316171}	316171 IMAGE Consortium	W12633
194 A	3 021CA1EF	IMAGE EST {IMAGE:316922}	316922 IMAGE Consortium	W11384
195 A	5 021CA1EF	Enolase 3, beta muscle {IMAGE:316967}	316967 IMAGE Consortium	W11965
196 A	7 021CA1EF	EST {IMAGE:317561}	317561 IMAGE Consortium	W12981
197 A	9 021CA1EF	ESTs {IMAGE:317976}	317976 IMAGE Consortium	W11545
198 A	11 021CA1EF	ESTs {IMAGE:317868}	317868 IMAGE Consortium	W12937
199 C	1 021CA1EF	ESTs, Weakly similar to T04A8.4 [C.elegans] {IMAGE:336133}	336133 IMAGE Consortium	W18106
200 C	3 021CA1EF	Mus musculus mRNA for synaptogyrin 2 {IMAGE:333102}	333102 IMAGE Consortium	W15720
201 C	5 021CA1EF	IMAGE EST {IMAGE:333508}	333508 IMAGE Consortium	W33974
202 C	7 021CA1EF	ACETYLCHOLINE RECEPTOR PROTEIN, GAMMA CHAIN PRECURSOR {IMAGE:333948}	333948 IMAGE Consortium	W16128



203 C	9 021CA1EF	ESTs, Moderately similar to probable calcium-binding protein [H.sapiens] {IMAGE:337667}	337667 IMAGE Consortium	W18735
204 C	11 021CA1EF	Mus musculus peroxisomal/mitochondrial dienyoyl-CoA isomerase ECH1p (Ech1) mRNA, complete cds {IMAGE:338088}	338088 IMAGE Consortium	W29607
205 E	1 021CA1EF	ESTs {IMAGE:333037}	333037 IMAGE Consortium	W15681
206 E	3 021CA1EF	ESTs {IMAGE:337203}	337203 IMAGE Consortium	W36867
207 E	5 021CA1EF	IMAGE EST {IMAGE:337096}	337096 IMAGE Consortium	W20765
208 E	7 021CA1EF	ESTs {IMAGE:337077}	337077 IMAGE Consortium	W20763
209 E	9 021CA1EF	ESTs {IMAGE:337452}	337452 IMAGE Consortium	W18661
210 E	11 021CA1EF	IMAGE EST {IMAGE:338162}	338162 IMAGE Consortium	W29797
211 G	1 021CA1EF	ESTs {IMAGE:338152}	338152 IMAGE Consortium	W29532
212 G	3 021CA1EF	ESTs {IMAGE:334906}	334906 IMAGE Consortium	W17831
213 G	5 021CA1EF	IMAGE EST {IMAGE:331400}	331400 IMAGE Consortium	AA052341
214 G	7 021CA1EF	ESTs, Moderately similar to LETHAL(2) GIANT LARVAE PROTEIN [Drosophila melanogaster] {IMAGE:331968}	331968 IMAGE Consortium	W08134
215 G	9 021CA1EF	ESTs {IMAGE:331695}	331695 IMAGE Consortium	W14338
216 G	11 021CA1EF	IMAGE EST {IMAGE:335943}	335943 IMAGE Consortium	W17800
217 A	1 021JA1EG	ESTs, Weakly similar to coded for by C. elegans cDNA yk173c12.5 [C.elegans] {IMAGE:338002}	338002 IMAGE Consortium	W18908
218 A	3 021JA1EG	ESTs {IMAGE:347906}	347906 IMAGE Consortium	W29916
219 A	5 021JA1EG	Cytokine inducible SH2-containing protein {IMAGE:335058}	335058 IMAGE Consortium	W36940
220 A	7 021JA1EG	ESTs, Weakly similar to olfactomedin [M.musculus] {IMAGE:331480}	331480 IMAGE Consortium	W14353
221 A	9 021JA1EG	ESTs, Weakly similar to HYPOTHETICAL 86.9 KD PROTEIN ZK945.3 IN CHROMOSOME II [Caenorhabditis elegans] {IMAGE:331974}	331974 IMAGE Consortium	W08137
222 A	11 021JA1EG	IMAGE EST {IMAGE:331818}	331818 IMAGE Consortium	W08069
223 C	1 021JA1EG	IMAGE EST {IMAGE:347917}	347917 IMAGE Consortium	W29582
224 C	3 021JA1EG	Mus musculus mRNA for legumain {IMAGE:331457}	331457 IMAGE Consortium	W14837
225 C	5 021JA1EG	Mus musculus C57BL/6J ribosomal protein S28 mRNA, complete cds {IMAGE:335746}	335746 IMAGE Consortium	W20956
226 C	7 021JA1EG	ESTs {IMAGE:332100}	332100 IMAGE Consortium	W08169
227 C	9 021JA1EG	IMAGE EST {IMAGE:336515}	336515 IMAGE Consortium	W18516
228 C	11 021JA1EG	ESTs {IMAGE:336306}	336306 IMAGE Consortium	W18265
229 E	1 021JA1EG	Mus musculus von Ebner minor salivary gland protein mRNA, complete cds {IMAGE:335127}	335127 IMAGE Consortium	W17563
230 E	3 021JA1EG	IMAGE EST {IMAGE:335795}	335795 IMAGE Consortium	W34121
231 E	5 021JA1EG	IMAGE EST {IMAGE:332388}	332388 IMAGE Consortium	W07943
232 E	7 021JA1EG	ESTs, Weakly similar to GS2 PROTEIN [H.sapiens] {IMAGE:336564}	336564 IMAGE Consortium	W36670
233 E	9 021JA1EG	Neuroblastoma ras oncogene {IMAGE:333030}	333030 IMAGE Consortium	W15662
234 E	11 021JA1EG	Laminin, alpha 2 {IMAGE:333498}	333498 IMAGE Consortium	W18311

235 G	1	021JA1EG	IMAGE EST {IMAGE:350336}	350336	IMAGE Consortium	W34967
236 G	3	021JA1EG	ESTs, Weakly similar to cDNA EST EMBL:M75904 comes from this gene [C.elegans] {IMAGE:354212}	354212	IMAGE Consortium	W44032
237 G	5	021JA1EG	Mouse NLRR-1 mRNA for leucine-rich-repeat protein, complete cds {IMAGE:351947}	351947	IMAGE Consortium	W40832
238 G	7	021JA1EG	Procollagen, type VI, alpha 1 {IMAGE:352450}	352450	IMAGE Consortium	W33786
239 G	9	021JA1EG	IMAGE EST {IMAGE:350432}	350432	IMAGE Consortium	W34824
240 G	11	021JA1EG	IMAGE EST {IMAGE:350158}	350158	IMAGE Consortium	W34715
241 A	1	021QA1EH	ESTs {IMAGE:352357}	352357	IMAGE Consortium	W33732
242 A	3	021QA1EH	Mus musculus Eph-and Elk-related kinase (eek) mRNA, complete cds {IMAGE:352644}	352644	IMAGE Consortium	W34187
243 A	5	021QA1EH	ESTs, Highly similar to SOH1 PROTEIN [Saccharomyces cerevisiae] {IMAGE:349747}	349747	IMAGE Consortium	W30130
244 A	7	021QA1EH	IMAGE EST {IMAGE:350243}	350243	IMAGE Consortium	W35879
245 A	9	021QA1EH	IMAGE EST {IMAGE:350881}	350881	IMAGE Consortium	W40994
246 A	11	021QA1EH	IMAGE EST {IMAGE:354215}	354215	IMAGE Consortium	W44037
247 C	1	021QA1EH	ESTs {IMAGE:354344}	354344	IMAGE Consortium	W42331
248 C	3	021QA1EH	IMAGE EST {IMAGE:352610}	352610	IMAGE Consortium	W34106
249 C	5	021QA1EH	ESTs, Highly similar to HAM1 PROTEIN [Saccharomyces cerevisiae] {IMAGE:352783}	352783	IMAGE Consortium	W34474
250 C	7	021QA1EH	ESTs {IMAGE:353507}	353507	IMAGE Consortium	W40894
251 C	9	021QA1EH	ESTs {IMAGE:351090}	351090	IMAGE Consortium	W41083
252 C	11	021QA1EH	IMAGE EST {IMAGE:352743}	352743	IMAGE Consortium	W34390
253 E	1	021QA1EH	ESTs {IMAGE:352785}	352785	IMAGE Consortium	W34475
254 E	3	021QA1EH	ESTs, Highly similar to DELTA 1-PYRROLINE-5-CARBOXYLATE SYNTHETASE [Vigna aconitifolia] {IMAGE:353317}	353317	IMAGE Consortium	W41878
255 E	5	021QA1EH	ESTs {IMAGE:350423}	350423	IMAGE Consortium	W40954
256 E	7	021QA1EH	IMAGE EST {IMAGE:351028}	351028	IMAGE Consortium	W35091
257 E	9	021QA1EH	IMAGE EST {IMAGE:354506}	354506	IMAGE Consortium	W45825
258 E	11	021QA1EH	ESTs, Weakly similar to P68-LIKE PROTEIN [Saccharomyces cerevisiae] {IMAGE:352409}	352409	IMAGE Consortium	W33833
259 G	1	021QA1EH	Mus musculus putative serine/threonine protein kinase MAK-V (Mak-v) mRNA, complete cds {IMAGE:353232}	353232	IMAGE Consortium	W41683
260 G	3	021QA1EH	Dlk1-like homolog (Drosophila) {IMAGE:350180}	350180	IMAGE Consortium	W34721
261 G	5	021QA1EH	EST {IMAGE:350863}	350863	IMAGE Consortium	W40989
262 G	7	021QA1EH	IMAGE EST {IMAGE:351045}	351045	IMAGE Consortium	W41210
263 G	9	021QA1EH	IMAGE EST {IMAGE:351420}	351420	IMAGE Consortium	W41309
264 G	11	021QA1EH	IMAGE EST {IMAGE:352601}	352601	IMAGE Consortium	W34486
265 A	1	021XA1EI	IMAGE EST {IMAGE:367915}	367915	IMAGE Consortium	W54533
266 A	3	021XA1EI	IMAGE EST {IMAGE:368264}	368264	IMAGE Consortium	W54145
267 A	5	021XA1EI	IMAGE EST {IMAGE:372454}	372454	IMAGE Consortium	W57116
268 A	7	021XA1EI	IMAGE EST {IMAGE:367447}	367447	IMAGE Consortium	AA052637

269 A	9 021XA1EI	ESTs, Highly similar to NADH-UBIQUINONE OXIDOREDUCTASE B17 SUBUNIT [Bos taurus] {IMAGE:367651}	367651 IMAGE Consortium	W54448
270 A	11 021XA1EI	CALPONIN H1, SMOOTH MUSCLE {IMAGE:373793}	373793 IMAGE Consortium	W64636
271 C	1 021XA1EI	IMAGE EST {IMAGE:367495}	367495 IMAGE Consortium	W54610
272 C	3 021XA1EI	IMAGE EST {IMAGE:367877}	367877 IMAGE Consortium	W53908
273 C	5 021XA1EI	IMAGE EST {IMAGE:367869}	367869 IMAGE Consortium	W54520
274 C	7 021XA1EI	IMAGE EST {IMAGE:373967}	373967 IMAGE Consortium	W62513
275 C	9 021XA1EI	IMAGE EST {IMAGE:355356}	355356 IMAGE Consortium	W48335
276 C	11 021XA1EI	IMAGE EST {IMAGE:367000}	367000 IMAGE Consortium	W54375
277 E	1 021XA1EI	ESTs {IMAGE:373242}	373242 IMAGE Consortium	W57265
278 E	3 021XA1EI	IMAGE EST {IMAGE:373521}	373521 IMAGE Consortium	W57234
279 E	5 021XA1EI	ESTs {IMAGE:368173}	368173 IMAGE Consortium	W54099
280 E	7 021XA1EI	ESTs {IMAGE:387151}	387151 IMAGE Consortium	W65734
281 E	9 021XA1EI	ESTs {IMAGE:387359}	387359 IMAGE Consortium	W65705
282 E	11 021XA1EI	ESTs, Weakly similar to HYPOTHETICAL 6.3 KD PROTEIN ZK652.2 IN CHROMOSOME III [C.elegans] {IMAGE:385341}	385341 IMAGE Consortium	W74865
283 G	1 021XA1EI	ESTs {IMAGE:374725}	374725 IMAGE Consortium	W59264
284 G	3 021XA1EI	Epithelial membrane protein 1 {IMAGE:385084}	385084 IMAGE Consortium	W64798
285 G	5 021XA1EI	IMAGE EST {IMAGE:385318}	385318 IMAGE Consortium	W63815
286 G	7 021XA1EI	IMAGE EST {IMAGE:388625}	388625 IMAGE Consortium	W70734
287 G	9 021XA1EI	ESTs, Moderately similar to PEROXISOME BIOSYNTHESIS PROTEIN PAY4 [Yarrowia lipolytica] {IMAGE:386328}	386328 IMAGE Consortium	W65135
288 G	11 021XA1EI	ESTs, Weakly similar to 3-OXOACYL-[ACYL-CARRIER PROTEIN] REDUCTASE [Escherichia coli] {IMAGE:386555}	386555 IMAGE Consortium	W65003
289 A	1 0214A1EJ	ESTs {IMAGE:350300}	350300 IMAGE Consortium	W41570
290 A	3 0214A1EJ	IMAGE EST {IMAGE:351288}	351288 IMAGE Consortium	W41563
291 A	5 0214A1EJ	Mus musculus RNA polymerase I largest subunit (RPA1) mRNA, complete cds {IMAGE:354709}	354709 IMAGE Consortium	W46057
292 A	7 0214A1EJ	ESTs, Moderately similar to R151.6 [C.elegans] {IMAGE:355887}	355887 IMAGE Consortium	W48017
293 A	9 0214A1EJ	Cytochrome P450, 2e1, ethanol inducible {IMAGE:367313}	367313 IMAGE Consortium	W50771
294 A	11 0214A1EJ	ESTs {IMAGE:367785}	367785 IMAGE Consortium	W53989
295 C	1 0214A1EJ	Glutamic acid decarboxylase 1 {IMAGE:374236}	374236 IMAGE Consortium	W59173
296 C	3 0214A1EJ	DEOXYCYTIDINE KINASE {IMAGE:373964}	373964 IMAGE Consortium	W62508
297 C	5 0214A1EJ	ESTs {IMAGE:355454}	355454 IMAGE Consortium	W48305
298 C	7 0214A1EJ	IMAGE EST {IMAGE:367123}	367123 IMAGE Consortium	W54582
299 C	9 0214A1EJ	IMAGE EST {IMAGE:367445}	367445 IMAGE Consortium	W50706
300 C	11 0214A1EJ	ESTs {IMAGE:367487}	367487 IMAGE Consortium	W54436
301 E	1 0214A1EJ	IMAGE EST {IMAGE:367765}	367765 IMAGE Consortium	W53959
302 E	3 0214A1EJ	Mouse leukotriene A-4 hydrolase mRNA, complete cds {IMAGE:374156}	374156 IMAGE Consortium	W63957
303 E	5 0214A1EJ	IMAGE EST {IMAGE:355025}	355025 IMAGE Consortium	W47839

304 E	7 0214A1EJ	IMAGE EST {IMAGE:355449}	355449 IMAGE Consortium	W48309
305 E	9 0214A1EJ	Laminin, gamma 2 {IMAGE:355793}	355793 IMAGE Consortium	W49392
306 E	11 0214A1EJ	ESTs {IMAGE:373112}	373112 IMAGE Consortium	W62546
307 G	1 0214A1EJ	ESTs {IMAGE:373693}	373693 IMAGE Consortium	W64547
308 G	3 0214A1EJ	IMAGE EST {IMAGE:374228}	374228 IMAGE Consortium	W63981
309 G	5 0214A1EJ	ESTs {IMAGE:367317}	367317 IMAGE Consortium	W50773
310 G	7 0214A1EJ	Biglycan {IMAGE:367435}	367435 IMAGE Consortium	W54287
311 G	9 0214A1EJ	IMAGE EST {IMAGE:374148}	374148 IMAGE Consortium	W59002
312 G	11 0214A1EJ	ESTs, Weakly similar to ORF YGR038w [S.cerevisiae] {IMAGE:354801}	354801 IMAGE Consortium	W44283
313 A	1 021BA1EK	Myeloid differentiation primary response gene 118 {IMAGE:386412}	386412 IMAGE Consortium	W64388
314 A	3 021BA1EK	Jumonji {IMAGE:386593}	386593 IMAGE Consortium	W65025
315 A	5 021BA1EK	ESTs {IMAGE:386901}	386901 IMAGE Consortium	W65570
316 A	7 021BA1EK	ESTs, Weakly similar to nuclear factor YC [M.musculus] {IMAGE:386962}	386962 IMAGE Consortium	W65723
317 A	9 021BA1EK	ESTs, Highly similar to GOLIATH PROTEIN [Drosophila melanogaster] {IMAGE:388008}	388008 IMAGE Consortium	W70414
318 A	11 021BA1EK	IMAGE EST {IMAGE:385063}	385063 IMAGE Consortium	W64752
319 C	1 021BA1EK	IMAGE EST {IMAGE:374581}	374581 IMAGE Consortium	W59380
320 C	3 021BA1EK	IMAGE EST {IMAGE:387034}	387034 IMAGE Consortium	W65713
321 C	5 021BA1EK	IMAGE EST {IMAGE:387493}	387493 IMAGE Consortium	W65940
322 C	7 021BA1EK	IMAGE EST {IMAGE:385033}	385033 IMAGE Consortium	W64594
323 C	9 021BA1EK	ESTs, Highly similar to SINGLE-STRANDED DNA-BINDING PROTEIN MSSP-1 [Homo sapiens] {IMAGE:388290}	388290 IMAGE Consortium	W66636
324 C	11 021BA1EK	Mus musculus NEDD8-conjugating enzyme (Uba3) mRNA, complete cds {IMAGE:385740}	385740 IMAGE Consortium	W62229
325 E	1 021BA1EK	ESTs {IMAGE:385383}	385383 IMAGE Consortium	W70855
326 E	3 021BA1EK	Mus musculus A10 mRNA, partial cds {IMAGE:385441}	385441 IMAGE Consortium	W61383
327 E	5 021BA1EK	ESTs {IMAGE:385970}	385970 IMAGE Consortium	W62316
328 E	7 021BA1EK	Mus musculus neural visinin-like protein 3 (NVP-3) mRNA, complete cds {IMAGE:389446}	389446 IMAGE Consortium	W70646
329 E	9 021BA1EK	ESTs {IMAGE:386486}	386486 IMAGE Consortium	W64982
330 E	11 021BA1EK	IMAGE EST {IMAGE:390307}	390307 IMAGE Consortium	W74928
331 G	1 021BA1EK	Mus musculus apoptosis activator Mtd (Mtd) mRNA, complete cds {IMAGE:386559}	386559 IMAGE Consortium	W65005
332 G	3 021BA1EK	IMAGE EST {IMAGE:390236}	390236 IMAGE Consortium	W74907
333 G	5 021BA1EK	IMAGE EST {IMAGE:387243}	387243 IMAGE Consortium	W65748
334 G	7 021BA1EK	Mouse mRNA for AF1q, complete cds {IMAGE:374756}	374756 IMAGE Consortium	W61574
335 G	9 021BA1EK	ESTs {IMAGE:385380}	385380 IMAGE Consortium	W70850
336 G	11 021BA1EK	TEA domain family member 2 {IMAGE:388244}	388244 IMAGE Consortium	W66622
337 A	1 021IA1EL	Mus musculus putative histone deacetylase (HD1) mRNA, partial cds {IMAGE:374877}	374877 IMAGE Consortium	W62765

338 A	3 021IA1EL	ESTs, Highly similar to PUTATIVE REGULATORY PROTEIN TSC-22 [Mus musculus; Rattus norvegicus] {IMAGE:387449}	387449 IMAGE Consortium	W66757
339 A	5 021IA1EL	IMAGE EST {IMAGE:385035}	385035 IMAGE Consortium	W64595
340 A	7 021IA1EL	IMAGE EST {IMAGE:388477}	388477 IMAGE Consortium	W71008
341 A	9 021IA1EL	DNA ligase I, ATP-dependent {IMAGE:388245}	388245 IMAGE Consortium	W66626
342 A	11 021IA1EL	ESTs {IMAGE:385693}	385693 IMAGE Consortium	W62920
343 C	1 021IA1EL	ESTs {IMAGE:385141}	385141 IMAGE Consortium	W64732
344 C	3 021IA1EL	ESTs, Moderately similar to MEMBRANE-ASSOCIATED PROTEIN HEM-2 [Mus musculus] {IMAGE:388365}	388365 IMAGE Consortium	W66661
345 C	5 021IA1EL	IMAGE EST {IMAGE:385997}	385997 IMAGE Consortium	W62360
346 C	7 021IA1EL	Hairy and enhancer of split 1, (Drosophila) {IMAGE:386552}	386552 IMAGE Consortium	W64999
347 C	9 021IA1EL	IMAGE EST {IMAGE:386491}	386491 IMAGE Consortium	W65010
348 C	11 021IA1EL	Mus musculus Grb-2-related monocytic adapter protein (Mona) mRNA, complete cds {IMAGE:390410}	390410 IMAGE Consortium	W71812
349 E	1 021IA1EL	Homeo box A2 {IMAGE:374538}	374538 IMAGE Consortium	W63822
350 E	3 021IA1EL	Adenylate cyclase 7 {IMAGE:387280}	387280 IMAGE Consortium	W65619
351 E	5 021IA1EL	IMAGE EST {IMAGE:385371}	385371 IMAGE Consortium	W70852
352 E	7 021IA1EL	ESTs {IMAGE:388776}	388776 IMAGE Consortium	W71130
353 E	9 021IA1EL	IMAGE EST {IMAGE:385528}	385528 IMAGE Consortium	W64177
354 E	11 021IA1EL	Mus musculus growth/differentiation factor 1 (GDF-1) mRNA, complete cds {IMAGE:386194}	386194 IMAGE Consortium	W65054
355 G	1 021IA1EL	IMAGE EST {IMAGE:401213}	401213 IMAGE Consortium	W98091
356 G	3 021IA1EL	ESTs {IMAGE:404345}	404345 IMAGE Consortium	W82800
357 G	5 021IA1EL	Mus musculus rab6/rab5-associated protein (rab6) mRNA, partial cds {IMAGE:401958}	401958 IMAGE Consortium	W77711
358 G	7 021IA1EL	ESTs, Highly similar to HETEROGENEOUS NUCLEAR RIBONUCLEOPROTEIN 27C [Drosophila melanogaster] {IMAGE:402547}	402547 IMAGE Consortium	W80268
359 G	9 021IA1EL	IMAGE EST {IMAGE:402995}	402995 IMAGE Consortium	W81775
360 G	11 021IA1EL	ESTs {IMAGE:403084}	403084 IMAGE Consortium	W81999
361 A	1 021PA1EM	IMAGE EST {IMAGE:403550}	403550 IMAGE Consortium	W82188
362 A	3 021PA1EM	ESTs, Weakly similar to LONG-CHAIN-FATTY-ACID--COA LIGASE, BRAIN ISOZYME [R.norvegicus] {IMAGE:404162}	404162 IMAGE Consortium	W82464
363 A	5 021PA1EM	IMAGE EST {IMAGE:404198}	404198 IMAGE Consortium	W82549
364 A	7 021PA1EM	IMAGE EST {IMAGE:404233}	404233 IMAGE Consortium	W82617
365 A	9 021PA1EM	IMAGE EST {IMAGE:404510}	404510 IMAGE Consortium	W83102
366 A	11 021PA1EM	ESTs, Highly similar to RIBOPHORIN I PRECURSOR [Rattus norvegicus] {IMAGE:405823}	405823 IMAGE Consortium	W83009
367 C	1 021PA1EM	Ancient ubiquitous protein {IMAGE:405768}	405768 IMAGE Consortium	W82989
368 C	3 021PA1EM	ESTs {IMAGE:402401}	402401 IMAGE Consortium	W78377
369 C	5 021PA1EM	ESTs, Weakly similar to PROBABLE RNA HELICASE IN WAPA-LICT INTERGENIC REGION [Bacillus subtilis] {IMAGE:403381}	403381 IMAGE Consortium	W82059

370 C	7 021PA1EM	IMAGE EST {IMAGE:403735}	403735 IMAGE Consortium	W82219
371 C	9 021PA1EM	IMAGE EST {IMAGE:400432}	400432 IMAGE Consortium	W77330
372 C	11 021PA1EM	ESTs, Highly similar to p82 [H.sapiens] {IMAGE:401305}	401305 IMAGE Consortium	W98251
373 E	1 021PA1EM	IMAGE EST {IMAGE:403453}	403453 IMAGE Consortium	W81953
374 E	3 021PA1EM	IMAGE EST {IMAGE:400504}	400504 IMAGE Consortium	W76719
375 E	5 021PA1EM	ESTs {IMAGE:403864}	403864 IMAGE Consortium	W82300
376 E	7 021PA1EM	Insulin-like growth factor binding protein 1 {IMAGE:404595}	404595 IMAGE Consortium	W83086
377 E	9 021PA1EM	Mus musculus neural precursor cell expressed developmentally downregulated Nedd9 (Nedd9) mRNA, complete cds {IMAGE:404536}	404536 IMAGE Consortium	W82737
378 E	11 021PA1EM	IMAGE EST {IMAGE:402641}	402641 IMAGE Consortium	W79958
379 G	1 021PA1EM	IMAGE EST {IMAGE:405842}	405842 IMAGE Consortium	W83011
380 G	3 021PA1EM	IMAGE EST {IMAGE:418701}	418701 IMAGE Consortium	W96997
381 G	5 021PA1EM	ESTs {IMAGE:419138}	419138 IMAGE Consortium	W88001
382 G	7 021PA1EM	IMAGE EST {IMAGE:419003}	419003 IMAGE Consortium	W89268
383 G	9 021PA1EM	ESTs, Highly similar to teg292 protein [M.musculus] {IMAGE:407480}	407480 IMAGE Consortium	W89998
384 G	11 021PA1EM	Mesenchyme homeobox 1 {IMAGE:419684}	419684 IMAGE Consortium	W89502
385 A	1 021WA1EN	M.musculus ART3 gene {IMAGE:404348}	404348 IMAGE Consortium	W82798
386 A	3 021WA1EN	ESTs {IMAGE:402667}	402667 IMAGE Consortium	W79994
387 A	5 021WA1EN	IMAGE EST {IMAGE:403068}	403068 IMAGE Consortium	W81793
388 A	7 021WA1EN	Aryl-hydrocarbon receptor {IMAGE:403384}	403384 IMAGE Consortium	W82873
389 A	9 021WA1EN	IMAGE EST {IMAGE:403682}	403682 IMAGE Consortium	W82201
390 A	11 021WA1EN	ESTs {IMAGE:403523}	403523 IMAGE Consortium	W82182
391 C	1 021WA1EN	IMAGE EST {IMAGE:402757}	402757 IMAGE Consortium	W80109
392 C	3 021WA1EN	Protamine 3 {IMAGE:400035}	400035 IMAGE Consortium	W77113
393 C	5 021WA1EN	IMAGE EST {IMAGE:403667}	403667 IMAGE Consortium	W82278
394 C	7 021WA1EN	IMAGE EST {IMAGE:403823}	403823 IMAGE Consortium	W82759
395 C	9 021WA1EN	IMAGE EST {IMAGE:401192}	401192 IMAGE Consortium	W98130
396 C	11 021WA1EN	IMAGE EST {IMAGE:401773}	401773 IMAGE Consortium	W78544
397 E	1 021WA1EN	ESTs {IMAGE:404058}	404058 IMAGE Consortium	W82574
398 E	3 021WA1EN	ESTs {IMAGE:404339}	404339 IMAGE Consortium	W82690
399 E	5 021WA1EN	ESTs, Weakly similar to similar to beta-mannosyltransferase [C.elegans] {IMAGE:404421}	404421 IMAGE Consortium	W82843
400 E	7 021WA1EN	CALPONIN H2, SMOOTH MUSCLE {IMAGE:402348}	402348 IMAGE Consortium	W78589
401 E	9 021WA1EN	IMAGE EST {IMAGE:403003}	403003 IMAGE Consortium	W81856
402 E	11 021WA1EN	Mus musculus mRNA for oxysterol-binding protein, complete cds {IMAGE:403493}	403493 IMAGE Consortium	W81823
403 G	1 021WA1EN	Cystatin 3 {IMAGE:402614}	402614 IMAGE Consortium	W78651
404 G	3 021WA1EN	IMAGE EST {IMAGE:403062}	403062 IMAGE Consortium	W81917
405 G	5 021WA1EN	ESTs {IMAGE:403449}	403449 IMAGE Consortium	W82080
406 G	7 021WA1EN	IMAGE EST {IMAGE:403478}	403478 IMAGE Consortium	W82131
407 G	9 021WA1EN	IMAGE EST {IMAGE:403946}	403946 IMAGE Consortium	W82313

408 G	11 021WA1EN	ESTs {IMAGE:403933}	403933 IMAGE Consortium	W82178
409 A	1 0213A1EO	PEPTIDYL-PROLYL CIS-TRANS ISOMERASE C {IMAGE:406947}	406947 IMAGE Consortium	W83904
410 A	3 0213A1EO	IMAGE EST {IMAGE:407081}	407081 IMAGE Consortium	W84179
411 A	5 0213A1EO	M.musculus p16K gene for 16 kDa protein {IMAGE:407552}	407552 IMAGE Consortium	W88130
412 A	7 0213A1EO	MutS homolog 2 (E. coli) {IMAGE:419771}	419771 IMAGE Consortium	W91178
413 A	9 0213A1EO	ESTs, Weakly similar to predicted using Genefinder [C.elegans] {IMAGE:408379}	408379 IMAGE Consortium	W85373
414 A	11 0213A1EO	IMAGE EST {IMAGE:408542}	408542 IMAGE Consortium	W85052
415 C	1 0213A1EO	Carboxypeptidase E {IMAGE:407068}	407068 IMAGE Consortium	W83974
416 C	3 0213A1EO	ESTs {IMAGE:407202}	407202 IMAGE Consortium	W84227
417 C	5 0213A1EO	ESTs, Highly similar to c-myc binding protein [H.sapiens] {IMAGE:407556}	407556 IMAGE Consortium	W88132
418 C	7 0213A1EO	IMAGE EST {IMAGE:407924}	407924 IMAGE Consortium	W85158
419 C	9 0213A1EO	ESTs {IMAGE:408189}	408189 IMAGE Consortium	AA052699
420 C	11 0213A1EO	IMAGE EST {IMAGE:420149}	420149 IMAGE Consortium	AA052087
421 E	1 0213A1EO	Mus musculus growth arrest specific mRNA, clone 3546, complete cds {IMAGE:406745}	406745 IMAGE Consortium	W83881
422 E	3 0213A1EO	ESTs {IMAGE:419243}	419243 IMAGE Consortium	AA052194
423 E	5 0213A1EO	IMAGE EST {IMAGE:407558}	407558 IMAGE Consortium	W88133
424 E	7 0213A1EO	IMAGE EST {IMAGE:419944}	419944 IMAGE Consortium	W89598
425 E	9 0213A1EO	ESTs, Moderately similar to PUTATIVE MITOCHONDRIAL RIBOSOMAL PROTEIN S14 [Caenorhabditis elegans] {IMAGE:419614}	419614 IMAGE Consortium	W89487
426 E	11 0213A1EO	C-src tyrosine kinase {IMAGE:420269}	420269 IMAGE Consortium	W89819
427 G	1 0213A1EO	ESTs {IMAGE:406999}	406999 IMAGE Consortium	W83959
428 G	3 0213A1EO	Solute carrier family 4 (anion exchanger), member 1 {IMAGE:407275}	407275 IMAGE Consortium	W85633
429 G	5 0213A1EO	ESTs, Highly similar to RAS GTPASE-ACTIVATING-LIKE PROTEIN IQGAP1 [Homo sapiens] {IMAGE:408007}	408007 IMAGE Consortium	W84270
430 G	7 0213A1EO	ESTs, Highly similar to ENDOTHELIN-CONVERTING ENZYME 1 E-1 [Rattus norvegicus] {IMAGE:419633}	419633 IMAGE Consortium	W89492
431 G	9 0213A1EO	ESTs, Moderately similar to HYPOTHETICAL 21.5 KD PROTEIN C08B11.9 IN CHROMOSOME II [Caenorhabditis elegans] {IMAGE:409092}	409092 IMAGE Consortium	W87153
432 G	11 0213A1EO	IMAGE EST {IMAGE:418633}	418633 IMAGE Consortium	W99231
433 A	1 021AA1EP	ESTs {IMAGE:406991}	406991 IMAGE Consortium	W84169
434 A	3 021AA1EP	IMAGE EST {IMAGE:407489}	407489 IMAGE Consortium	W89978
435 A	5 021AA1EP	IMAGE EST {IMAGE:408435}	408435 IMAGE Consortium	W85480
436 A	7 021AA1EP	ESTs {IMAGE:408143}	408143 IMAGE Consortium	AA052697
437 A	9 021AA1EP	Mouse mRNA for the third largest RNA polymerase II subunit, complete cds {IMAGE:408599}	408599 IMAGE Consortium	W85217
438 A	11 021AA1EP	Glutathione S-transferase, pi 2 {IMAGE:408997}	408997 IMAGE Consortium	W87197
439 C	1 021AA1EP	Decay accelerating factor 1 {IMAGE:407096}	407096 IMAGE Consortium	W83988

440 C	3 021AA1EP	ESTs {IMAGE:407377}	407377 IMAGE Consortium	AA052165
441 C	5 021AA1EP	Cytochrome P450, 2e1, ethanol inducible {IMAGE:419957}	419957 IMAGE Consortium	W89513
442 C	7 021AA1EP	IMAGE EST {IMAGE:419921}	419921 IMAGE Consortium	W89594
443 C	9 021AA1EP	IMAGE EST {IMAGE:419855}	419855 IMAGE Consortium	W91241
444 C	11 021AA1EP	Transformation related protein 53 {IMAGE:409069}	409069 IMAGE Consortium	W87149
445 E	1 021AA1EP	Chemokine (C-X-C) receptor 4 {IMAGE:407569}	407569 IMAGE Consortium	W88094
446 E	3 021AA1EP	Transcription factor AP-2, gamma {IMAGE:419886}	419886 IMAGE Consortium	W89370
447 E	5 021AA1EP	Fas-associating protein with death domain {IMAGE:419798}	419798 IMAGE Consortium	W91183
448 E	7 021AA1EP	IMAGE EST {IMAGE:408697}	408697 IMAGE Consortium	AA052118
449 E	9 021AA1EP	X-linked adrenoleukodystrophy (ALD) gene homolog {IMAGE:420294}	420294 IMAGE Consortium	W89869
450 E	11 021AA1EP	Mus musculus friend of GATA-1 (FOG) mRNA, complete cds {IMAGE:418539}	418539 IMAGE Consortium	W99029
451 G	1 021AA1EP	EST {IMAGE:419518}	419518 IMAGE Consortium	W88205
452 G	3 021AA1EP	ESTs {IMAGE:419887}	419887 IMAGE Consortium	W89374
453 G	5 021AA1EP	IMAGE EST {IMAGE:408254}	408254 IMAGE Consortium	W85513
454 G	7 021AA1EP	IMAGE EST {IMAGE:408857}	408857 IMAGE Consortium	W85520
455 G	9 021AA1EP	Mus musculus HIC-5 mRNA, complete cds {IMAGE:418546}	418546 IMAGE Consortium	W99117
456 G	11 021AA1EP	ESTs {IMAGE:418790}	418790 IMAGE Consortium	W97092
457 A	1 021HA1EQ	Ryanodine receptor 1, skeletal muscle {IMAGE:422161}	422161 IMAGE Consortium	W97384
458 A	3 021HA1EQ	IMAGE EST {IMAGE:420494}	420494 IMAGE Consortium	AA052091
459 A	5 021HA1EQ	Mus musculus zipper protein kinase (ZPK) mRNA, complete cds {IMAGE:422079}	422079 IMAGE Consortium	W97442
460 A	7 021HA1EQ	IMAGE EST {IMAGE:422426}	422426 IMAGE Consortium	W97734
461 A	9 021HA1EQ	ESTs {IMAGE:421480}	421480 IMAGE Consortium	W99081
462 A	11 021HA1EQ	EST {IMAGE:423169}	423169 IMAGE Consortium	W97213
463 C	1 021HA1EQ	ESTs {IMAGE:421810}	421810 IMAGE Consortium	W91392
464 C	3 021HA1EQ	IMAGE EST {IMAGE:421885}	421885 IMAGE Consortium	W91416
465 C	5 021HA1EQ	Mus musculus cleavage and polyadenylation specificity factor (MCPSF) mRNA, complete cds {IMAGE:421579}	421579 IMAGE Consortium	W97292
466 C	7 021HA1EQ	Mouse mRNA for ras-GTPase-activating protein SH3-domain binding protein, complete cds {IMAGE:423772}	423772 IMAGE Consortium	W98559
467 C	9 021HA1EQ	IMAGE EST {IMAGE:422196}	422196 IMAGE Consortium	W97477
468 C	11 021HA1EQ	ESTs, Highly similar to RAS-RELATED PROTEIN RAC1 [Drosophila melanogaster] {IMAGE:422182}	422182 IMAGE Consortium	W97276
469 E	1 021HA1EQ	ESTs, Highly similar to PEPTIDYL-PROLYL CIS-TRANS ISOMERASE [Drosophila melanogaster] {IMAGE:420400}	420400 IMAGE Consortium	W89608
470 E	3 021HA1EQ	ESTs {IMAGE:422206}	422206 IMAGE Consortium	W97499
471 E	5 021HA1EQ	IMAGE EST {IMAGE:422452}	422452 IMAGE Consortium	W97741
472 E	7 021HA1EQ	Protein kinase C substrate 80K-H {IMAGE:421205}	421205 IMAGE Consortium	W91045
473 E	9 021HA1EQ	ESTs, Weakly similar to INTERFERON-INDUCED PROTEIN 6-16 PRECURSOR [Homo sapiens] {IMAGE:421476}	421476 IMAGE Consortium	W99140
474 E	11 021HA1EQ	ESTs {IMAGE:421361}	421361 IMAGE Consortium	W97033



475 G	1	021HA1EQ	ESTs {IMAGE:422477}	422477	IMAGE Consortium	W97427
476 G	3	021HA1EQ	ESTs, Weakly similar to C01F6.8 [C.elegans] {IMAGE:422667}	422667	IMAGE Consortium	W97837
477 G	5	021HA1EQ	ESTs {IMAGE:422946}	422946	IMAGE Consortium	W97929
478 G	7	021HA1EQ	IMAGE EST {IMAGE:423147}	423147	IMAGE Consortium	W97059
479 G	9	021HA1EQ	ESTs {IMAGE:423344}	423344	IMAGE Consortium	W98865
480 G	11	021HA1EQ	ESTs {IMAGE:421862}	421862	IMAGE Consortium	W91408
481 A	1	021OA1ER	IMAGE EST {IMAGE:420240}	420240	IMAGE Consortium	AA052715
482 A	3	021OA1ER	IMAGE EST {IMAGE:418564}	418564	IMAGE Consortium	W99034
483 A	5	021OA1ER	IMAGE EST {IMAGE:418527}	418527	IMAGE Consortium	W87066
484 A	7	021OA1ER	IMAGE EST {IMAGE:418531}	418531	IMAGE Consortium	W87120
485 A	9	021OA1ER	Myostatin {IMAGE:418993}	418993	IMAGE Consortium	AA052179
486 A	11	021OA1ER	Smoothened homolog (Drosophila) {IMAGE:419194}	419194	IMAGE Consortium	W87954
487 C	1	021OA1ER	ESTs, Moderately similar to SNAP45 subunit [H.sapiens] {IMAGE:409202}	409202	IMAGE Consortium	W87183
488 C	3	021OA1ER	IMAGE EST {IMAGE:420230}	420230	IMAGE Consortium	W89813
489 C	5	021OA1ER	Glycine transporter 1 {IMAGE:420070}	420070	IMAGE Consortium	W90900
490 C	7	021OA1ER	ESTs {IMAGE:420484}	420484	IMAGE Consortium	W90822
491 C	9	021OA1ER	IMAGE EST {IMAGE:420634}	420634	IMAGE Consortium	AA052147
492 C	11	021OA1ER	IMAGE EST {IMAGE:406426}	406426	IMAGE Consortium	W83426
493 E	1	021OA1ER	IMAGE EST {IMAGE:418732}	418732	IMAGE Consortium	W96980
494 E	3	021OA1ER	IMAGE EST {IMAGE:422204}	422204	IMAGE Consortium	W97498
495 E	5	021OA1ER	Mouse chromosome 6 BAC-284H12 (Research Genetics mouse BAC library) complete sequence {IMAGE:420853}	420853	IMAGE Consortium	W90961
496 E	7	021OA1ER	IMAGE EST {IMAGE:421443}	421443	IMAGE Consortium	W99064
497 E	9	021OA1ER	ESTs, Highly similar to PUTATIVE AMINOPEPTIDASE ZK353.6 IN CHROMOSOME III [Caenorhabditis elegans] {IMAGE:421148}	421148	IMAGE Consortium	W98973
498 E	11	021OA1ER	ESTs {IMAGE:421634}	421634	IMAGE Consortium	W99167
499 G	1	021OA1ER	Mus musculus glutathione synthetase type A1 mRNA, complete cds {IMAGE:421833}	421833	IMAGE Consortium	W91402
500 G	3	021OA1ER	IMAGE EST {IMAGE:423373}	423373	IMAGE Consortium	W98891
501 G	5	021OA1ER	IMAGE EST {IMAGE:423091}	423091	IMAGE Consortium	W97254
502 G	7	021OA1ER	IMAGE EST {IMAGE:422141}	422141	IMAGE Consortium	W97378
503 G	9	021OA1ER	Zinc finger protein 35 {IMAGE:422121}	422121	IMAGE Consortium	W97363
504 G	11	021OA1ER	Notch gene homolog 1, (Drosophila) {IMAGE:421911}	421911	IMAGE Consortium	W98998
505 A	1	021VA1ES	Mouse chromosome 6 BAC-284H12 (Research Genetics mouse BAC library) complete sequence {IMAGE:423368}	423368	IMAGE Consortium	W98889
506 A	3	021VA1ES	Mus musculus betaine homocysteine methyl transferase (BHMT) mRNA, complete cds {IMAGE:423156}	423156	IMAGE Consortium	W97296
507 A	5	021VA1ES	IMAGE EST {IMAGE:423065}	423065	IMAGE Consortium	W97245
508 A	7	021VA1ES	ESTs, Moderately similar to NADH-UBIQUINONE OXIDOREDUCTASE KFYI SUBUNIT PRECURSOR [Bos	423071	IMAGE Consortium	W97248

509 A	9 021VA1ES	taurus] {IMAGE:423071} ESTs, Highly similar to ALDEHYDE DEHYDROGENASE, MITOCHONDRIAL X PRECURSOR [Homo sapiens] {IMAGE:423605}	423605 IMAGE Consortium	W98919
510 A	11 021VA1ES	IMAGE EST {IMAGE:422035}	422035 IMAGE Consortium	W96841
511 C	1 021VA1ES	Cytochrome P450, 2e1, ethanol inducible {IMAGE:423749}	423749 IMAGE Consortium	W98556
512 C	3 021VA1ES	ESTs, Moderately similar to ENDOTHELIAL ACTIN- BINDING PROTEIN [Homo sapiens] {IMAGE:420569}	420569 IMAGE Consortium	W90909
513 C	5 021VA1ES	IMAGE EST {IMAGE:422231}	422231 IMAGE Consortium	W97572
514 C	7 021VA1ES	ESTs {IMAGE:421155}	421155 IMAGE Consortium	W91036
515 C	9 021VA1ES	Mus musculus viral envelope like protein (G7e) gene, complete cds {IMAGE:422669}	422669 IMAGE Consortium	W97838
516 C	11 021VA1ES	ESTs {IMAGE:421549}	421549 IMAGE Consortium	W97161
517 E	1 021VA1ES	ESTs, Highly similar to unknown [H.sapiens] {IMAGE:421797}	421797 IMAGE Consortium	W99114
518 E	3 021VA1ES	Mus musculus GCN5L1 gene, exon 1 and joined CDS {IMAGE:425639}	425639 IMAGE Consortium	AA000036
519 E	5 021VA1ES	Complement component 1, q subcomponent, c polypeptide {IMAGE:426010}	426010 IMAGE Consortium	AA000461
520 E	7 021VA1ES	ESTs, Highly similar to KINESIN-II 85 KD SUBUNIT [Strongylocentrotus purpuratus] {IMAGE:424705}	424705 IMAGE Consortium	AA000842
521 E	9 021VA1ES	IMAGE EST {IMAGE:424796}	424796 IMAGE Consortium	AA000581
522 E	11 021VA1ES	ESTs {IMAGE:425281}	425281 IMAGE Consortium	W99880
523 G	1 021VA1ES	ESTs {IMAGE:426199}	426199 IMAGE Consortium	AA002947
524 G	3 021VA1ES	IMAGE EST {IMAGE:424964}	424964 IMAGE Consortium	W99838
525 G	5 021VA1ES	Chaperonin subunit 3 (gamma) {IMAGE:426239}	426239 IMAGE Consortium	AA002864
526 G	7 021VA1ES	ESTs {IMAGE:425106}	425106 IMAGE Consortium	W99938
527 G	9 021VA1ES	ESTs {IMAGE:425425}	425425 IMAGE Consortium	AA000101
528 G	11 021VA1ES	ESTs {IMAGE:427091}	427091 IMAGE Consortium	AA003378
529 A	1 0212A1ET	IMAGE EST {IMAGE:425998}	425998 IMAGE Consortium	AA000862
530 A	3 0212A1ET	IMAGE EST {IMAGE:424629}	424629 IMAGE Consortium	AA000641
531 A	5 0212A1ET	ESTs {IMAGE:426633}	426633 IMAGE Consortium	AA002573
532 A	7 0212A1ET	IMAGE EST {IMAGE:425593}	425593 IMAGE Consortium	AA000347
533 A	9 0212A1ET	ESTs {IMAGE:427218}	427218 IMAGE Consortium	AA003081
534 A	11 0212A1ET	ESTs {IMAGE:423984}	423984 IMAGE Consortium	W91480
535 C	1 0212A1ET	Prostaglandin E receptor EP1 subtype {IMAGE:425156}	425156 IMAGE Consortium	W99842
536 C	3 0212A1ET	IMAGE EST {IMAGE:425450}	425450 IMAGE Consortium	AA000180
537 C	5 0212A1ET	IMAGE EST {IMAGE:427235}	427235 IMAGE Consortium	AA003040
538 C	7 0212A1ET	IMAGE EST {IMAGE:427075}	427075 IMAGE Consortium	AA003159
539 C	9 0212A1ET	ESTs, Highly similar to TAR DNA-binding protein-43 [H.sapiens] {IMAGE:427346}	427346 IMAGE Consortium	AA003149
540 C	11 0212A1ET	IMAGE EST {IMAGE:425785}	425785 IMAGE Consortium	AA000678
541 E	1 0212A1ET	IMAGE EST {IMAGE:426989}	426989 IMAGE Consortium	AA003421
542 E	3 0212A1ET	ESTs, Highly similar to PROBABLE UBIQUITIN CARBOXYL-	425458 IMAGE Consortium	AA000038

		TERMINAL HYDROLASE [Mus musculus] {IMAGE:425458}		
543 E	5 0212A1ET	IMAGE EST {IMAGE:425583}	425583 IMAGE Consortium	AA000022
544 E	7 0212A1ET	DNA polymerase delta 1, catalytic domain {IMAGE:424021}	424021 IMAGE Consortium	W91387
545 E	9 0212A1ET	IMAGE EST {IMAGE:425936}	425936 IMAGE Consortium	AA000816
546 E	11 0212A1ET	Cytochrome P450, 2e1, ethanol inducible {IMAGE:426043}	426043 IMAGE Consortium	AA002904
547 G	1 0212A1ET	ESTs, Highly similar to REPLICATION PROTEIN A 14 KD SUBUNIT [Homo sapiens] {IMAGE:425703}	425703 IMAGE Consortium	AA000318
548 G	3 0212A1ET	EST {IMAGE:425493}	425493 IMAGE Consortium	AA000266
549 G	5 0212A1ET	Mus musculus transcription factor PBX3b (PBX3b) mRNA, complete cds {IMAGE:425881}	425881 IMAGE Consortium	AA000829
550 G	7 0212A1ET	IMAGE EST {IMAGE:425745}	425745 IMAGE Consortium	AA000732
551 G	9 0212A1ET	IMAGE EST {IMAGE:426086}	426086 IMAGE Consortium	AA002851
552 G	11 0212A1ET	Protein kinase C, theta {IMAGE:424857}	424857 IMAGE Consortium	W98195
553 A	1 0219A1EU	IMAGE EST {IMAGE:439001}	439001 IMAGE Consortium	AA008242
554 A	3 0219A1EU	IMAGE EST {IMAGE:439419}	439419 IMAGE Consortium	AA004070
555 A	5 0219A1EU	IMAGE EST {IMAGE:439362}	439362 IMAGE Consortium	AA004120
556 A	7 0219A1EU	IMAGE EST {IMAGE:439684}	439684 IMAGE Consortium	AA014431
557 A	9 0219A1EU	IMAGE EST {IMAGE:438506}	438506 IMAGE Consortium	AA050778
558 A	11 0219A1EU	ESTs {IMAGE:438463}	438463 IMAGE Consortium	AA008251
559 C	1 0219A1EU	IMAGE EST {IMAGE:438016}	438016 IMAGE Consortium	AA007816
560 C	3 0219A1EU	ESTs, Highly similar to NUCLEOLAR PROTEIN NAP57 [R.norvegicus] {IMAGE:438128}	438128 IMAGE Consortium	AA008051
561 C	5 0219A1EU	IMAGE EST {IMAGE:438254}	438254 IMAGE Consortium	AA008279
562 C	7 0219A1EU	IMAGE EST {IMAGE:439927}	439927 IMAGE Consortium	AA008735
563 C	9 0219A1EU	IMAGE EST {IMAGE:439596}	439596 IMAGE Consortium	AA008664
564 C	11 0219A1EU	ESTs {IMAGE:438735}	438735 IMAGE Consortium	AA051129
565 E	1 0219A1EU	IMAGE EST {IMAGE:438296}	438296 IMAGE Consortium	AA008288
566 E	3 0219A1EU	IMAGE EST {IMAGE:439829}	439829 IMAGE Consortium	AA008529
567 E	5 0219A1EU	ESTs, Highly similar to ARGINYL-TRNA SYNTHETASE [Cricetulus longicaudatus] {IMAGE:439640}	439640 IMAGE Consortium	AA008699
568 E	7 0219A1EU	ESTs {IMAGE:427289}	427289 IMAGE Consortium	AA003258
569 E	9 0219A1EU	ESTs, Weakly similar to Enx-2 [M.musculus] {IMAGE:427559}	427559 IMAGE Consortium	AA002473
570 E	11 0219A1EU	ESTs {IMAGE:436999}	436999 IMAGE Consortium	AA002783
571 G	1 0219A1EU	Catenin src {IMAGE:439784}	439784 IMAGE Consortium	AA008582
572 G	3 0219A1EU	ESTs {IMAGE:427594}	427594 IMAGE Consortium	AA002543
573 G	5 0219A1EU	ESTs {IMAGE:438616}	438616 IMAGE Consortium	AA051009
574 G	7 0219A1EU	ESTs, Moderately similar to highly charged protein [H.sapiens] {IMAGE:437167}	437167 IMAGE Consortium	AA003443
575 G	9 0219A1EU	ESTs {IMAGE:437246}	437246 IMAGE Consortium	AA003541
576 G	11 0219A1EU	Cadherin 3 {IMAGE:437297}	437297 IMAGE Consortium	AA003679
577 A	1 021GA1EV	IMAGE EST {IMAGE:424192}	424192 IMAGE Consortium	W98228
578 A	3 021GA1EV	Mus musculus sphingosine kinase (SPHK1b) mRNA, complete cds {IMAGE:425961}	425961 IMAGE Consortium	AA000819

579 A	5 021GA1EV	IMAGE EST {IMAGE:426240}	426240 IMAGE Consortium	AA002801
580 A	7 021GA1EV	FMS-like tyrosine kinase 3 ligand {IMAGE:426471}	426471 IMAGE Consortium	AA002314
581 A	9 021GA1EV	Mus musculus mRNA for collagen a1(V), complete cds {IMAGE:425344}	425344 IMAGE Consortium	W99856
582 A	11 021GA1EV	IMAGE EST {IMAGE:426815}	426815 IMAGE Consortium	AA003199
583 C	1 021GA1EV	ESTs {IMAGE:426380}	426380 IMAGE Consortium	AA002887
584 C	3 021GA1EV	IMAGE EST {IMAGE:426570}	426570 IMAGE Consortium	AA002446
585 C	5 021GA1EV	IMAGE EST {IMAGE:427152}	427152 IMAGE Consortium	AA003356
586 C	7 021GA1EV	IMAGE EST {IMAGE:426967}	426967 IMAGE Consortium	AA003273
587 C	9 021GA1EV	ESTs, Highly similar to ZINC FINGER PROTEIN 7 [Homo sapiens] {IMAGE:427349}	427349 IMAGE Consortium	AA003172
588 C	11 021GA1EV	IMAGE EST {IMAGE:424402}	424402 IMAGE Consortium	W98502
589 E	1 021GA1EV	Myeloblastosis oncogene-like 2 {IMAGE:427239}	427239 IMAGE Consortium	AA003042
590 E	3 021GA1EV	IMAGE EST {IMAGE:425561}	425561 IMAGE Consortium	AA000229
591 E	5 021GA1EV	ESTs {IMAGE:424563}	424563 IMAGE Consortium	W98594
592 E	7 021GA1EV	Cell adhesion kinase {IMAGE:424307}	424307 IMAGE Consortium	W98395
593 E	9 021GA1EV	ESTs {IMAGE:424632}	424632 IMAGE Consortium	AA000501
594 E	11 021GA1EV	ESTs {IMAGE:426172}	426172 IMAGE Consortium	AA002994
595 G	1 021GA1EV	IMAGE EST {IMAGE:424348}	424348 IMAGE Consortium	W98625
596 G	3 021GA1EV	IMAGE EST {IMAGE:424403}	424403 IMAGE Consortium	W98506
597 G	5 021GA1EV	ESTs {IMAGE:426069}	426069 IMAGE Consortium	AA002913
598 G	7 021GA1EV	ESTs, Weakly similar to HYPOTHETICAL 36.7 KD PROTEIN C2F7.02C IN CHROMOSOME I [Schizosaccharomyces pombe] {IMAGE:426233}	426233 IMAGE Consortium	AA002951
599 G	9 021GA1EV	Neurofilament, light polypeptide {IMAGE:425315}	425315 IMAGE Consortium	W99951
600 G	11 021GA1EV	Kinesin heavy chain member 1B {IMAGE:426763}	426763 IMAGE Consortium	AA002775
601 A	1 021NA1EW	Mus musculus NSD1 protein mRNA, complete cds {IMAGE:438694}	438694 IMAGE Consortium	AA051096
602 A	3 021NA1EW	Smoothened homolog (Drosophila) {IMAGE:439010}	439010 IMAGE Consortium	AA008222
603 A	5 021NA1EW	IMAGE EST {IMAGE:438883}	438883 IMAGE Consortium	AA008191
604 A	7 021NA1EW	IMAGE EST {IMAGE:439273}	439273 IMAGE Consortium	AA003981
605 A	9 021NA1EW	T-COMPLEX PROTEIN 1, BETA SUBUNIT {IMAGE:439254}	439254 IMAGE Consortium	AA003927
606 A	11 021NA1EW	ESTs, Highly similar to HETEROGENEOUS NUCLEAR RIBONUCLEOPROTEIN 27C [Drosophila melanogaster] {IMAGE:437938}	437938 IMAGE Consortium	AA007804
607 C	1 021NA1EW	IMAGE EST {IMAGE:439130}	439130 IMAGE Consortium	AA003836
608 C	3 021NA1EW	IMAGE EST {IMAGE:438908}	438908 IMAGE Consortium	AA007975
609 C	5 021NA1EW	IMAGE EST {IMAGE:439297}	439297 IMAGE Consortium	AA003995
610 C	7 021NA1EW	IMAGE EST {IMAGE:439446}	439446 IMAGE Consortium	AA004080
611 C	9 021NA1EW	IMAGE EST {IMAGE:437699}	437699 IMAGE Consortium	AA007775
612 C	11 021NA1EW	IMAGE EST {IMAGE:437685}	437685 IMAGE Consortium	AA007828
613 E	1 021NA1EW	IMAGE EST {IMAGE:437473}	437473 IMAGE Consortium	AA003746
614 E	3 021NA1EW	ESTs, Highly similar to kinesin-related protein 2	437761 IMAGE Consortium	AA007891

615 E	5 021NA1EW	[R.norvegicus] {IMAGE:437761} ESTs, Highly similar to protein inhibitor of neuronal nitric oxide synthase [R.norvegicus] {IMAGE:438208}	438208 IMAGE Consortium	AA008134
616 E	7 021NA1EW	IMAGE EST {IMAGE:439713}	439713 IMAGE Consortium	AA014455
617 E	9 021NA1EW	IMAGE EST {IMAGE:439765}	439765 IMAGE Consortium	AA008580
618 E	11 021NA1EW	IMAGE EST {IMAGE:438778}	438778 IMAGE Consortium	AA051135
619 G	1 021NA1EW	IMAGE EST {IMAGE:438383}	438383 IMAGE Consortium	AA008024
620 G	3 021NA1EW	Intercellular adhesion molecule 2 {IMAGE:438581}	438581 IMAGE Consortium	AA050972
621 G	5 021NA1EW	ESTs {IMAGE:427333}	427333 IMAGE Consortium	AA003226
622 G	7 021NA1EW	IMAGE EST {IMAGE:437162}	437162 IMAGE Consortium	AA003440
623 G	9 021NA1EW	ESTs {IMAGE:437172}	437172 IMAGE Consortium	AA003528
624 G	11 021NA1EW	IMAGE EST {IMAGE:437110}	437110 IMAGE Consortium	AA003405
625 A	1 021UA1EX	IMAGE EST {IMAGE:437234}	437234 IMAGE Consortium	AA003454
626 A	3 021UA1EX	ESTs {IMAGE:439132}	439132 IMAGE Consortium	AA003837
627 A	5 021UA1EX	Microtubule-associated protein 4 {IMAGE:437523}	437523 IMAGE Consortium	AA003769
628 A	7 021UA1EX	ESTs {IMAGE:439323}	439323 IMAGE Consortium	AA004021
629 A	9 021UA1EX	IMAGE EST {IMAGE:439399}	439399 IMAGE Consortium	AA004013
630 A	11 021UA1EX	ESTs, Weakly similar to F29B9.4 [C.elegans] {IMAGE:437679}	437679 IMAGE Consortium	AA007773
631 C	1 021UA1EX	IMAGE EST {IMAGE:439447}	439447 IMAGE Consortium	AA004096
632 C	3 021UA1EX	IMAGE EST {IMAGE:437727}	437727 IMAGE Consortium	AA007752
633 C	5 021UA1EX	ESTs {IMAGE:439222}	439222 IMAGE Consortium	AA003955
634 C	7 021UA1EX	IMAGE EST {IMAGE:439803}	439803 IMAGE Consortium	AA008521
635 C	9 021UA1EX	ESTs {IMAGE:439735}	439735 IMAGE Consortium	AA014384
636 C	11 021UA1EX	Carbon catabolite repression 4 homolog (S. cerevisiae) {IMAGE:438774}	438774 IMAGE Consortium	AA051133
637 E	1 021UA1EX	IMAGE EST {IMAGE:439751}	439751 IMAGE Consortium	AA014587
638 E	3 021UA1EX	ESTs, Highly similar to ATP-DEPENDENT CLP PROTEASE ATP-BINDING SUBUNIT CLPX [Haemophilus influenzae] {IMAGE:441677}	441677 IMAGE Consortium	AA013832
639 E	5 021UA1EX	Mus musculus golgin-245 (olp-1) mRNA, complete cds {IMAGE:440038}	440038 IMAGE Consortium	AA009086
640 E	7 021UA1EX	IMAGE EST {IMAGE:442012}	442012 IMAGE Consortium	AA014445
641 E	9 021UA1EX	IMAGE EST {IMAGE:442021}	442021 IMAGE Consortium	AA014678
642 E	11 021UA1EX	IMAGE EST {IMAGE:441216}	441216 IMAGE Consortium	AA011829
643 G	1 021UA1EX	IMAGE EST {IMAGE:441788}	441788 IMAGE Consortium	AA013851
644 G	3 021UA1EX	IMAGE EST {IMAGE:440978}	440978 IMAGE Consortium	AA008911
645 G	5 021UA1EX	IMAGE EST {IMAGE:439744}	439744 IMAGE Consortium	AA014534
646 G	7 021UA1EX	Mouse beta-D-galactosidase fusion protein mRNA, complete cds {IMAGE:439652}	439652 IMAGE Consortium	AA008495
647 G	9 021UA1EX	IMAGE EST {IMAGE:441769}	441769 IMAGE Consortium	AA014004
648 G	11 021UA1EX	Mus musculus mRNA for mrg1 protein {IMAGE:442189}	442189 IMAGE Consortium	AA013772
649 A	1 021UA1EX	ESTs, Moderately similar to cDNA EST EMBL:Z14908 comes	442944 IMAGE Consortium	AA014165

		from this gene [C.elegans] {IMAGE:442944}		
650 A	3 0211A1EY	Glycoprotein galactosyltransferase beta 1, 4 {IMAGE:442858}	442858 IMAGE Consortium	AA014772
651 A	5 0211A1EY	ESTs, Moderately similar to KERATIN, TYPE II CYTOSKELETAL 5 [Homo sapiens] {IMAGE:445061}	445061 IMAGE Consortium	AA016810
652 A	7 0211A1EY	ESTs {IMAGE:444055}	444055 IMAGE Consortium	AA014901
653 A	9 0211A1EY	IMAGE EST {IMAGE:445707}	445707 IMAGE Consortium	AA017940
654 A	11 0211A1EY	ESTs, Highly similar to CATION-TRANSPORTING ATPASE PACL [Synechococcus pcc7942] {IMAGE:442484}	442484 IMAGE Consortium	AA015291
655 C	1 0211A1EY	Fos-like antigen 2 {IMAGE:445817}	445817 IMAGE Consortium	AA020223
656 C	3 0211A1EY	IMAGE EST {IMAGE:442639}	442639 IMAGE Consortium	AA015142
657 C	5 0211A1EY	Amyloid beta (A4) precursor-like protein 2 {IMAGE:444846}	444846 IMAGE Consortium	AA016732
658 C	7 0211A1EY	Mus musculus TFIIH transcription/DNA repair factor p52 subunit mRNA, complete cds {IMAGE:444718}	444718 IMAGE Consortium	AA016373
659 C	9 0211A1EY	IMAGE EST {IMAGE:445197}	445197 IMAGE Consortium	AA016827
660 C	11 0211A1EY	IMAGE EST {IMAGE:443638}	443638 IMAGE Consortium	AA013742
661 E	1 0211A1EY	IMAGE EST {IMAGE:443806}	443806 IMAGE Consortium	AA014846
662 E	3 0211A1EY	PROTEIN KINASE C, ZETA TYPE {IMAGE:442856}	442856 IMAGE Consortium	AA014771
663 E	5 0211A1EY	Glutamate dehydrogenase {IMAGE:442843}	442843 IMAGE Consortium	AA014196
664 E	7 0211A1EY	ESTs {IMAGE:444954}	444954 IMAGE Consortium	AA016761
665 E	9 0211A1EY	IMAGE EST {IMAGE:445426}	445426 IMAGE Consortium	AA017991
666 E	11 0211A1EY	IMAGE EST {IMAGE:445908}	445908 IMAGE Consortium	AA020034
667 G	1 0211A1EY	Mouse mRNA for uridine phosphorylase, complete cds {IMAGE:445862}	445862 IMAGE Consortium	AA020155
668 G	3 0211A1EY	IMAGE EST {IMAGE:442365}	442365 IMAGE Consortium	AA015532
669 G	5 0211A1EY	EST {IMAGE:442833}	442833 IMAGE Consortium	AA014745
670 G	7 0211A1EY	IMAGE EST {IMAGE:445560}	445560 IMAGE Consortium	AA017895
671 G	9 0211A1EY	IMAGE EST {IMAGE:443815}	443815 IMAGE Consortium	AA014091
672 G	11 0211A1EY	IMAGE EST {IMAGE:444010}	444010 IMAGE Consortium	AA014885
673 A	1 0218A1EZ	Mus musculus DNA polymerase epsilon small subunit mRNA, partial cds {IMAGE:439792}	439792 IMAGE Consortium	AA008627
674 A	3 0218A1EZ	ESTs, Moderately similar to unknown [H.sapiens] {IMAGE:440584}	440584 IMAGE Consortium	AA014820
675 A	5 0218A1EZ	ESTs, Highly similar to GASTRIN-RELEASING PEPTIDE PRECURSOR [Rattus norvegicus] {IMAGE:440937}	440937 IMAGE Consortium	AA008951
676 A	7 0218A1EZ	Hormone receptor {IMAGE:439773}	439773 IMAGE Consortium	AA008625
677 A	9 0218A1EZ	ESTs {IMAGE:440053}	440053 IMAGE Consortium	AA016507
678 A	11 0218A1EZ	Signal transducer and activator of transcription 3 {IMAGE:440571}	440571 IMAGE Consortium	AA014643
679 C	1 0218A1EZ	ESTs {IMAGE:440057}	440057 IMAGE Consortium	AA016594
680 C	3 0218A1EZ	IMAGE EST {IMAGE:440418}	440418 IMAGE Consortium	AA011778
681 C	5 0218A1EZ	ESTs {IMAGE:440941}	440941 IMAGE Consortium	AA008952
682 C	7 0218A1EZ	IMAGE EST {IMAGE:442527}	442527 IMAGE Consortium	AA015136
683 C	9 0218A1EZ	T-complex-associated testis expressed 1 {IMAGE:439654}	439654 IMAGE Consortium	AA008496

684 C	11 0218A1EZ	IMAGE EST {IMAGE:440121}	440121 IMAGE Consortium	AA016547
685 E	1 0218A1EZ	IMAGE EST {IMAGE:440938}	440938 IMAGE Consortium	AA008941
686 E	3 0218A1EZ	ESTs, Moderately similar to HYPOTHETICAL 97.6 KD PROTEIN IN SHP1-SEC17 INTERGENIC REGION [Saccharomyces cerevisiae] {IMAGE:441101}	441101 IMAGE Consortium	AA008976
687 E	5 0218A1EZ	IMAGE EST {IMAGE:442294}	442294 IMAGE Consortium	AA013725
688 E	7 0218A1EZ	IMAGE EST {IMAGE:442744}	442744 IMAGE Consortium	AA036470
689 E	9 0218A1EZ	IMAGE EST {IMAGE:443023}	443023 IMAGE Consortium	AA014227
690 E	11 0218A1EZ	IMAGE EST {IMAGE:443580}	443580 IMAGE Consortium	AA013553
691 G	1 0218A1EZ	ESTs {IMAGE:445864}	445864 IMAGE Consortium	AA020233
692 G	3 0218A1EZ	ESTs, Highly similar to hypothetical protein [M.musculus] {IMAGE:442415}	442415 IMAGE Consortium	AA015542
693 G	5 0218A1EZ	ESTs, Highly similar to ALPHA-2-MACROGLOBULIN PRECURSOR [Homo sapiens] {IMAGE:444715}	444715 IMAGE Consortium	AA016374
694 G	7 0218A1EZ	IMAGE EST {IMAGE:443395}	443395 IMAGE Consortium	AA015048
695 G	9 0218A1EZ	IMAGE EST {IMAGE:443596}	443596 IMAGE Consortium	AA015004
696 G	11 0218A1EZ	Mus musculus Pur-alpha mRNA, complete cds {IMAGE:445562}	445562 IMAGE Consortium	AA017896
697 A	1 0218A1F0	ESTs {IMAGE:455545}	455545 IMAGE Consortium	AA020557
698 A	3 0218A1F0	ESTs {IMAGE:457432}	457432 IMAGE Consortium	AA021844
699 A	5 0218A1F0	Suppressor of Ty 4 homolog (S. cerevisiae) {IMAGE:457853}	457853 IMAGE Consortium	AA021916
700 A	7 0218A1F0	IMAGE EST {IMAGE:456481}	456481 IMAGE Consortium	AA023385
701 A	9 0218A1F0	ESTs {IMAGE:445725}	445725 IMAGE Consortium	AA020017
702 A	11 0218A1F0	ESTs, Highly similar to ESTROGEN RECEPTOR [Gallus gallus] {IMAGE:457235}	457235 IMAGE Consortium	AA023625
703 C	1 0218A1F0	IMAGE EST {IMAGE:456492}	456492 IMAGE Consortium	AA023427
704 C	3 0218A1F0	RAB geranylgeranyl transferase, b subunit {IMAGE:455483}	455483 IMAGE Consortium	AA020526
705 C	5 0218A1F0	ESTs {IMAGE:456208}	456208 IMAGE Consortium	AA023256
706 C	7 0218A1F0	IMAGE EST {IMAGE:456493}	456493 IMAGE Consortium	AA023390
707 C	9 0218A1F0	ESTs {IMAGE:445847}	445847 IMAGE Consortium	AA020232
708 C	11 0218A1F0	IMAGE EST {IMAGE:457253}	457253 IMAGE Consortium	AA023595
709 E	1 0218A1F0	ESTs {IMAGE:457055}	457055 IMAGE Consortium	AA023720
710 E	3 0218A1F0	ESTs {IMAGE:456342}	456342 IMAGE Consortium	AA023338
711 E	5 0218A1F0	IMAGE EST {IMAGE:455408}	455408 IMAGE Consortium	AA020458
712 E	7 0218A1F0	IMAGE EST {IMAGE:457434}	457434 IMAGE Consortium	AA021845
713 E	9 0218A1F0	IMAGE EST {IMAGE:455800}	455800 IMAGE Consortium	AA023801
714 E	11 0218A1F0	ESTs {IMAGE:456295}	456295 IMAGE Consortium	AA023306
715 G	1 0218A1F0	ESTs {IMAGE:457071}	457071 IMAGE Consortium	AA023656
716 G	3 0218A1F0	ESTs {IMAGE:455401}	455401 IMAGE Consortium	AA020462
717 G	5 0218A1F0	IMAGE EST {IMAGE:457567}	457567 IMAGE Consortium	AA021943
718 G	7 0218A1F0	EST {IMAGE:455347}	455347 IMAGE Consortium	AA020470
719 G	9 0218A1F0	IMAGE EST {IMAGE:455930}	455930 IMAGE Consortium	AA023072
720 G	11 0218A1F0	Histocompatibility 2, K region {IMAGE:458067}	458067 IMAGE Consortium	AA030668

721 A	1	021FA1F1	ESTs, Weakly similar to TRANSCRIPTION INITIATION FACTOR TFIID 150 KD SUBUNIT [D.melanogaster] {IMAGE:457963}	457963	IMAGE Consortium	AA028778
722 A	3	021FA1F1	IMAGE EST {IMAGE:458457}	458457	IMAGE Consortium	AA024375
723 A	5	021FA1F1	ESTs {IMAGE:459137}	459137	IMAGE Consortium	AA024030
724 A	7	021FA1F1	ESTs {IMAGE:464099}	464099	IMAGE Consortium	AA028306
725 A	9	021FA1F1	Mus musculus mRNA for vesicular GABA and glycine transporter {IMAGE:463941}	463941	IMAGE Consortium	AA028280
726 A	11	021FA1F1	IMAGE EST {IMAGE:464209}	464209	IMAGE Consortium	AA028406
727 C	1	021FA1F1	IMAGE EST {IMAGE:458926}	458926	IMAGE Consortium	AA024251
728 C	3	021FA1F1	Serine protease inhibitor 4 {IMAGE:464497}	464497	IMAGE Consortium	AA033050
729 C	5	021FA1F1	IMAGE EST {IMAGE:459902}	459902	IMAGE Consortium	AA027589
730 C	7	021FA1F1	ESTs, Weakly similar to protein co-factor [M.musculus] {IMAGE:464250}	464250	IMAGE Consortium	AA033055
731 C	9	021FA1F1	ESTs {IMAGE:463334}	463334	IMAGE Consortium	AA027707
732 C	11	021FA1F1	ESTs {IMAGE:463002}	463002	IMAGE Consortium	AA030573
733 E	1	021FA1F1	ESTs {IMAGE:459639}	459639	IMAGE Consortium	AA027549
734 E	3	021FA1F1	IMAGE EST {IMAGE:464415}	464415	IMAGE Consortium	AA030512
735 E	5	021FA1F1	IMAGE EST {IMAGE:464442}	464442	IMAGE Consortium	AA030427
736 E	7	021FA1F1	IMAGE EST {IMAGE:464784}	464784	IMAGE Consortium	AA030098
737 E	9	021FA1F1	IMAGE EST {IMAGE:464863}	464863	IMAGE Consortium	AA030422
738 E	11	021FA1F1	ESTs {IMAGE:463656}	463656	IMAGE Consortium	AA028716
739 G	1	021FA1F1	IMAGE EST {IMAGE:463314}	463314	IMAGE Consortium	AA028670
740 G	3	021FA1F1	IMAGE EST {IMAGE:463592}	463592	IMAGE Consortium	AA030781
741 G	5	021FA1F1	Mus musculus BS4 peptide mRNA, complete cds {IMAGE:459091}	459091	IMAGE Consortium	AA024028
742 G	7	021FA1F1	IMAGE EST {IMAGE:459511}	459511	IMAGE Consortium	AA027452
743 G	9	021FA1F1	IMAGE EST {IMAGE:464498}	464498	IMAGE Consortium	AA033029
744 G	11	021FA1F1	Mus musculus thioredoxin-related protein mRNA, complete cds {IMAGE:459904}	459904	IMAGE Consortium	AA030348
745 A	1	021MA1F2	IMAGE EST {IMAGE:463889}	463889	IMAGE Consortium	AA028229
746 A	3	021MA1F2	IMAGE EST {IMAGE:459284}	459284	IMAGE Consortium	AA030294
747 A	5	021MA1F2	IMAGE EST {IMAGE:464381}	464381	IMAGE Consortium	AA030385
748 A	7	021MA1F2	ESTs {IMAGE:462987}	462987	IMAGE Consortium	AA027643
749 A	9	021MA1F2	IMAGE EST {IMAGE:463255}	463255	IMAGE Consortium	AA030379
750 A	11	021MA1F2	IMAGE EST {IMAGE:463046}	463046	IMAGE Consortium	AA027653
751 C	1	021MA1F2	Mus musculus mRNA for CtBP1 protein {IMAGE:467484}	467484	IMAGE Consortium	AA036604
752 C	3	021MA1F2	Mouse RNA helicase and RNA-dependent ATPase from the DEAD box family mRNA, complete cds {IMAGE:466224}	466224	IMAGE Consortium	AA033377
753 C	5	021MA1F2	IMAGE EST {IMAGE:466280}	466280	IMAGE Consortium	AA033362
754 C	7	021MA1F2	IMAGE EST {IMAGE:466190}	466190	IMAGE Consortium	AA033485
755 C	9	021MA1F2	ESTs {IMAGE:468635}	468635	IMAGE Consortium	AA014578
756 C	11	021MA1F2	ESTs, Highly similar to RABPHILIN-3A [Rattus norvegicus]	466503	IMAGE Consortium	AA030884



		{IMAGE:466503}		
757 E	1	021MA1F2	IMAGE EST {IMAGE:468019}	468019 IMAGE Consortium AA015563
758 E	3	021MA1F2	IMAGE EST {IMAGE:466391}	466391 IMAGE Consortium AA034761
759 E	5	021MA1F2	ESTs, Highly similar to PROFILIN II [Homo sapiens] {IMAGE:464995}	464995 IMAGE Consortium AA032658
760 E	7	021MA1F2	MAD homolog 4 (Drosophila) {IMAGE:466551}	466551 IMAGE Consortium AA030901
761 E	9	021MA1F2	IMAGE EST {IMAGE:466678}	466678 IMAGE Consortium AA031159
762 E	11	021MA1F2	Myeloperoxidase {IMAGE:465816}	465816 IMAGE Consortium AA033466
763 G	1	021MA1F2	ESTs, Highly similar to transcription factor TFIID chain TAFII28 [H.sapiens] {IMAGE:467027}	467027 IMAGE Consortium AA034616
764 G	3	021MA1F2	IMAGE EST {IMAGE:465984}	465984 IMAGE Consortium AA031051
765 G	5	021MA1F2	IMAGE EST {IMAGE:465679}	465679 IMAGE Consortium AA032882
766 G	7	021MA1F2	IMAGE EST {IMAGE:467559}	467559 IMAGE Consortium AA036463
767 G	9	021MA1F2	EST {IMAGE:466225}	466225 IMAGE Consortium AA033381
768 G	11	021MA1F2	ESTs, Weakly similar to F42F12.4 [C.elegans] {IMAGE:468356}	468356 IMAGE Consortium AA014552
769 A	1	021TA1F3	Insulin-like growth factor 2 receptor {IMAGE:464200}	464200 IMAGE Consortium AA028475
770 A	3	021TA1F3	IMAGE EST {IMAGE:462995}	462995 IMAGE Consortium AA027647
771 A	5	021TA1F3	IMAGE EST {IMAGE:464530}	464530 IMAGE Consortium AA030081
772 A	7	021TA1F3	IMAGE EST {IMAGE:463586}	463586 IMAGE Consortium AA028677
773 A	9	021TA1F3	ESTs, Highly similar to COATOMER BETA SUBUNIT [Rattus norvegicus] {IMAGE:463386}	463386 IMAGE Consortium AA030639
774 A	11	021TA1F3	Mus musculus mRNA for sFlt-1, complete cds {IMAGE:458943}	458943 IMAGE Consortium AA024210
775 C	1	021TA1F3	ESTs, Highly similar to BETA CRYSTALLIN B1 [Rattus norvegicus] {IMAGE:464120}	464120 IMAGE Consortium AA028309
776 C	3	021TA1F3	IMAGE EST {IMAGE:459858}	459858 IMAGE Consortium AA030324
777 C	5	021TA1F3	ESTs {IMAGE:464548}	464548 IMAGE Consortium AA030497
778 C	7	021TA1F3	IMAGE EST {IMAGE:464650}	464650 IMAGE Consortium AA030271
779 C	9	021TA1F3	IMAGE EST {IMAGE:463477}	463477 IMAGE Consortium AA028628
780 C	11	021TA1F3	ESTs {IMAGE:464936}	464936 IMAGE Consortium AA031241
781 E	1	021TA1F3	IMAGE EST {IMAGE:463020}	463020 IMAGE Consortium AA027467
782 E	3	021TA1F3	ESTs {IMAGE:463621}	463621 IMAGE Consortium AA030810
783 E	5	021TA1F3	IMAGE EST {IMAGE:463651}	463651 IMAGE Consortium AA027730
784 E	7	021TA1F3	ESTs {IMAGE:458992}	458992 IMAGE Consortium AA024158
785 E	9	021TA1F3	Mus musculus SH3-containing protein SH3P2 mRNA, partial cds {IMAGE:458781}	458781 IMAGE Consortium AA024088
786 E	11	021TA1F3	ESTs, Moderately similar to TUBULIN--TYROSINE LIGASE [Bos taurus] {IMAGE:459276}	459276 IMAGE Consortium AA030256
787 G	1	021TA1F3	ESTs, Weakly similar to ZINC FINGER PROTEIN HF.12 [Homo sapiens] {IMAGE:463723}	463723 IMAGE Consortium AA028784
788 G	3	021TA1F3	Microtubule-associated protein tau {IMAGE:463954}	463954 IMAGE Consortium AA028410
789 G	5	021TA1F3	IMAGE EST {IMAGE:459493}	459493 IMAGE Consortium AA027444

790 G	7 021TA1F3	Early lymphoid specific transcription factor {IMAGE:459676}	459676 IMAGE Consortium	AA027561
791 G	9 021TA1F3	IMAGE EST {IMAGE:464504}	464504 IMAGE Consortium	AA033097
792 G	11 021TA1F3	IMAGE EST {IMAGE:463346}	463346 IMAGE Consortium	AA027699
793 A	1 0210A1F4	Bone morphogenetic protein 1 {IMAGE:466306}	466306 IMAGE Consortium	AA033292
794 A	3 0210A1F4	ESTs, Highly similar to ENDOTHELIAL ACTIN-BINDING PROTEIN [Homo sapiens] {IMAGE:466292}	466292 IMAGE Consortium	AA033393
795 A	5 0210A1F4	IMAGE EST {IMAGE:465085}	465085 IMAGE Consortium	AA032550
796 A	7 0210A1F4	Acetylcholinesterase {IMAGE:466476}	466476 IMAGE Consortium	AA030863
797 A	9 0210A1F4	IMAGE EST {IMAGE:465560}	465560 IMAGE Consortium	AA032862
798 A	11 0210A1F4	IMAGE EST {IMAGE:466909}	466909 IMAGE Consortium	AA031232
799 C	1 0210A1F4	IMAGE EST {IMAGE:465133}	465133 IMAGE Consortium	AA032723
800 C	3 0210A1F4	IMAGE EST {IMAGE:466644}	466644 IMAGE Consortium	AA030999
801 C	5 0210A1F4	ESTs, Weakly similar to HYPOTHETICAL 31.9 KD PROTEIN IN GOG5-CLG1 INTERGENIC REGION [S.cerevisiae] {IMAGE:465848}	465848 IMAGE Consortium	AA032797
802 C	7 0210A1F4	IMAGE EST {IMAGE:467850}	467850 IMAGE Consortium	AA036542
803 C	9 0210A1F4	IMAGE EST {IMAGE:466403}	466403 IMAGE Consortium	AA032381
804 C	11 0210A1F4	IMAGE EST {IMAGE:466592}	466592 IMAGE Consortium	AA030919
805 E	1 0210A1F4	IMAGE EST {IMAGE:466803}	466803 IMAGE Consortium	AA031075
806 E	3 0210A1F4	IMAGE EST {IMAGE:466616}	466616 IMAGE Consortium	AA030942
807 E	5 0210A1F4	IMAGE EST {IMAGE:465657}	465657 IMAGE Consortium	AA033443
808 E	7 0210A1F4	IMAGE EST {IMAGE:465352}	465352 IMAGE Consortium	AA032438
809 E	9 0210A1F4	Mus musculus EGF repeat transmembrane protein mRNA, complete cds {IMAGE:465986}	465986 IMAGE Consortium	AA031052
810 E	11 0210A1F4	IMAGE EST {IMAGE:467219}	467219 IMAGE Consortium	AA034883
811 G	1 0210A1F4	ESTs, Weakly similar to T03G11.6 gene product [C.elegans] {IMAGE:467379}	467379 IMAGE Consortium	AA036624
812 G	3 0210A1F4	Mus musculus C-terminal binding protein 2 mRNA, complete cds {IMAGE:467669}	467669 IMAGE Consortium	AA036337
813 G	5 0210A1F4	IMAGE EST {IMAGE:466255}	466255 IMAGE Consortium	AA033308
814 G	7 0210A1F4	IMAGE EST {IMAGE:466385}	466385 IMAGE Consortium	AA034758
815 G	9 0210A1F4	IMAGE EST {IMAGE:466438}	466438 IMAGE Consortium	AA034771
816 G	11 0210A1F4	Mus musculus syntaxin 7 (Syn7) mRNA, complete cds {IMAGE:468472}	468472 IMAGE Consortium	AA014394
817 A	1 0217A1F5	ESTs, Weakly similar to EUKARYOTIC INITIATION FACTOR 4B [Homo sapiens] {IMAGE:466333}	466333 IMAGE Consortium	AA033370
818 A	3 0217A1F5	IMAGE EST {IMAGE:466199}	466199 IMAGE Consortium	AA032344
819 A	5 0217A1F5	ESTs {IMAGE:465628}	465628 IMAGE Consortium	AA033201
820 A	7 0217A1F5	IMAGE EST {IMAGE:466905}	466905 IMAGE Consortium	AA031230
821 A	9 0217A1F5	IMAGE EST {IMAGE:467134}	467134 IMAGE Consortium	AA034829
822 A	11 0217A1F5	Mus musculus rbm3 mRNA, complete cds {IMAGE:467238}	467238 IMAGE Consortium	AA034857
823 C	1 0217A1F5	IMAGE EST {IMAGE:474156}	474156 IMAGE Consortium	AA038665
824 C	3 0217A1F5	Mus musculus BS4 peptide mRNA, complete cds	468813 IMAGE ConsoGtium	AA035926

		{IMAGE:468813}		
825 C	5	0217A1F5	IMAGE EST {IMAGE:472304}	472304 IMAGE Consortium AA036323
826 C	7	0217A1F5	ESTs, Highly similar to CREATINE KINASE, SARCOMERIC MITOCHONDRIAL PRECURSOR [Rattus norvegicus] {IMAGE:472860}	472860 IMAGE Consortium AA038095
827 C	9	0217A1F5	IMAGE EST {IMAGE:473745}	473745 IMAGE Consortium AA038412
828 C	11	0217A1F5	IMAGE EST {IMAGE:474094}	474094 IMAGE Consortium AA038717
829 E	1	0217A1F5	ESTs, Highly similar to ERGIC-53 PROTEIN PRECURSOR [Homo sapiens] {IMAGE:469029}	469029 IMAGE Consortium AA036111
830 E	3	0217A1F5	Mus musculus Cek7 ligand mRNA, complete cds {IMAGE:472400}	472400 IMAGE Consortium AA036231
831 E	5	0217A1F5	ESTs, Highly similar to PEPSINOGEN F PRECURSOR [Oryctolagus cuniculus] {IMAGE:472743}	472743 IMAGE Consortium AA038137
832 E	7	0217A1F5	IMAGE EST {IMAGE:474107}	474107 IMAGE Consortium AA038649
833 E	9	0217A1F5	ESTs {IMAGE:468571}	468571 IMAGE Consortium AA014335
834 E	11	0217A1F5	GATA-binding protein 1 {IMAGE:468938}	468938 IMAGE Consortium AA036275
835 G	1	0217A1F5	IMAGE EST {IMAGE:472445}	472445 IMAGE Consortium AA036156
836 G	3	0217A1F5	IMAGE EST {IMAGE:472805}	472805 IMAGE Consortium AA037997
837 G	5	0217A1F5	IMAGE EST {IMAGE:473401}	473401 IMAGE Consortium AA038556
838 G	7	0217A1F5	ESTs {IMAGE:475586}	475586 IMAGE Consortium AA050193
839 G	9	0217A1F5	IMAGE EST {IMAGE:468954}	468954 IMAGE Consortium AA036086
840 G	11	0217A1F5	ESTs {IMAGE:472783}	472783 IMAGE Consortium AA037991
841 A	1	021EA1F6	IMAGE EST {IMAGE:476408}	476408 IMAGE Consortium AA050650
842 A	3	021EA1F6	ESTs {IMAGE:477828}	477828 IMAGE Consortium AA048602
843 A	5	021EA1F6	Mus musculus BAF57 (Baf57) gene, complete cds {IMAGE:476738}	476738 IMAGE Consortium AA050900
844 A	7	021EA1F6	ESTs {IMAGE:476577}	476577 IMAGE Consortium AA050235
845 A	9	021EA1F6	ESTs, Weakly similar to serine threonine kinase [M.musculus] {IMAGE:476687}	476687 IMAGE Consortium AA050680
846 A	11	021EA1F6	Excision repair 1 {IMAGE:475399}	475399 IMAGE Consortium AA041710
847 C	1	021EA1F6	ESTs {IMAGE:476086}	476086 IMAGE Consortium AA049861
848 C	3	021EA1F6	IMAGE EST {IMAGE:478178}	478178 IMAGE Consortium AA049616
849 C	5	021EA1F6	IMAGE EST {IMAGE:478159}	478159 IMAGE Consortium AA049611
850 C	7	021EA1F6	IMAGE EST {IMAGE:476809}	476809 IMAGE Consortium AA050511
851 C	9	021EA1F6	EST {IMAGE:476961}	476961 IMAGE Consortium AA048029
852 C	11	021EA1F6	ESTs {IMAGE:476972}	476972 IMAGE Consortium AA048101
853 E	1	021EA1F6	RADIXIN {IMAGE:476178}	476178 IMAGE Consortium AA050683
854 E	3	021EA1F6	ESTs, Highly similar to HYPOTHETICAL 66.5 KD PROTEIN F02A9.5 IN CHROMOSOME III [Caenorhabditis elegans] {IMAGE:476754}	476754 IMAGE Consortium AA050443
855 E	5	021EA1F6	Cytochrome P450, 2e1, ethanol inducible {IMAGE:475992}	475992 IMAGE Consortium AA049906
856 E	7	021EA1F6	Suppressor of Ty 6 (S. cerevisiae) homolog {IMAGE:477535}	477535 IMAGE Consortium AA048444
857 E	9	021EA1F6	DNA POLYMERASE ALPHA/PRIMASE ASSOCIATED	476123 IMAGE Consortium AA050378

		SUBUNIT {IMAGE:476123}		
858 E	11 021EA1F6	EST {IMAGE:476278}	476278 IMAGE Consortium	AA050573
859 G	1 021EA1F6	Mus musculus mRNA for CIRP, complete cds {IMAGE:475679}	475679 IMAGE Consortium	AA050300
860 G	3 021EA1F6	IMAGE EST {IMAGE:476367}	476367 IMAGE Consortium	AA050610
861 G	5 021EA1F6	ESTs {IMAGE:477988}	477988 IMAGE Consortium	AA049432
862 G	7 021EA1F6	IMAGE EST {IMAGE:477002}	477002 IMAGE Consortium	AA048118
863 G	9 021EA1F6	ESTs, Weakly similar to cytoplasmic protein Ndr1 [M.musculus] {IMAGE:475403}	475403 IMAGE Consortium	AA041825
864 G	11 021EA1F6	ESTs {IMAGE:476987}	476987 IMAGE Consortium	AA047985
865 A	1 021LA1F7	IMAGE EST {IMAGE:474208}	474208 IMAGE Consortium	AA038743
866 A	3 021LA1F7	IMAGE EST {IMAGE:468886}	468886 IMAGE Consortium	AA036176
867 A	5 021LA1F7	ESTs {IMAGE:472664}	472664 IMAGE Consortium	AA037977
868 A	7 021LA1F7	IMAGE EST {IMAGE:473802}	473802 IMAGE Consortium	AA038392
869 A	9 021LA1F7	IMAGE EST {IMAGE:468816}	468816 IMAGE Consortium	AA035895
870 A	11 021LA1F7	Mus musculus stanniocalcin precursor, mRNA, complete cds {IMAGE:474298}	474298 IMAGE Consortium	AA038788
871 C	1 021LA1F7	ESTs {IMAGE:469037}	469037 IMAGE Consortium	AA036370
872 C	3 021LA1F7	IMAGE EST {IMAGE:468970}	468970 IMAGE Consortium	AA035994
873 C	5 021LA1F7	IMAGE EST {IMAGE:469047}	469047 IMAGE Consortium	AA036223
874 C	7 021LA1F7	IMAGE EST {IMAGE:472320}	472320 IMAGE Consortium	AA036212
875 C	9 021LA1F7	IMAGE EST {IMAGE:472691}	472691 IMAGE Consortium	AA038156
876 C	11 021LA1F7	IMAGE EST {IMAGE:474044}	474044 IMAGE Consortium	AA038694
877 E	1 021LA1F7	Cytochrome P450, 2e1, ethanol inducible {IMAGE:476836}	476836 IMAGE Consortium	AA050328
878 E	3 021LA1F7	ESTs {IMAGE:477056}	477056 IMAGE Consortium	AA048139
879 E	5 021LA1F7	IMAGE EST {IMAGE:476917}	476917 IMAGE Consortium	AA048016
880 E	7 021LA1F7	IMAGE EST {IMAGE:476899}	476899 IMAGE Consortium	AA050888
881 E	9 021LA1F7	Mus musculus bystin mRNA, complete cds {IMAGE:475670}	475670 IMAGE Consortium	AA044532
882 E	11 021LA1F7	IMAGE EST {IMAGE:476400}	476400 IMAGE Consortium	AA049991
883 G	1 021LA1F7	IMAGE EST {IMAGE:476989}	476989 IMAGE Consortium	AA047986
884 G	3 021LA1F7	ESTs {IMAGE:477217}	477217 IMAGE Consortium	AA048224
885 G	5 021LA1F7	Mus musculus Mlark mRNA, complete cds {IMAGE:476312}	476312 IMAGE Consortium	AA050727
886 G	7 021LA1F7	IMAGE EST {IMAGE:477588}	477588 IMAGE Consortium	AA048459
887 G	9 021LA1F7	ESTs {IMAGE:477881}	477881 IMAGE Consortium	AA049408
888 G	11 021LA1F7	IMAGE EST {IMAGE:478058}	478058 IMAGE Consortium	AA049294
889 A	1 021SA1F8	Thymidine kinase 1 {IMAGE:475451}	475451 IMAGE Consortium	AA041834
890 A	3 021SA1F8	ESTs, Highly similar to CARBOXYPEPTIDASE B PRECURSOR [Rattus norvegicus] {IMAGE:477035}	477035 IMAGE Consortium	AA048136
891 A	5 021SA1F8	MYELOID DIFFERENTIATION PRIMARY RESPONSE PROTEIN MYD116 {IMAGE:475803}	475803 IMAGE Consortium	AA050417
892 A	7 021SA1F8	ESTs {IMAGE:475929}	475929 IMAGE Consortium	AA050341
893 A	9 021SA1F8	IMAGE EST {IMAGE:476340}	476340 IMAGE Consortium	AA050647
894 A	11 021SA1F8	ESTs, Highly similar to angio-associated migratory cell	477784 IMAGE Consortium	AA049376

		protein [H.sapiens] {IMAGE:477784}		
895 C	1 021SA1F8	Paired mesoderm homeobox 1 {IMAGE:476293}	476293 IMAGE Consortium	AA050726
896 C	3 021SA1F8	CD151 antigen {IMAGE:476416}	476416 IMAGE Consortium	AA050218
897 C	5 021SA1F8	Phosphoribosylglycinamide formyltransferase {IMAGE:477593}	477593 IMAGE Consortium	AA048560
898 C	7 021SA1F8	ESTs {IMAGE:476431}	476431 IMAGE Consortium	AA050723
899 C	9 021SA1F8	IMAGE EST {IMAGE:476661}	476661 IMAGE Consortium	AA050851
900 C	11 021SA1F8	Cytochrome P450, 2e1, ethanol inducible {IMAGE:478053}	478053 IMAGE Consortium	AA049507
901 E	1 021SA1F8	Mus musculus mRNA for DEDD protein {IMAGE:478020}	478020 IMAGE Consortium	AA049467
902 E	3 021SA1F8	Pleiotrophin {IMAGE:478168}	478168 IMAGE Consortium	AA049699
903 E	5 021SA1F8	IMAGE EST {IMAGE:478101}	478101 IMAGE Consortium	AA049584
904 E	7 021SA1F8	Beta-spectrin 3 {IMAGE:478354}	478354 IMAGE Consortium	AA049581
905 E	9 021SA1F8	Bcl2-like {IMAGE:478723}	478723 IMAGE Consortium	AA051441
906 E	11 021SA1F8	IMAGE EST {IMAGE:480196}	480196 IMAGE Consortium	AA058055
907 G	1 021SA1F8	IMAGE EST {IMAGE:479159}	479159 IMAGE Consortium	AA049330
908 G	3 021SA1F8	IMAGE EST {IMAGE:478378}	478378 IMAGE Consortium	AA049768
909 G	5 021SA1F8	Cytochrome P450, 2e1, ethanol inducible {IMAGE:479558}	479558 IMAGE Consortium	AA048980
910 G	7 021SA1F8	IMAGE EST {IMAGE:478388}	478388 IMAGE Consortium	AA049946
911 G	9 021SA1F8	IMAGE EST {IMAGE:479985}	479985 IMAGE Consortium	AA051577
912 G	11 021SA1F8	IMAGE EST {IMAGE:478932}	478932 IMAGE Consortium	AA049055
913 A	1 021ZA1F9	Mus musculus mRNA for SAP18, Sin3-associated-polypeptide 18 {IMAGE:479654}	479654 IMAGE Consortium	AA049308
914 A	3 021ZA1F9	Guanine nucleotide binding protein, alpha o {IMAGE:480073}	480073 IMAGE Consortium	AA051709
915 A	5 021ZA1F9	ESTs {IMAGE:478927}	478927 IMAGE Consortium	AA048938
916 A	7 021ZA1F9	IMAGE EST {IMAGE:478795}	478795 IMAGE Consortium	AA051236
917 A	9 021ZA1F9	ESTs {IMAGE:479186}	479186 IMAGE Consortium	AA049125
918 A	11 021ZA1F9	ESTs {IMAGE:479314}	479314 IMAGE Consortium	AA048763
919 C	1 021ZA1F9	IMAGE EST {IMAGE:478843}	478843 IMAGE Consortium	AA048808
920 C	3 021ZA1F9	Mouse fast skeletal troponin C (sTnC) gene, complete cds {IMAGE:479354}	479354 IMAGE Consortium	AA048790
921 C	5 021ZA1F9	Mus musculus Bik-like killer protein (Bik) mRNA, complete cds {IMAGE:480492}	480492 IMAGE Consortium	AA064058
922 C	7 021ZA1F9	EST {IMAGE:479369}	479369 IMAGE Consortium	AA049041
923 C	9 021ZA1F9	IMAGE EST {IMAGE:478416}	478416 IMAGE Consortium	AA049778
924 C	11 021ZA1F9	IMAGE EST {IMAGE:478614}	478614 IMAGE Consortium	AA050068
925 E	1 021ZA1F9	IMAGE EST {IMAGE:479764}	479764 IMAGE Consortium	AA051544
926 E	3 021ZA1F9	ESTs {IMAGE:478580}	478580 IMAGE Consortium	AA051338
927 E	5 021ZA1F9	Neurotrophic tyrosine kinase, receptor, type 3 {IMAGE:478848}	478848 IMAGE Consortium	AA051250
928 E	7 021ZA1F9	ESTs, Highly similar to THIMET OLIGOPEPTIDASE [Rattus norvegicus] {IMAGE:480082}	480082 IMAGE Consortium	AA051777
929 E	9 021ZA1F9	IMAGE EST {IMAGE:479387}	479387 IMAGE Consortium	AA048754
930 E	11 021ZA1F9	IMAGE EST {IMAGE:479247}	479247 IMAGE Consortium	AA048730

931 G	1 021ZA1F9	ESTs {IMAGE:479435}	479435 IMAGE Consortium	AA049249
932 G	3 021ZA1F9	IMAGE EST {IMAGE:480284}	480284 IMAGE Consortium	AA058097
933 G	5 021ZA1F9	ESTs {IMAGE:478680}	478680 IMAGE Consortium	AA051371
934 G	7 021ZA1F9	ESTs {IMAGE:479741}	479741 IMAGE Consortium	AA051541
935 G	9 021ZA1F9	ESTs {IMAGE:478693}	478693 IMAGE Consortium	AA051406
936 G	11 021ZA1F9	Cytochrome P450, 2e1, ethanol inducible {IMAGE:480122}	480122 IMAGE Consortium	AA051721
937 A	1 021JA1FA	IMAGE EST {IMAGE:483662}	483662 IMAGE Consortium	AA061365
938 A	3 021JA1FA	IMAGE EST {IMAGE:483381}	483381 IMAGE Consortium	AA061870
939 A	5 021JA1FA	ESTs, Moderately similar to HYPOTHETICAL 97.6 KD PROTEIN IN SHP1-SEC17 INTERGENIC REGION [Saccharomyces cerevisiae] {IMAGE:482177}	482177 IMAGE Consortium	AA059768
940 A	7 021JA1FA	ESTs {IMAGE:480697}	480697 IMAGE Consortium	AA064069
941 A	9 021JA1FA	IMAGE EST {IMAGE:480656}	480656 IMAGE Consortium	AA064127
942 A	11 021JA1FA	IMAGE EST {IMAGE:482371}	482371 IMAGE Consortium	AA063754
943 C	1 021JA1FA	IMAGE EST {IMAGE:483442}	483442 IMAGE Consortium	AA061721
944 C	3 021JA1FA	IMAGE EST {IMAGE:481926}	481926 IMAGE Consortium	AA059926
945 C	5 021JA1FA	ESTs {IMAGE:482027}	482027 IMAGE Consortium	AA059892
946 C	7 021JA1FA	IMAGE EST {IMAGE:482058}	482058 IMAGE Consortium	AA059979
947 C	9 021JA1FA	Mus musculus Stra13 mRNA, complete cds {IMAGE:480896}	480896 IMAGE Consortium	AA064241
948 C	11 021JA1FA	IMAGE EST {IMAGE:480876}	480876 IMAGE Consortium	AA064236
949 E	1 021JA1FA	Synaptobrevin 2 {IMAGE:483687}	483687 IMAGE Consortium	AA061343
950 E	3 021JA1FA	ESTs {IMAGE:483597}	483597 IMAGE Consortium	AA061662
951 E	5 021JA1FA	Mus musculus diacylglycerol acyltransferase (Dgat) mRNA, complete cds {IMAGE:483777}	483777 IMAGE Consortium	AA061439
952 E	7 021JA1FA	Metallothionein 1 {IMAGE:480920}	480920 IMAGE Consortium	AA064247
953 E	9 021JA1FA	ESTs {IMAGE:480900}	480900 IMAGE Consortium	AA064243
954 E	11 021JA1FA	EST {IMAGE:480859}	480859 IMAGE Consortium	AA060540
955 G	1 021JA1FA	ESTs {IMAGE:482142}	482142 IMAGE Consortium	AA059644
956 G	3 021JA1FA	Delta-like 1 homolog (Drosophila) {IMAGE:481908}	481908 IMAGE Consortium	AA059653
957 G	5 021JA1FA	ESTs {IMAGE:481867}	481867 IMAGE Consortium	AA059812
958 G	7 021JA1FA	IMAGE EST {IMAGE:480986}	480986 IMAGE Consortium	AA059557
959 G	9 021JA1FA	ESTs {IMAGE:480729}	480729 IMAGE Consortium	AA064165
960 G	11 021JA1FA	ESTs {IMAGE:480996}	480996 IMAGE Consortium	AA060191
961 A	1 021QA1FB	Mus musculus GTPase-activating protein GAP111 mRNA, complete cds {IMAGE:478472}	478472 IMAGE Consortium	AA050000
962 A	3 021QA1FB	IMAGE EST {IMAGE:478387}	478387 IMAGE Consortium	AA049948
963 A	5 021QA1FB	IMAGE EST {IMAGE:478485}	478485 IMAGE Consortium	AA050028
964 A	7 021QA1FB	ESTs {IMAGE:480107}	480107 IMAGE Consortium	AA051789
965 A	9 021QA1FB	IMAGE EST {IMAGE:480255}	480255 IMAGE Consortium	AA058149
966 A	11 021QA1FB	IMAGE EST {IMAGE:480166}	480166 IMAGE Consortium	AA058021
967 C	1 021QA1FB	IMAGE EST {IMAGE:479699}	479699 IMAGE Consortium	AA051607
968 C	3 021QA1FB	EST {IMAGE:478435}	478435 IMAGE Consortium	AA049963
969 C	5 021QA1FB	ESTs {IMAGE:478509}	478509 IMAGE Consortium	AA050035

970 C	7 021QA1FB	ESTs {IMAGE:478826}	478826 IMAGE Consortium	AA051246
971 C	9 021QA1FB	IMAGE EST {IMAGE:478763}	478763 IMAGE Consortium	AA050999
972 C	11 021QA1FB	ESTs {IMAGE:480295}	480295 IMAGE Consortium	AA058209
973 E	1 021QA1FB	Protein kinase, cAMP dependent, catalytic, alpha {IMAGE:478782}	478782 IMAGE Consortium	AA051181
974 E	3 021QA1FB	ESTs, Highly similar to INTERFERON-INDUCIBLE PROTEIN [Rattus norvegicus] {IMAGE:479027}	479027 IMAGE Consortium	AA048837
975 E	5 021QA1FB	ESTs, Highly similar to MITOCHONDRIAL RESPIRATORY CHAIN COMPLEXES ASSEMBLY PROTEIN RCA1 [Saccharomyces cerevisiae] {IMAGE:480263}	480263 IMAGE Consortium	AA058153
976 E	7 021QA1FB	ESTs, Moderately similar to KIAA0110 [H.sapiens] {IMAGE:479455}	479455 IMAGE Consortium	AA048950
977 E	9 021QA1FB	ESTs, Highly similar to NEDD-4 PROTEIN [Homo sapiens] {IMAGE:479436}	479436 IMAGE Consortium	AA049246
978 E	11 021QA1FB	IMAGE EST {IMAGE:480599}	480599 IMAGE Consortium	AA059521
979 G	1 021QA1FB	Metallothionein 1 {IMAGE:480068}	480068 IMAGE Consortium	AA051654
980 G	3 021QA1FB	IMAGE EST {IMAGE:479432}	479432 IMAGE Consortium	AA049244
981 G	5 021QA1FB	IMAGE EST {IMAGE:479269}	479269 IMAGE Consortium	AA048539
982 G	7 021QA1FB	Mus musculus type VI collagen alpha 3 subunit mRNA, complete cds {IMAGE:480620}	480620 IMAGE Consortium	AA059524
983 G	9 021QA1FB	IMAGE EST {IMAGE:481008}	481008 IMAGE Consortium	AA059563
984 G	11 021QA1FB	ESTs, Highly similar to PHENYLALANYL-TRNA SYNTHETASE MITOCHONDRIAL PRECURSOR [Saccharomyces cerevisiae] {IMAGE:480738}	480738 IMAGE Consortium	AA064206
985 A	1 021XA1FC	IMAGE EST {IMAGE:481956}	481956 IMAGE Consortium	AA059636
986 A	3 021XA1FC	ESTs {IMAGE:480747}	480747 IMAGE Consortium	AA060703
987 A	5 021XA1FC	IMAGE EST {IMAGE:482366}	482366 IMAGE Consortium	AA060983
988 A	7 021XA1FC	Mouse mRNA for pituitary adenylate cyclase activating polypeptide {IMAGE:480980}	480980 IMAGE Consortium	AA060249
989 A	9 021XA1FC	IMAGE EST {IMAGE:482677}	482677 IMAGE Consortium	AA061997
990 A	11 021XA1FC	IMAGE EST {IMAGE:482955}	482955 IMAGE Consortium	AA060806
991 C	1 021XA1FC	ESTs {IMAGE:484348}	484348 IMAGE Consortium	AA080239
992 C	3 021XA1FC	ESTs, Highly similar to FRUCTOSE-1,6-BISPHOSPHATASE [Rattus norvegicus] {IMAGE:482534}	482534 IMAGE Consortium	AA063840
993 C	5 021XA1FC	IMAGE EST {IMAGE:482592}	482592 IMAGE Consortium	AA061834
994 C	7 021XA1FC	ESTs {IMAGE:482910}	482910 IMAGE Consortium	AA062133
995 C	9 021XA1FC	IMAGE EST {IMAGE:482773}	482773 IMAGE Consortium	AA062035
996 C	11 021XA1FC	Amyloid beta (A4) precursor protein-binding, family A, member 2 {IMAGE:481145}	481145 IMAGE Consortium	AA060361
997 E	1 021XA1FC	ESTs {IMAGE:482664}	482664 IMAGE Consortium	AA061989
998 E	3 021XA1FC	ESTs {IMAGE:482606}	482606 IMAGE Consortium	AA061937
999 E	5 021XA1FC	ESTs {IMAGE:481198}	481198 IMAGE Consortium	AA060376
1000 E	7 021XA1FC	ESTs, Highly similar to T-CADHERIN PRECURSOR [Gallus gallus] {IMAGE:482972}	482972 IMAGE Consortium	AA061041

1001 E	9 021XA1FC	IMAGE EST {IMAGE:481526}	481526 IMAGE Consortium	AA060624
1002 E	11 021XA1FC	Mus musculus Cdc42/Rac effector kinase PAK-A (Paka) mRNA, complete cds {IMAGE:483684}	483684 IMAGE Consortium	AA061378
1003 G	1 021XA1FC	ESTs {IMAGE:481087}	481087 IMAGE Consortium	AA064288
1004 G	3 021XA1FC	Myeloid/lymphoid or mixed-lineage leukemia {IMAGE:481236}	481236 IMAGE Consortium	AA060386
1005 G	5 021XA1FC	Kinesin heavy chain member 4 {IMAGE:481146}	481146 IMAGE Consortium	AA060358
1006 G	7 021XA1FC	IMAGE EST {IMAGE:481390}	481390 IMAGE Consortium	AA060554
1007 G	9 021XA1FC	EST {IMAGE:483212}	483212 IMAGE Consortium	AA063802
1008 G	11 021XA1FC	IMAGE EST {IMAGE:483361}	483361 IMAGE Consortium	AA059943
1009 A	1 0214A1FD	ATP-binding cassette 1 {IMAGE:482369}	482369 IMAGE Consortium	AA063753
1010 A	3 0214A1FD	IMAGE EST {IMAGE:482689}	482689 IMAGE Consortium	AA062006
1011 A	5 0214A1FD	ESTs {IMAGE:482865}	482865 IMAGE Consortium	AA062139
1012 A	7 0214A1FD	IMAGE EST {IMAGE:482755}	482755 IMAGE Consortium	AA061913
1013 A	9 0214A1FD	Procollagen, type IX, alpha 1 {IMAGE:481151}	481151 IMAGE Consortium	AA060364
1014 A	11 0214A1FD	IMAGE EST {IMAGE:483649}	483649 IMAGE Consortium	AA061366
1015 C	1 0214A1FD	Uncoupling protein 3, mitochondrial {IMAGE:482847}	482847 IMAGE Consortium	AA062091
1016 C	3 0214A1FD	IMAGE EST {IMAGE:483006}	483006 IMAGE Consortium	AA060858
1017 C	5 0214A1FD	ESTs {IMAGE:483578}	483578 IMAGE Consortium	AA061744
1018 C	7 0214A1FD	IMAGE EST {IMAGE:481773}	481773 IMAGE Consortium	AA060184
1019 C	9 0214A1FD	ESTs {IMAGE:483476}	483476 IMAGE Consortium	AA060036
1020 C	11 0214A1FD	ESTs, Moderately similar to HYPOTHETICAL 28.5 KD PROTEIN ZK1236.7 IN CHROMOSOME III [Caenorhabditis elegans] {IMAGE:483751}	483751 IMAGE Consortium	AA061407
1021 E	1 0214A1FD	ESTs {IMAGE:483150}	483150 IMAGE Consortium	AA060910
1022 E	3 0214A1FD	IMAGE EST {IMAGE:483189}	483189 IMAGE Consortium	AA063798
1023 E	5 0214A1FD	IMAGE EST {IMAGE:483626}	483626 IMAGE Consortium	AA061285
1024 E	7 0214A1FD	EST {IMAGE:483585}	483585 IMAGE Consortium	AA061905
1025 E	9 0214A1FD	ESTs {IMAGE:483572}	483572 IMAGE Consortium	AA061650
1026 E	11 0214A1FD	IMAGE EST {IMAGE:483775}	483775 IMAGE Consortium	AA061438
1027 G	1 0214A1FD	ESTs {IMAGE:493671}	493671 IMAGE Consortium	AA098264
1028 G	3 0214A1FD	IMAGE EST {IMAGE:573193}	573193 IMAGE Consortium	AA119297
1029 G	5 0214A1FD	Mus musculus ubiquitin-conjugating enzyme UbcM2 mRNA, complete cds {IMAGE:570902}	570902 IMAGE Consortium	AA110280
1030 G	7 0214A1FD	IMAGE EST {IMAGE:484280}	484280 IMAGE Consortium	AA080234
1031 G	9 0214A1FD	Mus musculus RalBP1-associated EH domain protein Reps1 (reps1) mRNA, complete cds {IMAGE:484194}	484194 IMAGE Consortium	AA073700
1032 G	11 0214A1FD	ESTs {IMAGE:492408}	492408 IMAGE Consortium	AA097181
1033 A	1 021BA1FE	ESTs, Highly similar to PROBABLE CALCIUM-BINDING PROTEIN PMP41 [Mus musculus] {IMAGE:572282}	572282 IMAGE Consortium	AA110246
1034 A	3 021BA1FE	ESTs {IMAGE:572201}	572201 IMAGE Consortium	AA105240
1035 A	5 021BA1FE	IMAGE EST {IMAGE:572934}	572934 IMAGE Consortium	AA117039
1036 A	7 021BA1FE	IMAGE EST {IMAGE:493558}	493558 IMAGE Consortium	AA087123



1037 A	9 021BA1FE	ESTs {IMAGE:573195}	573195 IMAGE Consortium	AA119298
1038 A	11 021BA1FE	ESTs {IMAGE:573470}	573470 IMAGE Consortium	AA118626
1039 C	1 021BA1FE	IMAGE EST {IMAGE:572943}	572943 IMAGE Consortium	AA111556
1040 C	3 021BA1FE	ESTs {IMAGE:573093}	573093 IMAGE Consortium	AA110645
1041 C	5 021BA1FE	ESTs {IMAGE:573243}	573243 IMAGE Consortium	AA119313
1042 C	7 021BA1FE	ESTs, Moderately similar to unknown [H.sapiens] {IMAGE:573226}	573226 IMAGE Consortium	AA119354
1043 C	9 021BA1FE	IMAGE EST {IMAGE:484485}	484485 IMAGE Consortium	AA073913
1044 C	11 021BA1FE	Calmodulin 3 {IMAGE:571789}	571789 IMAGE Consortium	AA109041
1045 E	1 021BA1FE	ESTs {IMAGE:570612}	570612 IMAGE Consortium	AA108670
1046 E	3 021BA1FE	IMAGE EST {IMAGE:571067}	571067 IMAGE Consortium	AA108457
1047 E	5 021BA1FE	ESTs, Highly similar to PROTEASOME THETA CHAIN [Rattus norvegicus] {IMAGE:571569}	571569 IMAGE Consortium	AA108678
1048 E	7 021BA1FE	Mus musculus zeta proteasome chain mRNA, complete cds {IMAGE:572285}	572285 IMAGE Consortium	AA110250
1049 E	9 021BA1FE	Mus musculus tetraspan TM4SF (Tspan-6) mRNA, complete cds {IMAGE:572156}	572156 IMAGE Consortium	AA105224
1050 E	11 021BA1FE	IMAGE EST {IMAGE:493683}	493683 IMAGE Consortium	AA087204
1051 G	1 021BA1FE	Hemoglobin alpha, adult chain 1 {IMAGE:571819}	571819 IMAGE Consortium	AA109900
1052 G	3 021BA1FE	IMAGE EST {IMAGE:492972}	492972 IMAGE Consortium	AA097471
1053 G	5 021BA1FE	ESTs {IMAGE:493122}	493122 IMAGE Consortium	AA097379
1054 G	7 021BA1FE	ESTs {IMAGE:572881}	572881 IMAGE Consortium	AA110713
1055 G	9 021BA1FE	Mus musculus alpha-actinin 3 (Actn3) mRNA, complete cds {IMAGE:493675}	493675 IMAGE Consortium	AA098356
1056 G	11 021BA1FE	IMAGE EST {IMAGE:573384}	573384 IMAGE Consortium	AA118700
1057 A	1 021IA1FF	ESTs, Moderately similar to phosphate cyclase [H.sapiens] {IMAGE:573205}	573205 IMAGE Consortium	AA119351
1058 A	3 021IA1FF	IMAGE EST {IMAGE:571348}	571348 IMAGE Consortium	AA108848
1059 A	5 021IA1FF	ESTs {IMAGE:492456}	492456 IMAGE Consortium	AA097497
1060 A	7 021IA1FF	Cytochrome P450, 2e1, ethanol inducible {IMAGE:492407}	492407 IMAGE Consortium	AA097387
1061 A	9 021IA1FF	IMAGE EST {IMAGE:572256}	572256 IMAGE Consortium	AA110195
1062 A	11 021IA1FF	IMAGE EST {IMAGE:493165}	493165 IMAGE Consortium	AA097784
1063 C	1 021IA1FF	ESTs, Weakly similar to GLUTATHIONE S-TRANSFERASE, MITOCHONDRIAL [Rattus norvegicus] {IMAGE:571334}	571334 IMAGE Consortium	AA108741
1064 C	3 021IA1FF	ESTs, Moderately similar to P120 PROTEIN [Mus musculus] {IMAGE:571753}	571753 IMAGE Consortium	AA108926
1065 C	5 021IA1FF	ESTs, Highly similar to EUKARYOTIC INITIATION FACTOR 4A-LIKE NUK-34 [Homo sapiens] {IMAGE:571841}	571841 IMAGE Consortium	AA109923
1066 C	7 021IA1FF	Myeloid differentiation primary response gene 88 {IMAGE:572317}	572317 IMAGE Consortium	AA110412
1067 C	9 021IA1FF	Mus musculus cyclin D-interacting myb-like protein (Dmp1) mRNA, complete cds {IMAGE:572713}	572713 IMAGE Consortium	AA111662
1068 C	11 021IA1FF	IMAGE EST {IMAGE:493442}	493442 IMAGE Consortium	AA086909
1069 E	1 021IA1FF	CDC-like kinase {IMAGE:571779}	571779 IMAGE Consortium	AA108933

1070 E	3 021IA1FF	ESTs, Highly similar to SWI/SNF complex 170 KDa subunit [H.sapiens] {IMAGE:572000}	572000 IMAGE Consortium	AA108369
1071 E	5 021IA1FF	ESTs {IMAGE:493256}	493256 IMAGE Consortium	AA098158
1072 E	7 021IA1FF	ESTs {IMAGE:493238}	493238 IMAGE Consortium	AA098154
1073 E	9 021IA1FF	ESTs, Highly similar to ALDEHYDE DEHYDROGENASE, E3 ISOZYME [Homo sapiens] {IMAGE:572614}	572614 IMAGE Consortium	AA109914
1074 E	11 021IA1FF	IMAGE EST {IMAGE:572869}	572869 IMAGE Consortium	AA110805
1075 G	1 021IA1FF	IMAGE EST {IMAGE:492945}	492945 IMAGE Consortium	AA097466
1076 G	3 021IA1FF	IMAGE EST {IMAGE:493263}	493263 IMAGE Consortium	AA098164
1077 G	5 021IA1FF	IMAGE EST {IMAGE:572463}	572463 IMAGE Consortium	AA110679
1078 G	7 021IA1FF	IMAGE EST {IMAGE:572471}	572471 IMAGE Consortium	AA109938
1079 G	9 021IA1FF	ESTs, Highly similar to H-2 class I histocompatibility antigen T15 [M.musculus] {IMAGE:493481}	493481 IMAGE Consortium	AA086935
1080 G	11 021IA1FF	IMAGE EST {IMAGE:493605}	493605 IMAGE Consortium	AA087141
1081 A	1 021PA1FG	IMAGE EST {IMAGE:572604}	572604 IMAGE Consortium	AA111001
1082 A	3 021PA1FG	ESTs, Highly similar to 6-PYRUVOYL TETRAHYDROBIOPTERIN SYNTHASE PRECURSOR [Rattus norvegicus] {IMAGE:572953}	572953 IMAGE Consortium	AA117045
1083 A	5 021PA1FG	ESTs {IMAGE:493460}	493460 IMAGE Consortium	AA086835
1084 A	7 021PA1FG	IMAGE EST {IMAGE:573413}	573413 IMAGE Consortium	AA118754
1085 A	9 021PA1FG	IMAGE EST {IMAGE:573539}	573539 IMAGE Consortium	AA119321
1086 A	11 021PA1FG	ESTs {IMAGE:573449}	573449 IMAGE Consortium	AA111120
1087 C	1 021PA1FG	IMAGE EST {IMAGE:576477}	576477 IMAGE Consortium	AA117414
1088 C	3 021PA1FG	IMAGE EST {IMAGE:576797}	576797 IMAGE Consortium	AA123843
1089 C	5 021PA1FG	IMAGE EST {IMAGE:573285}	573285 IMAGE Consortium	AA110956
1090 C	7 021PA1FG	ESTs {IMAGE:575319}	575319 IMAGE Consortium	AA125133
1091 C	9 021PA1FG	IMAGE EST {IMAGE:575347}	575347 IMAGE Consortium	AA119278
1092 C	11 021PA1FG	Mus musculus polypeptide GalNAc transferase-T1 (ppGaNTase-T1) mRNA, complete cds {IMAGE:575760}	575760 IMAGE Consortium	AA124836
1093 E	1 021PA1FG	IMAGE EST {IMAGE:575051}	575051 IMAGE Consortium	AA123923
1094 E	3 021PA1FG	IMAGE EST {IMAGE:576924}	576924 IMAGE Consortium	AA122429
1095 E	5 021PA1FG	IMAGE EST {IMAGE:576835}	576835 IMAGE Consortium	AA120034
1096 E	7 021PA1FG	IMAGE EST {IMAGE:573581}	573581 IMAGE Consortium	AA119375
1097 E	9 021PA1FG	ESTs, Weakly similar to R11A8.7 [C.elegans] {IMAGE:575492}	575492 IMAGE Consortium	AA119523
1098 E	11 021PA1FG	ESTs, Highly similar to KIAA0370 [H.sapiens] {IMAGE:575497}	575497 IMAGE Consortium	AA125005
1099 G	1 021PA1FG	Mus musculus FAN protein mRNA, complete cds {IMAGE:576941}	576941 IMAGE Consortium	AA122528
1100 G	3 021PA1FG	ESTs, Moderately similar to syncollin [R.norvegicus] {IMAGE:576931}	576931 IMAGE Consortium	AA116528
1101 G	5 021PA1FG	X-ray repair complementing defective repair in Chinese hamster cells 5 {IMAGE:573725}	573725 IMAGE Consortium	AA118775
1102 G	7 021PA1FG	IMAGE EST {IMAGE:573588}	573588 IMAGE Consortium	AA118762

1103 G	9 021PA1FG	IMAGE EST {IMAGE:573595}	573595 IMAGE Consortium	AA111710
1104 G	11 021PA1FG	Tissue specific transplantation antigen P198-7 {IMAGE:574251}	574251 IMAGE Consortium	AA119255
1105 A	1 021WA1FH	Mus musculus 76 kDa tyrosine phosphoprotein SLP-76 mRNA, complete cds {IMAGE:576989}	576989 IMAGE Consortium	AA122543
1106 A	3 021WA1FH	ESTs {IMAGE:576901}	576901 IMAGE Consortium	AA122684
1107 A	5 021WA1FH	ESTs {IMAGE:575265}	575265 IMAGE Consortium	AA125395
1108 A	7 021WA1FH	ESTs {IMAGE:573589}	573589 IMAGE Consortium	AA118765
1109 A	9 021WA1FH	IMAGE EST {IMAGE:574236}	574236 IMAGE Consortium	AA119402
1110 A	11 021WA1FH	IMAGE EST {IMAGE:574294}	574294 IMAGE Consortium	AA111070
1111 C	1 021WA1FH	IMAGE EST {IMAGE:577021}	577021 IMAGE Consortium	AA124355
1112 C	3 021WA1FH	ESTs {IMAGE:576885}	576885 IMAGE Consortium	AA116480
1113 C	5 021WA1FH	ESTs, Highly similar to H-2 class I histocompatibility antigen T15 [M.musculus] {IMAGE:575122}	575122 IMAGE Consortium	AA124999
1114 C	7 021WA1FH	IMAGE EST {IMAGE:575320}	575320 IMAGE Consortium	AA119248
1115 C	9 021WA1FH	MAP KINASE PHOSPHATASE-1 {IMAGE:575665}	575665 IMAGE Consortium	AA125367
1116 C	11 021WA1FH	ESTs, Weakly similar to synaptotagmin VII [H.sapiens] {IMAGE:575700}	575700 IMAGE Consortium	AA123008
1117 E	1 021WA1FH	Transporter 1, ABC (ATP binding cassette) {IMAGE:576775}	576775 IMAGE Consortium	AA123837
1118 E	3 021WA1FH	IMAGE EST {IMAGE:576686}	576686 IMAGE Consortium	AA124417
1119 E	5 021WA1FH	IMAGE EST {IMAGE:577005}	577005 IMAGE Consortium	AA116465
1120 E	7 021WA1FH	IMAGE EST {IMAGE:573848}	573848 IMAGE Consortium	AA119048
1121 E	9 021WA1FH	ESTs {IMAGE:573807}	573807 IMAGE Consortium	AA118979
1122 E	11 021WA1FH	ESTs {IMAGE:574134}	574134 IMAGE Consortium	AA118838
1123 G	1 021WA1FH	IMAGE EST {IMAGE:575283}	575283 IMAGE Consortium	AA125302
1124 G	3 021WA1FH	Mus musculus syntaxin 7 (Syn7) mRNA, complete cds {IMAGE:573479}	573479 IMAGE Consortium	AA111166
1125 G	5 021WA1FH	ESTs, Moderately similar to KIAA0253 [H.sapiens] {IMAGE:573609}	573609 IMAGE Consortium	AA118848
1126 G	7 021WA1FH	IMAGE EST {IMAGE:573741}	573741 IMAGE Consortium	AA110909
1127 G	9 021WA1FH	ESTs {IMAGE:574069}	574069 IMAGE Consortium	AA119074
1128 G	11 021WA1FH	ESTs {IMAGE:574241}	574241 IMAGE Consortium	AA111232
1129 A	1 0213A1FI	A disintegrin and metalloprotease domain (ADAM) 8 {IMAGE:582054}	582054 IMAGE Consortium	AA154162
1130 A	3 0213A1FI	ESTs {IMAGE:578345}	578345 IMAGE Consortium	AA120130
1131 A	5 0213A1FI	ESTs {IMAGE:579699}	579699 IMAGE Consortium	AA116727
1132 A	7 0213A1FI	Mus musculus aldehyde dehydrogenase (ALDH2) mRNA, nuclear gene encoding mitochondrial protein, complete cds {IMAGE:579570}	579570 IMAGE Consortium	AA122975
1133 A	9 0213A1FI	ESTs {IMAGE:579998}	579998 IMAGE Consortium	AA137535
1134 A	11 0213A1FI	IMAGE EST {IMAGE:580142}	580142 IMAGE Consortium	AA137967
1135 C	1 0213A1FI	IMAGE EST {IMAGE:580166}	580166 IMAGE Consortium	AA138018
1136 C	3 0213A1FI	IMAGE EST {IMAGE:580810}	580810 IMAGE Consortium	AA146027

1137 C	5 0213A1FI	IMAGE EST {IMAGE:580945}	580945 IMAGE Consortium	AA139826
1138 C	7 0213A1FI	Mus musculus developmentally regulated repeat (Dr-3), mRNA {IMAGE:581169}	581169 IMAGE Consortium	AA139621
1139 C	9 0213A1FI	IMAGE EST {IMAGE:577601}	577601 IMAGE Consortium	AA138052
1140 C	11 0213A1FI	IMAGE EST {IMAGE:577820}	577820 IMAGE Consortium	AA116221
1141 E	1 0213A1FI	IMAGE EST {IMAGE:577488}	577488 IMAGE Consortium	AA139205
1142 E	3 0213A1FI	IMAGE EST {IMAGE:577592}	577592 IMAGE Consortium	AA139270
1143 E	5 0213A1FI	ESTs {IMAGE:577739}	577739 IMAGE Consortium	AA139223
1144 E	7 0213A1FI	ESTs {IMAGE:577903}	577903 IMAGE Consortium	AA122863
1145 E	9 0213A1FI	ESTs {IMAGE:577844}	577844 IMAGE Consortium	AA116261
1146 E	11 0213A1FI	ESTs {IMAGE:577276}	577276 IMAGE Consortium	AA183727
1147 G	1 0213A1FI	Carbonic anhydrase 2 {IMAGE:579391}	579391 IMAGE Consortium	AA122925
1148 G	3 0213A1FI	Glyceraldehyde-3-phosphate dehydrogenase {IMAGE:579715}	579715 IMAGE Consortium	AA122891
1149 G	5 0213A1FI	ESTs, Weakly similar to DIHA [D.melanogaster] {IMAGE:580000}	580000 IMAGE Consortium	AA122471
1150 G	7 0213A1FI	ESTs {IMAGE:577109}	577109 IMAGE Consortium	AA124476
1151 G	9 0213A1FI	IMAGE EST {IMAGE:580763}	580763 IMAGE Consortium	AA145877
1152 G	11 0213A1FI	ESTs, Highly similar to PROBABLE PEPTIDYL-PROLYL CIS-TRANS ISOMERASE C21E11.05C [Schizosaccharomyces pombe] {IMAGE:577501}	577501 IMAGE Consortium	AA139313
1153 A	1 021AA1FJ	ESTs, Highly similar to CLATHRIN COAT ASSEMBLY PROTEIN AP47 HOMOLOG 1 [R.norvegicus] {IMAGE:573648}	573648 IMAGE Consortium	AA119389
1154 A	3 021AA1FJ	IMAGE EST {IMAGE:573633}	573633 IMAGE Consortium	AA118891
1155 A	5 021AA1FJ	IMAGE EST {IMAGE:573808}	573808 IMAGE Consortium	AA110902
1156 A	7 021AA1FJ	IMAGE EST {IMAGE:575502}	575502 IMAGE Consortium	AA125004
1157 A	9 021AA1FJ	ESTs {IMAGE:574285}	574285 IMAGE Consortium	AA117657
1158 A	11 021AA1FJ	IMAGE EST {IMAGE:574348}	574348 IMAGE Consortium	AA119538
1159 C	1 021AA1FJ	IMAGE EST {IMAGE:573577}	573577 IMAGE Consortium	AA119373
1160 C	3 021AA1FJ	D-E-A-D (aspartate-glutamate-alanine-aspartate) box polypeptide 5 {IMAGE:573801}	573801 IMAGE Consortium	AA118976
1161 C	5 021AA1FJ	Ceroid lipofuscinosis, neuronal 3, juvenile (Batten, Spielmeyer-Vogt disease) {IMAGE:575303}	575303 IMAGE Consortium	AA119043
1162 C	7 021AA1FJ	Myeloblastosis oncogene {IMAGE:575702}	575702 IMAGE Consortium	AA123009
1163 C	9 021AA1FJ	HEMATOPOIETIC CELL PROTEIN-TYROSINE PHOSPHATASE 70Z-PEP {IMAGE:574608}	574608 IMAGE Consortium	AA124804
1164 C	11 021AA1FJ	ESTs, Highly similar to L-SERINE DEHYDRATASE [Rattus norvegicus] {IMAGE:574588}	574588 IMAGE Consortium	AA124751
1165 E	1 021AA1FJ	IMAGE EST {IMAGE:573993}	573993 IMAGE Consortium	AA118985
1166 E	3 021AA1FJ	T-COMPLEX PROTEIN 1, EPSILON SUBUNIT {IMAGE:576003}	576003 IMAGE Consortium	AA120241
1167 E	5 021AA1FJ	ESTs {IMAGE:574379}	574379 IMAGE Consortium	AA119653
1168 E	7 021AA1FJ	IMAGE EST {IMAGE:574625}	574625 IMAGE Consortium	AA118383

1169 E	9 021AA1FJ	IMAGE EST {IMAGE:574832}	574832 IMAGE Consortium	AA124811
1170 E	11 021AA1FJ	CD39 antigen-like 4 {IMAGE:574894}	574894 IMAGE Consortium	AA120757
1171 G	1 021AA1FJ	IMAGE EST {IMAGE:576089}	576089 IMAGE Consortium	AA124508
1172 G	3 021AA1FJ	ESTs {IMAGE:574476}	574476 IMAGE Consortium	AA125030
1173 G	5 021AA1FJ	IMAGE EST {IMAGE:574411}	574411 IMAGE Consortium	AA118284
1174 G	7 021AA1FJ	ESTs {IMAGE:574802}	574802 IMAGE Consortium	AA124495
1175 G	9 021AA1FJ	PROTEIN PHOSPHATASE 2C ALPHA ISOFORM {IMAGE:574952}	574952 IMAGE Consortium	AA125276
1176 G	11 021AA1FJ	IMAGE EST {IMAGE:574984}	574984 IMAGE Consortium	AA118266
1177 A	1 021HA1FK	IMAGE EST {IMAGE:582000}	582000 IMAGE Consortium	AA153001
1178 A	3 021HA1FK	IMAGE EST {IMAGE:581717}	581717 IMAGE Consortium	AA152639
1179 A	5 021HA1FK	IMAGE EST {IMAGE:581893}	581893 IMAGE Consortium	AA152763
1180 A	7 021HA1FK	IMAGE EST {IMAGE:581950}	581950 IMAGE Consortium	AA153285
1181 A	9 021HA1FK	ESTs, Highly similar to similar to human DNA-binding protein 5. [H.sapiens] {IMAGE:582135}	582135 IMAGE Consortium	AA154465
1182 A	11 021HA1FK	IMAGE EST {IMAGE:582535}	582535 IMAGE Consortium	AA153116
1183 C	1 021HA1FK	IMAGE EST {IMAGE:578097}	578097 IMAGE Consortium	AA124744
1184 C	3 021HA1FK	ESTs {IMAGE:577846}	577846 IMAGE Consortium	AA116262
1185 C	5 021HA1FK	IMAGE EST {IMAGE:582067}	582067 IMAGE Consortium	AA145773
1186 C	7 021HA1FK	ESTs {IMAGE:582984}	582984 IMAGE Consortium	AA153685
1187 C	9 021HA1FK	IMAGE EST {IMAGE:577146}	577146 IMAGE Consortium	AA116532
1188 C	11 021HA1FK	IMAGE EST {IMAGE:581069}	581069 IMAGE Consortium	AA139948
1189 E	1 021HA1FK	ESTs, Highly similar to PROBABLE CALCIUM-BINDING PROTEIN PMP41 [Mus musculus] {IMAGE:578399}	578399 IMAGE Consortium	AA116237
1190 E	3 021HA1FK	Epoxide hydrolase 2, cytoplasmic {IMAGE:579349}	579349 IMAGE Consortium	AA124202
1191 E	5 021HA1FK	Kidney androgen regulated protein {IMAGE:579721}	579721 IMAGE Consortium	AA116857
1192 E	7 021HA1FK	IMAGE EST {IMAGE:580257}	580257 IMAGE Consortium	AA137369
1193 E	9 021HA1FK	Mouse mRNA for ribosomal protein S12 {IMAGE:577267}	577267 IMAGE Consortium	AA116548
1194 E	11 021HA1FK	ESTs {IMAGE:577498}	577498 IMAGE Consortium	AA139308
1195 G	1 021HA1FK	ESTs {IMAGE:577594}	577594 IMAGE Consortium	AA139271
1196 G	3 021HA1FK	ESTs {IMAGE:581906}	581906 IMAGE Consortium	AA154452
1197 G	5 021HA1FK	Mus musculus Supl15h gene {IMAGE:582028}	582028 IMAGE Consortium	AA154156
1198 G	7 021HA1FK	IMAGE EST {IMAGE:578439}	578439 IMAGE Consortium	AA123363
1199 G	9 021HA1FK	Imprinted polyspecific membrane transporter-like gene 1 {IMAGE:579339}	579339 IMAGE Consortium	AA122896
1200 G	11 021HA1FK	ESTs {IMAGE:579399}	579399 IMAGE Consortium	AA124250
1201 A	1 021OA1FL	ESTs {IMAGE:582692}	582692 IMAGE Consortium	AA144144
1202 A	3 021OA1FL	IMAGE EST {IMAGE:597035}	597035 IMAGE Consortium	AA145508
1203 A	5 021OA1FL	IMAGE EST {IMAGE:597118}	597118 IMAGE Consortium	AA139063
1204 A	7 021OA1FL	ESTs, Weakly similar to HYPOTHETICAL 38.5 KD PROTEIN IN SUI2-TDH2 INTERGENIC REGION [Saccharomyces cerevisiae] {IMAGE:597416}	597416 IMAGE Consortium	AA153206
1205 A	9 021OA1FL	IMAGE EST {IMAGE:597711}	597711 IMAGE Consortium	AA154089

1206 A	11 021OA1FL	IMAGE EST {IMAGE:597668}	597668 IMAGE Consortium	AA144421
1207 C	1 021OA1FL	ESTs {IMAGE:596976}	596976 IMAGE Consortium	AA145316
1208 C	3 021OA1FL	ESTs {IMAGE:596754}	596754 IMAGE Consortium	AA138332
1209 C	5 021OA1FL	ESTs {IMAGE:583256}	583256 IMAGE Consortium	AA154888
1210 C	7 021OA1FL	Uroporphyrinogen III synthase {IMAGE:583476}	583476 IMAGE Consortium	AA142673
1211 C	9 021OA1FL	IMAGE EST {IMAGE:583753}	583753 IMAGE Consortium	AA153012
1212 C	11 021OA1FL	IMAGE EST {IMAGE:583831}	583831 IMAGE Consortium	AA152954
1213 E	1 021OA1FL	IMAGE EST {IMAGE:596361}	596361 IMAGE Consortium	AA138862
1214 E	3 021OA1FL	Mus musculus aldehyde dehydrogenase Ahd-2-like mRNA, complete cds {IMAGE:596348}	596348 IMAGE Consortium	AA122814
1215 E	5 021OA1FL	IMAGE EST {IMAGE:596665}	596665 IMAGE Consortium	AA146510
1216 E	7 021OA1FL	IMAGE EST {IMAGE:596965}	596965 IMAGE Consortium	AA145398
1217 E	9 021OA1FL	IMAGE EST {IMAGE:597012}	597012 IMAGE Consortium	AA145367
1218 E	11 021OA1FL	IMAGE EST {IMAGE:597095}	597095 IMAGE Consortium	AA138966
1219 G	1 021OA1FL	ESTs {IMAGE:596243}	596243 IMAGE Consortium	AA138786
1220 G	3 021OA1FL	ESTs, Weakly similar to Maxp1 [R.norvegicus] {IMAGE:598598}	598598 IMAGE Consortium	AA161846
1221 G	5 021OA1FL	IMAGE EST {IMAGE:596678}	596678 IMAGE Consortium	AA146282
1222 G	7 021OA1FL	IMAGE EST {IMAGE:596971}	596971 IMAGE Consortium	AA138835
1223 G	9 021OA1FL	Mus musculus Major Histocompatibility Locus class II region {IMAGE:597054}	597054 IMAGE Consortium	AA145469
1224 G	11 021OA1FL	IMAGE EST {IMAGE:597037}	597037 IMAGE Consortium	AA145509
1225 A	1 021VA1FM	Mus musculus radio-resistance/chemo-resistance/cell cycle checkpoint control protein (Rad9) mRNA, complete cds {IMAGE:583606}	583606 IMAGE Consortium	AA146142
1226 A	3 021VA1FM	IMAGE EST {IMAGE:598440}	598440 IMAGE Consortium	AA162169
1227 A	5 021VA1FM	ESTs {IMAGE:598175}	598175 IMAGE Consortium	AA163717
1228 A	7 021VA1FM	ESTs {IMAGE:596863}	596863 IMAGE Consortium	AA145136
1229 A	9 021VA1FM	IMAGE EST {IMAGE:598956}	598956 IMAGE Consortium	AA168646
1230 A	11 021VA1FM	ESTs {IMAGE:616786}	616786 IMAGE Consortium	AA174594
1231 C	1 021VA1FM	ESTs, Weakly similar to SKD1 PROTEIN [Mus musculus] {IMAGE:596358}	596358 IMAGE Consortium	AA138500
1232 C	3 021VA1FM	Cyclin E {IMAGE:598832}	598832 IMAGE Consortium	AA168532
1233 C	5 021VA1FM	Mouse protein disulfide isomerase-related protein (ERp72) mRNA, complete cds {IMAGE:599044}	599044 IMAGE Consortium	AA168956
1234 C	7 021VA1FM	ESTs {IMAGE:598983}	598983 IMAGE Consortium	AA168656
1235 C	9 021VA1FM	ESTs {IMAGE:597201}	597201 IMAGE Consortium	AA145326
1236 C	11 021VA1FM	ESTs {IMAGE:583522}	583522 IMAGE Consortium	AA142685
1237 E	1 021VA1FM	IMAGE EST {IMAGE:596230}	596230 IMAGE Consortium	AA122440
1238 E	3 021VA1FM	Testis specific protein kinase 1 {IMAGE:598713}	598713 IMAGE Consortium	AA163997
1239 E	5 021VA1FM	ESTs, Moderately similar to HYPOTHETICAL PROTEIN KIAA0081 [H.sapiens] {IMAGE:598674}	598674 IMAGE Consortium	AA163274
1240 E	7 021VA1FM	IMAGE EST {IMAGE:599031}	599031 IMAGE Consortium	AA172822

1241 E	9	021VA1FM	ESTs {IMAGE:616706}	616706 IMAGE Consortium	AA174289
1242 E	11	021VA1FM	ESTs {IMAGE:597164}	597164 IMAGE Consortium	AA139041
1243 G	1	021VA1FM	IMAGE EST {IMAGE:596478}	596478 IMAGE Consortium	AA153793
1244 G	3	021VA1FM	Mus musculus mRNA for origin recognition complex subunit 4 {IMAGE:598833}	598833 IMAGE Consortium	AA168536
1245 G	5	021VA1FM	ESTs {IMAGE:599069}	599069 IMAGE Consortium	AA168967
1246 G	7	021VA1FM	ESTs {IMAGE:599055}	599055 IMAGE Consortium	AA172847
1247 G	9	021VA1FM	ESTs {IMAGE:616669}	616669 IMAGE Consortium	AA174857
1248 G	11	021VA1FM	IMAGE EST {IMAGE:583632}	583632 IMAGE Consortium	AA143993
1249 A	1	0212A1FN	IMAGE EST {IMAGE:598387}	598387 IMAGE Consortium	AA155021
1250 A	3	0212A1FN	Mus musculus heterogenous nuclear ribonucleoprotein A2/B1 (hnRNP A2/B1) mRNA, complete cds {IMAGE:598779}	598779 IMAGE Consortium	AA168387
1251 A	5	0212A1FN	ESTs, Moderately similar to probable transcription regulator NT fin12 [M.musculus] {IMAGE:598718}	598718 IMAGE Consortium	AA163996
1252 A	7	0212A1FN	ESTs {IMAGE:599024}	599024 IMAGE Consortium	AA172774
1253 A	9	0212A1FN	IMAGE EST {IMAGE:599058}	599058 IMAGE Consortium	AA162244
1254 A	11	0212A1FN	ESTs {IMAGE:616831}	616831 IMAGE Consortium	AA174503
1255 C	1	0212A1FN	IMAGE EST {IMAGE:598157}	598157 IMAGE Consortium	AA161805
1256 C	3	0212A1FN	ESTs, Weakly similar to FOCAL ADHESION KINASE [Homo sapiens] {IMAGE:598214}	598214 IMAGE Consortium	AA162543
1257 C	5	0212A1FN	IMAGE EST {IMAGE:598411}	598411 IMAGE Consortium	AA155071
1258 C	7	0212A1FN	B-CELL ANTIGEN RECEPTOR COMPLEX ASSOCIATED PROTEIN BETA-CHAIN PRECURSOR {IMAGE:596470}	596470 IMAGE Consortium	AA122794
1259 C	9	0212A1FN	ESTs {IMAGE:596620}	596620 IMAGE Consortium	AA146494
1260 C	11	0212A1FN	IMAGE EST {IMAGE:596637}	596637 IMAGE Consortium	AA138267
1261 E	1	0212A1FN	IMAGE EST {IMAGE:597959}	597959 IMAGE Consortium	AA163668
1262 E	3	0212A1FN	Mus musculus TNF-related apoptosis inducing ligand TRAIL mRNA, complete cds {IMAGE:598349}	598349 IMAGE Consortium	AA163835
1263 E	5	0212A1FN	ESTs {IMAGE:598454}	598454 IMAGE Consortium	AA161771
1264 E	7	0212A1FN	ESTs {IMAGE:598244}	598244 IMAGE Consortium	AA163735
1265 E	9	0212A1FN	ESTs {IMAGE:598566}	598566 IMAGE Consortium	AA162564
1266 E	11	0212A1FN	ESTs {IMAGE:598696}	598696 IMAGE Consortium	AA163456
1267 G	1	0212A1FN	ESTs, Moderately similar to hypothetical protein [M.musculus] {IMAGE:598090}	598090 IMAGE Consortium	AA266639
1268 G	3	0212A1FN	ESTs, Moderately similar to ORF [H.sapiens] {IMAGE:595881}	595881 IMAGE Consortium	AA138086
1269 G	5	0212A1FN	ESTs {IMAGE:598479}	598479 IMAGE Consortium	AA161823
1270 G	7	0212A1FN	IG ALPHA CHAIN C REGION {IMAGE:596604}	596604 IMAGE Consortium	AA146478
1271 G	9	0212A1FN	ESTs {IMAGE:596351}	596351 IMAGE Consortium	AA122819
1272 G	11	0212A1FN	ESTs, Highly similar to ZINC FINGER PROTEIN 131 [H.sapiens] {IMAGE:596767}	596767 IMAGE Consortium	AA146544
1273 A	1	0219A1FO	ESTs {IMAGE:598456}	598456 IMAGE Consortium	AA163309
1274 A	3	0219A1FO	IMAGE EST {IMAGE:596562}	596562 IMAGE Consortium	AA138315

1275 A	5 0219A1FO	IMAGE EST {IMAGE:598842}	598842 IMAGE Consortium	AA162777
1276 A	7 0219A1FO	ESTs {IMAGE:599093}	599093 IMAGE Consortium	AA169001
1277 A	9 0219A1FO	ESTs, Weakly similar to similar to beta transducin proteins containing TRP-ASP domains [C.elegans] {IMAGE:599127}	599127 IMAGE Consortium	AA154732
1278 A	11 0219A1FO	CD20 antigen {IMAGE:616709}	616709 IMAGE Consortium	AA174292
1279 C	1 0219A1FO	IMAGE EST {IMAGE:618446}	618446 IMAGE Consortium	AA175361
1280 C	3 0219A1FO	ESTs {IMAGE:620452}	620452 IMAGE Consortium	AA178118
1281 C	5 0219A1FO	IG ALPHA CHAIN C REGION {IMAGE:620655}	620655 IMAGE Consortium	AA177218
1282 C	7 0219A1FO	ESTs {IMAGE:619000}	619000 IMAGE Consortium	AA170604
1283 C	9 0219A1FO	IMAGE EST {IMAGE:617222}	617222 IMAGE Consortium	AA174781
1284 C	11 0219A1FO	ESTs, Highly similar to NUCLEOPROTEIN TPR [H.sapiens] {IMAGE:619482}	619482 IMAGE Consortium	AA172597
1285 E	1 0219A1FO	ESTs {IMAGE:621006}	621006 IMAGE Consortium	AA178446
1286 E	3 0219A1FO	ESTs {IMAGE:620870}	620870 IMAGE Consortium	AA178361
1287 E	5 0219A1FO	Interleukin 7 receptor {IMAGE:619392}	619392 IMAGE Consortium	AA175538
1288 E	7 0219A1FO	ESTs {IMAGE:617402}	617402 IMAGE Consortium	AA174434
1289 E	9 0219A1FO	IMAGE EST {IMAGE:617552}	617552 IMAGE Consortium	AA174867
1290 E	11 0219A1FO	Interleukin 3 receptor, beta chain 2 {IMAGE:617370}	617370 IMAGE Consortium	AA170386
1291 G	1 0219A1FO	Signal transducer and activator of transcription 4 {IMAGE:618949}	618949 IMAGE Consortium	AA175477
1292 G	3 0219A1FO	IMAGE EST {IMAGE:616771}	616771 IMAGE Consortium	AA170506
1293 G	5 0219A1FO	ESTs {IMAGE:619632}	619632 IMAGE Consortium	AA175784
1294 G	7 0219A1FO	ESTs {IMAGE:619483}	619483 IMAGE Consortium	AA172601
1295 G	9 0219A1FO	ESTs, Weakly similar to hairless protein [M.musculus] {IMAGE:617514}	617514 IMAGE Consortium	AA170351
1296 G	11 0219A1FO	ESTs {IMAGE:619718}	619718 IMAGE Consortium	AA175808
1297 A	1 021GA1FP	IMAGE EST {IMAGE:619117}	619117 IMAGE Consortium	AA175501
1298 A	3 021GA1FP	ESTs, Weakly similar to predicted using Genefinder [C.elegans] {IMAGE:617247}	617247 IMAGE Consortium	AA174789
1299 A	5 021GA1FP	ESTs {IMAGE:617467}	617467 IMAGE Consortium	AA170481
1300 A	7 021GA1FP	IMAGE EST {IMAGE:618052}	618052 IMAGE Consortium	AA175044
1301 A	9 021GA1FP	IMAGE EST {IMAGE:617919}	617919 IMAGE Consortium	AA175422
1302 A	11 021GA1FP	ESTs {IMAGE:620075}	620075 IMAGE Consortium	AA175880
1303 C	1 021GA1FP	ESTs {IMAGE:617500}	617500 IMAGE Consortium	AA174590
1304 C	3 021GA1FP	Myeloid/lymphoid or mixed-lineage leukemia {IMAGE:617388}	617388 IMAGE Consortium	AA174521
1305 C	5 021GA1FP	IMAGE EST {IMAGE:619809}	619809 IMAGE Consortium	AA177856
1306 C	7 021GA1FP	IMAGE EST {IMAGE:619934}	619934 IMAGE Consortium	AA178003
1307 C	9 021GA1FP	ESTs {IMAGE:619799}	619799 IMAGE Consortium	AA172929
1308 C	11 021GA1FP	ESTs {IMAGE:620189}	620189 IMAGE Consortium	AA177914
1309 E	1 021GA1FP	ESTs, Highly similar to SON PROTEIN [Homo sapiens] {IMAGE:619100}	619100 IMAGE Consortium	AA170564
1310 E	3 021GA1FP	ESTs {IMAGE:619418}	619418 IMAGE Consortium	AA175547



1311 E	5	021GA1FP	IMAGE EST {IMAGE:617724}	617724	IMAGE Consortium	AA174910
1312 E	7	021GA1FP	ESTs {IMAGE:617814}	617814	IMAGE Consortium	AA175102
1313 E	9	021GA1FP	IMAGE EST {IMAGE:617944}	617944	IMAGE Consortium	AA170323
1314 E	11	021GA1FP	CD79a antigen {IMAGE:618268}	618268	IMAGE Consortium	AA175325
1315 G	1	021GA1FP	ESTs {IMAGE:617050}	617050	IMAGE Consortium	AA184088
1316 G	3	021GA1FP	ESTs, Highly similar to HMG-box containing protein 1 [R.norvegicus] {IMAGE:619406}	619406	IMAGE Consortium	AA175582
1317 G	5	021GA1FP	ESTs {IMAGE:619728}	619728	IMAGE Consortium	AA175984
1318 G	7	021GA1FP	ESTs {IMAGE:619710}	619710	IMAGE Consortium	AA175980
1319 G	9	021GA1FP	IMAGE EST {IMAGE:617992}	617992	IMAGE Consortium	AA170335
1320 G	11	021GA1FP	MAD homolog 1 (Drosophila) {IMAGE:620333}	620333	IMAGE Consortium	AA177814
1321 A	1	021NA1FQ	Mus musculus GTPase-activating protein GAP111 mRNA, complete cds {IMAGE:622459}	622459	IMAGE Consortium	AA177361
1322 A	3	021NA1FQ	IMAGE EST {IMAGE:636082}	636082	IMAGE Consortium	AA189450
1323 A	5	021NA1FQ	IMAGE EST {IMAGE:636361}	636361	IMAGE Consortium	AA183016
1324 A	7	021NA1FQ	Mus musculus C57BL/6J Sec61 protein complex gamma subunit mRNA, complete cds {IMAGE:621483}	621483	IMAGE Consortium	AA178745
1325 A	9	021NA1FQ	IMAGE EST {IMAGE:621725}	621725	IMAGE Consortium	AA183901
1326 A	11	021NA1FQ	ESTs {IMAGE:621784}	621784	IMAGE Consortium	AA177339
1327 C	1	021NA1FQ	IMAGE EST {IMAGE:621353}	621353	IMAGE Consortium	AA177330
1328 C	3	021NA1FQ	IMAGE EST {IMAGE:621630}	621630	IMAGE Consortium	AA183684
1329 C	5	021NA1FQ	ESTs {IMAGE:621952}	621952	IMAGE Consortium	AA177431
1330 C	7	021NA1FQ	ESTs {IMAGE:634681}	634681	IMAGE Consortium	AA154787
1331 C	9	021NA1FQ	IMAGE EST {IMAGE:622182}	622182	IMAGE Consortium	AA183431
1332 C	11	021NA1FQ	Interleukin 1 receptor, type I {IMAGE:622288}	622288	IMAGE Consortium	AA177717
1333 E	1	021NA1FQ	Janus kinase 2 {IMAGE:621226}	621226	IMAGE Consortium	AA178495
1334 E	3	021NA1FQ	IMAGE EST {IMAGE:634549}	634549	IMAGE Consortium	AA186042
1335 E	5	021NA1FQ	IMAGE EST {IMAGE:621742}	621742	IMAGE Consortium	AA177584
1336 E	7	021NA1FQ	ESTs {IMAGE:622036}	622036	IMAGE Consortium	AA183831
1337 E	9	021NA1FQ	ESTs {IMAGE:621991}	621991	IMAGE Consortium	AA183803
1338 E	11	021NA1FQ	ESTs {IMAGE:622574}	622574	IMAGE Consortium	AA184309
1339 G	1	021NA1FQ	IMAGE EST {IMAGE:634739}	634739	IMAGE Consortium	AA163173
1340 G	3	021NA1FQ	IMAGE EST {IMAGE:634962}	634962	IMAGE Consortium	AA155444
1341 G	5	021NA1FQ	IMAGE EST {IMAGE:635240}	635240	IMAGE Consortium	AA189716
1342 G	7	021NA1FQ	ESTs {IMAGE:635299}	635299	IMAGE Consortium	AA189781
1343 G	9	021NA1FQ	ESTs {IMAGE:622748}	622748	IMAGE Consortium	AA177611
1344 G	11	021NA1FQ	ESTs {IMAGE:634590}	634590	IMAGE Consortium	AA185940
1345 A	1	021UA1FR	IMAGE EST {IMAGE:617290}	617290	IMAGE Consortium	AA174313
1346 A	3	021UA1FR	ESTs {IMAGE:617010}	617010	IMAGE Consortium	AA163405
1347 A	5	021UA1FR	Forkhead homolog 14, (Drosophila) {IMAGE:619776}	619776	IMAGE Consortium	AA176045
1348 A	7	021UA1FR	IMAGE EST {IMAGE:619830}	619830	IMAGE Consortium	AA176061
1349 A	9	021UA1FR	IMAGE EST {IMAGE:619931}	619931	IMAGE Consortium	AA178005
1350 A	11	021UA1FR	ESTs, Highly similar to MYOSIN HEAVY CHAIN, CARDIAC	620239	IMAGE Consortium	AA178015

		MUSCLE BETA ISOFORM [Homo sapiens] {IMAGE:620239}		
1351 C	1	021UA1FR ESTs {IMAGE:619527}	619527	IMAGE Consortium AA175664
1352 C	3	021UA1FR IMAGE EST {IMAGE:617701}	617701	IMAGE Consortium AA174906
1353 C	5	021UA1FR IMAGE EST {IMAGE:617736}	617736	IMAGE Consortium AA175116
1354 C	7	021UA1FR ESTs {IMAGE:617744}	617744	IMAGE Consortium AA175203
1355 C	9	021UA1FR ESTs {IMAGE:620311}	620311	IMAGE Consortium AA177968
1356 C	11	021UA1FR ESTs {IMAGE:618281}	618281	IMAGE Consortium AA170595
1357 E	1	021UA1FR ESTs {IMAGE:617359}	617359	IMAGE Consortium AA174428
1358 E	3	021UA1FR IMAGE EST {IMAGE:617462}	617462	IMAGE Consortium AA174702
1359 E	5	021UA1FR ESTs {IMAGE:617976}	617976	IMAGE Consortium AA175845
1360 E	7	021UA1FR ESTs, Highly similar to COMPLEMENT C1R COMPONENT PRECURSOR [Homo sapiens] {IMAGE:617816}	617816	IMAGE Consortium AA175226
1361 E	9	021UA1FR IMAGE EST {IMAGE:618155}	618155	IMAGE Consortium AA175346
1362 E	11	021UA1FR EST {IMAGE:618401}	618401	IMAGE Consortium AA170693
1363 G	1	021UA1FR IMAGE EST {IMAGE:620240}	620240	IMAGE Consortium AA178066
1364 G	3	021UA1FR ESTs {IMAGE:620547}	620547	IMAGE Consortium AA178286
1365 G	5	021UA1FR ESTs {IMAGE:620932}	620932	IMAGE Consortium AA177178
1366 G	7	021UA1FR ESTs {IMAGE:621068}	621068	IMAGE Consortium AA178142
1367 G	9	021UA1FR IMAGE EST {IMAGE:634106}	634106	IMAGE Consortium AA184612
1368 G	11	021UA1FR IMAGE EST {IMAGE:621352}	621352	IMAGE Consortium AA177326
1369 A	1	0211A1FS ESTs {IMAGE:622762}	622762	IMAGE Consortium AA184259
1370 A	3	0211A1FS IMAGE EST {IMAGE:622893}	622893	IMAGE Consortium AA177702
1371 A	5	0211A1FS Mus musculus mRNA for Rab33B, complete cds {IMAGE:621389}	621389	IMAGE Consortium AA178463
1372 A	7	0211A1FS ESTs {IMAGE:621562}	621562	IMAGE Consortium AA178796
1373 A	9	0211A1FS ESTs, Moderately similar to rtpv-1 [H.sapiens] {IMAGE:634591}	634591	IMAGE Consortium AA185944
1374 A	11	0211A1FS ESTs {IMAGE:634413}	634413	IMAGE Consortium AA185238
1375 C	1	0211A1FS IMAGE EST {IMAGE:621563}	621563	IMAGE Consortium AA178800
1376 C	3	0211A1FS Mus musculus WSB-1 mRNA, complete cds {IMAGE:621575}	621575	IMAGE Consortium AA177596
1377 C	5	0211A1FS ESTs {IMAGE:621715}	621715	IMAGE Consortium AA177791
1378 C	7	0211A1FS ESTs {IMAGE:621815}	621815	IMAGE Consortium AA177352
1379 C	9	0211A1FS ESTs {IMAGE:634790}	634790	IMAGE Consortium AA163180
1380 C	11	0211A1FS Zinc finger protein X-linked {IMAGE:634652}	634652	IMAGE Consortium AA155377
1381 E	1	0211A1FS ESTs {IMAGE:621696}	621696	IMAGE Consortium AA183881
1382 E	3	0211A1FS ESTs {IMAGE:621611}	621611	IMAGE Consortium AA183794
1383 E	5	0211A1FS IMAGE EST {IMAGE:621739}	621739	IMAGE Consortium AA177586
1384 E	7	0211A1FS ESTs {IMAGE:621960}	621960	IMAGE Consortium AA183747
1385 E	9	0211A1FS Growth factor receptor bound protein 2 {IMAGE:622019}	622019	IMAGE Consortium AA183927
1386 E	11	0211A1FS IMAGE EST {IMAGE:622267}	622267	IMAGE Consortium AA177714
1387 G	1	0211A1FS IMAGE EST {IMAGE:621207}	621207	IMAGE Consortium AA178493
1388 G	3	0211A1FS IMAGE EST {IMAGE:634616}	634616	IMAGE Consortium AA185874
1389 G	5	0211A1FS IMAGE EST {IMAGE:634504}	634504	IMAGE Consortium AA185701

1390 G	7 0211A1FS	IMAGE EST {IMAGE:634583}	634583 IMAGE Consortium	AA189649
1391 G	9 0211A1FS	IMAGE EST {IMAGE:634898}	634898 IMAGE Consortium	AA163513
1392 G	11 0211A1FS	IMAGE EST {IMAGE:634671}	634671 IMAGE Consortium	AA155546
1393 A	1 0218A1FT	ESTs {IMAGE:634401}	634401 IMAGE Consortium	AA185446
1394 A	3 0218A1FT	ESTs {IMAGE:634529}	634529 IMAGE Consortium	AA185711
1395 A	5 0218A1FT	ESTs {IMAGE:634632}	634632 IMAGE Consortium	AA154590
1396 A	7 0218A1FT	Lymphocyte antigen 6 complex, locus D {IMAGE:634779}	634779 IMAGE Consortium	AA163336
1397 A	9 0218A1FT	Trans-acting transcription factor 1 {IMAGE:622236}	622236 IMAGE Consortium	AA183548
1398 A	11 0218A1FT	IMAGE EST {IMAGE:622028}	622028 IMAGE Consortium	AA177505
1399 C	1 0218A1FT	ESTs {IMAGE:621820}	621820 IMAGE Consortium	AA184114
1400 C	3 0218A1FT	IMAGE EST {IMAGE:621925}	621925 IMAGE Consortium	AA183712
1401 C	5 0218A1FT	IMAGE EST {IMAGE:621835}	621835 IMAGE Consortium	AA177302
1402 C	7 0218A1FT	IMAGE EST {IMAGE:622226}	622226 IMAGE Consortium	AA183459
1403 C	9 0218A1FT	ESTs {IMAGE:621966}	621966 IMAGE Consortium	AA183749
1404 C	11 0218A1FT	IMAGE EST {IMAGE:634797}	634797 IMAGE Consortium	AA154681
1405 E	1 0218A1FT	ESTs, Weakly similar to T08A11.2 [C.elegans] {IMAGE:639405}	639405 IMAGE Consortium	AA200809
1406 E	3 0218A1FT	Ganglioside expression 2 {IMAGE:637290}	637290 IMAGE Consortium	AA185869
1407 E	5 0218A1FT	ESTs {IMAGE:637444}	637444 IMAGE Consortium	AA185052
1408 E	7 0218A1FT	Histocompatibility 2, Q region locus 7 {IMAGE:637849}	637849 IMAGE Consortium	AA186012
1409 E	9 0218A1FT	Mus musculus mRNA for Ariadne-2 protein {IMAGE:640052}	640052 IMAGE Consortium	AA197417
1410 E	11 0218A1FT	IMAGE EST {IMAGE:640734}	640734 IMAGE Consortium	AA203878
1411 G	1 0218A1FT	ESTs {IMAGE:637891}	637891 IMAGE Consortium	AA189583
1412 G	3 0218A1FT	Cytochrome P450, 2e1, ethanol inducible {IMAGE:640711}	640711 IMAGE Consortium	AA203874
1413 G	5 0218A1FT	ESTs {IMAGE:638302}	638302 IMAGE Consortium	AA200877
1414 G	7 0218A1FT	IMAGE EST {IMAGE:638651}	638651 IMAGE Consortium	AA200144
1415 G	9 0218A1FT	ESTs {IMAGE:641507}	641507 IMAGE Consortium	AA200473
1416 G	11 0218A1FT	ESTs {IMAGE:636308}	636308 IMAGE Consortium	AA182050
1417 A	1 021FA1FU	IMAGE EST {IMAGE:639962}	639962 IMAGE Consortium	AA197427
1418 A	3 021FA1FU	IMAGE EST {IMAGE:639975}	639975 IMAGE Consortium	AA197599
1419 A	5 021FA1FU	IMAGE EST {IMAGE:637882}	637882 IMAGE Consortium	AA189810
1420 A	7 021FA1FU	ESTs {IMAGE:638427}	638427 IMAGE Consortium	AA217016
1421 A	9 021FA1FU	Mus musculus T-cell receptor gamma-chain mRNA, complete cds, clone MNG1 {IMAGE:640763}	640763 IMAGE Consortium	AA203978
1422 A	11 021FA1FU	Growth factor independent 1 {IMAGE:640751}	640751 IMAGE Consortium	AA204603
1423 C	1 021FA1FU	IMAGE EST {IMAGE:637536}	637536 IMAGE Consortium	AA184744
1424 C	3 021FA1FU	IMAGE EST {IMAGE:640177}	640177 IMAGE Consortium	AA197541
1425 C	5 021FA1FU	ESTs {IMAGE:640096}	640096 IMAGE Consortium	AA197454
1426 C	7 021FA1FU	ESTs, Weakly similar to Similarity to B.subtilis tetracycline resistance protein [C.elegans] {IMAGE:640709}	640709 IMAGE Consortium	AA203873
1427 C	9 021FA1FU	ESTs {IMAGE:639220}	639220 IMAGE Consortium	AA200633
1428 C	11 021FA1FU	ESTs {IMAGE:636605}	636605 IMAGE Consortium	AA183294
1429 E	1 021FA1FU	Mus musculus secretory leukoprotease inhibitor mRNA,	638243 IMAGE Consortium	AA200339

		complete cds {IMAGE:638243}		
1430 E	3 021FA1FU	IMAGE EST {IMAGE:638419}	638419 IMAGE Consortium	AA200293
1431 E	5 021FA1FU	Mus musculus I-kappa B alpha chain mRNA, complete cds {IMAGE:641058}	641058 IMAGE Consortium	AA204405
1432 E	7 021FA1FU	ESTs {IMAGE:641524}	641524 IMAGE Consortium	AA200279
1433 E	9 021FA1FU	ESTs {IMAGE:641395}	641395 IMAGE Consortium	AA200392
1434 E	11 021FA1FU	ESTs {IMAGE:639356}	639356 IMAGE Consortium	AA200792
1435 G	1 021FA1FU	ESTs {IMAGE:640875}	640875 IMAGE Consortium	AA210513
1436 G	3 021FA1FU	ESTs {IMAGE:641106}	641106 IMAGE Consortium	AA204468
1437 G	5 021FA1FU	Granzyme B {IMAGE:636781}	636781 IMAGE Consortium	AA183327
1438 G	7 021FA1FU	SH2-B PH domain containing signaling mediator 1 {IMAGE:636884}	636884 IMAGE Consortium	AA189485
1439 G	9 021FA1FU	ESTs {IMAGE:637135}	637135 IMAGE Consortium	AA189690
1440 G	11 021FA1FU	ESTs {IMAGE:639645}	639645 IMAGE Consortium	AA197466
1441 A	1 021MA1FV	IMAGE EST {IMAGE:638819}	638819 IMAGE Consortium	AA200069
1442 A	3 021MA1FV	IMAGE EST {IMAGE:639009}	639009 IMAGE Consortium	AA200176
1443 A	5 021MA1FV	IMAGE EST {IMAGE:636453}	636453 IMAGE Consortium	AA182060
1444 A	7 021MA1FV	ESTs {IMAGE:636921}	636921 IMAGE Consortium	AA189445
1445 A	9 021MA1FV	IMAGE EST {IMAGE:636927}	636927 IMAGE Consortium	AA189448
1446 A	11 021MA1FV	IMAGE EST {IMAGE:639768}	639768 IMAGE Consortium	AA197519
1447 C	1 021MA1FV	ESTs {IMAGE:638783}	638783 IMAGE Consortium	AA199345
1448 C	3 021MA1FV	IMAGE EST {IMAGE:639105}	639105 IMAGE Consortium	AA199573
1449 C	5 021MA1FV	ESTs {IMAGE:636526}	636526 IMAGE Consortium	AA185295
1450 C	7 021MA1FV	Mus musculus ES2 protein (Es2) mRNA, complete cds {IMAGE:636658}	636658 IMAGE Consortium	AA189473
1451 C	9 021MA1FV	IMAGE EST {IMAGE:636640}	636640 IMAGE Consortium	AA204588
1452 C	11 021MA1FV	ESTs {IMAGE:637233}	637233 IMAGE Consortium	AA189900
1453 E	1 021MA1FV	HpaII tiny fragments locus 9c {IMAGE:641594}	641594 IMAGE Consortium	AA209823
1454 E	3 021MA1FV	ESTs {IMAGE:639215}	639215 IMAGE Consortium	AA200989
1455 E	5 021MA1FV	IMAGE EST {IMAGE:639273}	639273 IMAGE Consortium	AA199055
1456 E	7 021MA1FV	ESTs {IMAGE:637202}	637202 IMAGE Consortium	AA189705
1457 E	9 021MA1FV	IMAGE EST {IMAGE:637305}	637305 IMAGE Consortium	AA190040
1458 E	11 021MA1FV	ESTs {IMAGE:639998}	639998 IMAGE Consortium	AA197603
1459 G	1 021MA1FV	Mus musculus FK506-binding protein 25 homolog (mFKBP25) mRNA, partial cds {IMAGE:636888}	636888 IMAGE Consortium	AA182992
1460 G	3 021MA1FV	ESTs {IMAGE:636730}	636730 IMAGE Consortium	AA189425
1461 G	5 021MA1FV	IMAGE EST {IMAGE:639306}	639306 IMAGE Consortium	AA200721
1462 G	7 021MA1FV	ESTs {IMAGE:637007}	637007 IMAGE Consortium	AA185012
1463 G	9 021MA1FV	ESTs {IMAGE:637976}	637976 IMAGE Consortium	AA189999
1464 G	11 021MA1FV	ESTs {IMAGE:640611}	640611 IMAGE Consortium	AA203985
1465 A	1 021TA1FW	IMAGE EST {IMAGE:644533}	644533 IMAGE Consortium	AA208818
1466 A	3 021TA1FW	IMAGE EST {IMAGE:644850}	644850 IMAGE Consortium	AA210272
1467 A	5 021TA1FW	ESTs {IMAGE:652544}	652544 IMAGE Consortium	AA217333

1468 A	7 021TA1FW	IMAGE EST {IMAGE:642299}	642299 IMAGE Consortium	AA212558
1469 A	9 021TA1FW	ESTs {IMAGE:642114}	642114 IMAGE Consortium	AA210196
1470 A	11 021TA1FW	IMAGE EST {IMAGE:642506}	642506 IMAGE Consortium	AA185477
1471 C	1 021TA1FW	ESTs {IMAGE:644320}	644320 IMAGE Consortium	AA204530
1472 C	3 021TA1FW	IMAGE EST {IMAGE:644887}	644887 IMAGE Consortium	AA210606
1473 C	5 021TA1FW	IMAGE EST {IMAGE:641854}	641854 IMAGE Consortium	AA209932
1474 C	7 021TA1FW	IMAGE EST {IMAGE:641936}	641936 IMAGE Consortium	AA210024
1475 C	9 021TA1FW	IMAGE EST {IMAGE:642204}	642204 IMAGE Consortium	AA212239
1476 C	11 021TA1FW	ESTs {IMAGE:652717}	652717 IMAGE Consortium	AA217264
1477 E	1 021TA1FW	IMAGE EST {IMAGE:644371}	644371 IMAGE Consortium	AA204518
1478 E	3 021TA1FW	ESTs {IMAGE:644840}	644840 IMAGE Consortium	AA209544
1479 E	5 021TA1FW	IMAGE EST {IMAGE:644998}	644998 IMAGE Consortium	AA210405
1480 E	7 021TA1FW	ESTs {IMAGE:652506}	652506 IMAGE Consortium	AA217452
1481 E	9 021TA1FW	ESTs {IMAGE:652758}	652758 IMAGE Consortium	AA217199
1482 E	11 021TA1FW	IMAGE EST {IMAGE:653189}	653189 IMAGE Consortium	AA217174
1483 G	1 021TA1FW	ESTs {IMAGE:652498}	652498 IMAGE Consortium	AA217255
1484 G	3 021TA1FW	IMAGE EST {IMAGE:652507}	652507 IMAGE Consortium	AA217455
1485 G	5 021TA1FW	IMAGE EST {IMAGE:652735}	652735 IMAGE Consortium	AA217196
1486 G	7 021TA1FW	ESTs {IMAGE:642394}	642394 IMAGE Consortium	AA184806
1487 G	9 021TA1FW	Mus musculus gamma2-adaptin (G2ad) mRNA, complete cds {IMAGE:642332}	642332 IMAGE Consortium	AA185250
1488 G	11 021TA1FW	ESTs {IMAGE:642836}	642836 IMAGE Consortium	AA185271
1489 A	1 0210A1FX	ESTs {IMAGE:642408}	642408 IMAGE Consortium	AA184879
1490 A	3 0210A1FX	ESTs {IMAGE:642467}	642467 IMAGE Consortium	AA190123
1491 A	5 0210A1FX	ESTs {IMAGE:642549}	642549 IMAGE Consortium	AA184778
1492 A	7 0210A1FX	IMAGE EST {IMAGE:653484}	653484 IMAGE Consortium	AA215268
1493 A	9 0210A1FX	ESTs {IMAGE:643489}	643489 IMAGE Consortium	AA204118
1494 A	11 0210A1FX	Mus musculus formin binding protein 21 mRNA, complete cds {IMAGE:658427}	658427 IMAGE Consortium	AA198329
1495 C	1 0210A1FX	IMAGE EST {IMAGE:643420}	643420 IMAGE Consortium	AA190093
1496 C	3 0210A1FX	ESTs {IMAGE:643095}	643095 IMAGE Consortium	AA189917
1497 C	5 0210A1FX	Mus musculus transcription factor Sp3 mRNA, partial cds {IMAGE:643587}	643587 IMAGE Consortium	AA204627
1498 C	7 0210A1FX	IMAGE EST {IMAGE:658260}	658260 IMAGE Consortium	AA198345
1499 C	9 0210A1FX	ESTs {IMAGE:641698}	641698 IMAGE Consortium	AA209964
1500 C	11 0210A1FX	IMAGE EST {IMAGE:641775}	641775 IMAGE Consortium	AA210085
1501 E	1 0210A1FX	ESTs, Weakly similar to TROPOMYOSIN, FIBROBLAST ISOFORM 2 [M.musculus] {IMAGE:643158}	643158 IMAGE Consortium	AA189695
1502 E	3 0210A1FX	ESTs, Highly similar to nucleoporin Nup84 [R.norvegicus] {IMAGE:643755}	643755 IMAGE Consortium	AA203895
1503 E	5 0210A1FX	ESTs, Weakly similar to HYPOTHETICAL PROTEIN F44G4.1 IN CHROMOSOME II [C.elegans] {IMAGE:658190}	658190 IMAGE Consortium	AA198227
1504 E	7 0210A1FX	ESTs {IMAGE:641746}	641746 IMAGE Consortium	AA210069

1505 E	9 0210A1FX	IMAGE EST {IMAGE:641871}	641871 IMAGE Consortium	AA209884
1506 E	11 0210A1FX	ESTs {IMAGE:642145}	642145 IMAGE Consortium	AA212164
1507 G	1 0210A1FX	ESTs {IMAGE:658334}	658334 IMAGE Consortium	AA198396
1508 G	3 0210A1FX	IMAGE EST {IMAGE:644977}	644977 IMAGE Consortium	AA210357
1509 G	5 0210A1FX	IMAGE EST {IMAGE:644941}	644941 IMAGE Consortium	AA210377
1510 G	7 0210A1FX	ESTs {IMAGE:652515}	652515 IMAGE Consortium	AA217217
1511 G	9 0210A1FX	IMAGE EST {IMAGE:652811}	652811 IMAGE Consortium	AA217625
1512 G	11 0210A1FX	IMAGE EST {IMAGE:653365}	653365 IMAGE Consortium	AA240403
1513 A	1 0217A1FY	IMAGE EST {IMAGE:832632}	832632 IMAGE Consortium	AA423757
1514 A	3 0217A1FY	ESTs, Weakly similar to T13H5.5 [C.elegans] {IMAGE:848702}	848702 IMAGE Consortium	AA445435
1515 A	5 0217A1FY	IMAGE EST {IMAGE:846037}	846037 IMAGE Consortium	AA416235
1516 A	7 0217A1FY	Mus musculus polypeptide GalNAc transferase-T1 (ppGaNTase-T1) mRNA, complete cds {IMAGE:849524}	849524 IMAGE Consortium	AA457927
1517 A	9 0217A1FY	IMAGE EST {IMAGE:850178}	850178 IMAGE Consortium	AA451403
1518 A	11 0217A1FY	IMAGE EST {IMAGE:850093}	850093 IMAGE Consortium	AA450659
1519 C	1 0217A1FY	ESTs, Weakly similar to weak similarity with many protein kinases [C.elegans] {IMAGE:851311}	851311 IMAGE Consortium	AA461641
1520 C	3 0217A1FY	IMAGE EST {IMAGE:851230}	851230 IMAGE Consortium	AA462936
1521 C	5 0217A1FY	Cyclin G {IMAGE:849762}	849762 IMAGE Consortium	AA458196
1522 C	7 0217A1FY	IMAGE EST {IMAGE:850281}	850281 IMAGE Consortium	AA451505
1523 C	9 0217A1FY	Complement component 2 (within H-2S) {IMAGE:851201}	851201 IMAGE Consortium	AA462869
1524 C	11 0217A1FY	ESTs, Moderately similar to metastasis-associated protein mta-1 [R.norvegicus] {IMAGE:851326}	851326 IMAGE Consortium	AA461637
1525 E	1 0217A1FY	ATP-binding cassette 2 {IMAGE:850420}	850420 IMAGE Consortium	AA451270
1526 E	3 0217A1FY	ESTs {IMAGE:850931}	850931 IMAGE Consortium	AA462270
1527 E	5 0217A1FY	Mus musculus CC chemokine ABCD-1 mRNA, complete cds {IMAGE:851752}	851752 IMAGE Consortium	AA522165
1528 E	7 0217A1FY	ESTs {IMAGE:851030}	851030 IMAGE Consortium	AA462524
1529 E	9 0217A1FY	Nuclear factor of activated T-cells, cytoplasmic 2 {IMAGE:850163}	850163 IMAGE Consortium	AA451395
1530 E	11 0217A1FY	Cell division cycle 25A {IMAGE:851054}	851054 IMAGE Consortium	AA462544
1531 G	1 0217A1FY	IMAGE EST {IMAGE:874592}	874592 IMAGE Consortium	AA472426
1532 G	3 0217A1FY	IMAGE EST {IMAGE:871179}	871179 IMAGE Consortium	AA462784
1533 G	5 0217A1FY	ESTs, Moderately similar to PIAS3 [M.musculus] {IMAGE:872973}	872973 IMAGE Consortium	AA466489
1534 G	7 0217A1FY	ESTs, Moderately similar to COMPLEMENT C6 PRECURSOR [H.sapiens] {IMAGE:873972}	873972 IMAGE Consortium	AA472154
1535 G	9 0217A1FY	ESTs {IMAGE:874436}	874436 IMAGE Consortium	AA475037
1536 G	11 0217A1FY	IMAGE EST {IMAGE:872220}	872220 IMAGE Consortium	AA472217
1537 A	1 021EA1FZ	IMAGE EST {IMAGE:642209}	642209 IMAGE Consortium	AA212320
1538 A	3 021EA1FZ	ESTs {IMAGE:642649}	642649 IMAGE Consortium	AA185415
1539 A	5 021EA1FZ	ESTs, Moderately similar to T10C6.i [C.elegans]	642424 IMAGE Consortium	AA189182

		{IMAGE:642424}		
1540 A	7 021EA1FZ	ESTs {IMAGE:642407}	642407 IMAGE Consortium	AA185258
1541 A	9 021EA1FZ	ESTs, Highly similar to zinc finger protein C2H2-171 [H.sapiens] {IMAGE:642975}	642975 IMAGE Consortium	AA185217
1542 A	11 021EA1FZ	ESTs {IMAGE:643183}	643183 IMAGE Consortium	AA185909
1543 C	1 021EA1FZ	IMAGE EST {IMAGE:642250}	642250 IMAGE Consortium	AA212540
1544 C	3 021EA1FZ	IMAGE EST {IMAGE:641994}	641994 IMAGE Consortium	AA210258
1545 C	5 021EA1FZ	ESTs, Highly similar to MYOSIN HEAVY CHAIN, NONMUSCLE [Gallus gallus] {IMAGE:642434}	642434 IMAGE Consortium	AA184722
1546 C	7 021EA1FZ	Mus musculus CTP synthetase mRNA, complete cds {IMAGE:642329}	642329 IMAGE Consortium	AA185252
1547 C	9 021EA1FZ	IMAGE EST {IMAGE:643416}	643416 IMAGE Consortium	AA190091
1548 C	11 021EA1FZ	ESTs {IMAGE:643784}	643784 IMAGE Consortium	AA203922
1549 E	1 021EA1FZ	IMAGE EST {IMAGE:653161}	653161 IMAGE Consortium	AA217113
1550 E	3 021EA1FZ	IMAGE EST {IMAGE:642747}	642747 IMAGE Consortium	AA185127
1551 E	5 021EA1FZ	ESTs {IMAGE:653828}	653828 IMAGE Consortium	AA217718
1552 E	7 021EA1FZ	ESTs {IMAGE:643376}	643376 IMAGE Consortium	AA185931
1553 E	9 021EA1FZ	IMAGE EST {IMAGE:657691}	657691 IMAGE Consortium	AA199150
1554 E	11 021EA1FZ	IMAGE EST {IMAGE:643750}	643750 IMAGE Consortium	AA203779
1555 G	1 021EA1FZ	ESTs, Moderately similar to highly charged protein [H.sapiens] {IMAGE:847303}	847303 IMAGE Consortium	AA420156
1556 G	3 021EA1FZ	IMAGE EST {IMAGE:845839}	845839 IMAGE Consortium	AA416091
1557 G	5 021EA1FZ	ESTs {IMAGE:846560}	846560 IMAGE Consortium	AA433576
1558 G	7 021EA1FZ	ESTs {IMAGE:832614}	832614 IMAGE Consortium	AA423717
1559 G	9 021EA1FZ	ESTs {IMAGE:831647}	831647 IMAGE Consortium	AA466089
1560 G	11 021EA1FZ	IMAGE EST {IMAGE:848652}	848652 IMAGE Consortium	AA445405
1561 A	1 021EA1G0	IMAGE EST {IMAGE:864414}	864414 IMAGE Consortium	AA458171
1562 A	3 021EA1G0	IMAGE EST {IMAGE:874046}	874046 IMAGE Consortium	AA471737
1563 A	5 021EA1G0	ESTs {IMAGE:874419}	874419 IMAGE Consortium	AA475039
1564 A	7 021EA1G0	ESTs {IMAGE:870879}	870879 IMAGE Consortium	AA461727
1565 A	9 021EA1G0	BP-3 alloantigen {IMAGE:871565}	871565 IMAGE Consortium	AA462202
1566 A	11 021EA1G0	IMAGE EST {IMAGE:864295}	864295 IMAGE Consortium	AA458096
1567 C	1 021EA1G0	ESTs, Highly similar to CELLULAR APOPTOSIS SUSCEPTIBILITY PROTEIN [H.sapiens] {IMAGE:874148}	874148 IMAGE Consortium	AA471761
1568 C	3 021EA1G0	Extracellular matrix protein 1 {IMAGE:874833}	874833 IMAGE Consortium	AA474897
1569 C	5 021EA1G0	IMAGE EST {IMAGE:864552}	864552 IMAGE Consortium	AA458273
1570 C	7 021EA1G0	IMAGE EST {IMAGE:864344}	864344 IMAGE Consortium	AA458211
1571 C	9 021EA1G0	ESTs {IMAGE:870815}	870815 IMAGE Consortium	AA462005
1572 C	11 021EA1G0	ESTs, Highly similar to mitogen inducible gene mig-2 [H.sapiens] {IMAGE:864576}	864576 IMAGE Consortium	AA458292
1573 E	1 021EA1G0	Mus musculus ras-related protein (rab18) mRNA, complete cds {IMAGE:874352}	874352 IMAGE Consortium	AA474932
1574 E	3 021EA1G0	ESTs {IMAGE:874554}	874554 IMAGE Consortium	AA472321

1575 E	5 021EA1G0	Monokine induced by gamma interferon {IMAGE:874758}	874758 IMAGE Consortium	AA472492
1576 E	7 021EA1G0	IMAGE EST {IMAGE:871676}	871676 IMAGE Consortium	AA462348
1577 E	9 021EA1G0	IMAGE EST {IMAGE:873873}	873873 IMAGE Consortium	AA472038
1578 E	11 021EA1G0	ESTs, Highly similar to NADH-UBIQUINONE OXIDOREDUCTASE B14 SUBUNIT [Bos taurus] {IMAGE:871020}	871020 IMAGE Consortium	AA462323
1579 G	1 021EA1G0	IMAGE EST {IMAGE:891442}	891442 IMAGE Consortium	AA511060
1580 G	3 021EA1G0	Mus musculus aldo-keto reductase mRNA, complete cds {IMAGE:891261}	891261 IMAGE Consortium	AA510877
1581 G	5 021EA1G0	Mus musculus mRNA for peroxisomal integral membrane protein PMP34 {IMAGE:888563}	888563 IMAGE Consortium	AA498608
1582 G	7 021EA1G0	ESTs {IMAGE:890582}	890582 IMAGE Consortium	AA509736
1583 G	9 021EA1G0	Mus musculus G protein-coupled receptor kinase 5 (GRK5) mRNA, complete cds {IMAGE:875444}	875444 IMAGE Consortium	AA475346
1584 G	11 021EA1G0	ESTs {IMAGE:891203}	891203 IMAGE Consortium	AA510287
1585 A	1 021LA1G1	D-E-A-D (aspartate-glutamate-alanine-aspartate) box polypeptide 5 {IMAGE:876780}	876780 IMAGE Consortium	AA497429
1586 A	3 021LA1G1	Mus musculus translocase of inner mitochondrial membrane (Tim44) mRNA, nuclear gene encoding mitochondrial protein, complete cds {IMAGE:888708}	888708 IMAGE Consortium	AA498767
1587 A	5 021LA1G1	IMAGE EST {IMAGE:889469}	889469 IMAGE Consortium	AA498528
1588 A	7 021LA1G1	ESTs, Highly similar to PI 6.1 ESTERASE PRECURSOR [Rattus norvegicus] {IMAGE:890429}	890429 IMAGE Consortium	AA509566
1589 A	9 021LA1G1	ESTs {IMAGE:875684}	875684 IMAGE Consortium	AA475387
1590 A	11 021LA1G1	ESTs, Weakly similar to LL5 protein [R.norvegicus] {IMAGE:876535}	876535 IMAGE Consortium	AA475992
1591 C	1 021LA1G1	ESTs {IMAGE:902264}	902264 IMAGE Consortium	AA516913
1592 C	3 021LA1G1	IMAGE EST {IMAGE:902385}	902385 IMAGE Consortium	AA517261
1593 C	5 021LA1G1	IMAGE EST {IMAGE:890478}	890478 IMAGE Consortium	AA509851
1594 C	7 021LA1G1	ESTs {IMAGE:876311}	876311 IMAGE Consortium	AA475767
1595 C	9 021LA1G1	ESTs, Highly similar to ELONGATION FACTOR G, MITOCHONDRIAL PRECURSOR [Rattus norvegicus] {IMAGE:889470}	889470 IMAGE Consortium	AA498518
1596 C	11 021LA1G1	ESTs, Weakly similar to VILLIN [Gallus gallus] {IMAGE:875543}	875543 IMAGE Consortium	AA475482
1597 E	1 021LA1G1	Peroxisome proliferator activator receptor gamma {IMAGE:890915}	890915 IMAGE Consortium	AA511029
1598 E	3 021LA1G1	ESTs, Moderately similar to RAF SERINE/THREONINE-PROTEIN KINASE TRANSFORMING PROTEIN [Murine sarcoma virus 3611] {IMAGE:889607}	889607 IMAGE Consortium	AA498185
1599 E	5 021LA1G1	ESTs {IMAGE:890763}	890763 IMAGE Consortium	AA510492
1600 E	7 021LA1G1	ESTs {IMAGE:876257}	876257 IMAGE Consortium	AA476183
1601 E	9 021LA1G1	IMAGE EST {IMAGE:891229}	891229 IMAGE Consortium	AA510303
1602 E	11 021LA1G1	Kruppel zinc finger protein 2 {IMAGE:901909}	901909 IMAGE Consortium	AA517781



1603 G	1 021LA1G1	IMAGE EST {IMAGE:876297}	876297 IMAGE Consortium	AA475761
1604 G	3 021LA1G1	IMAGE EST {IMAGE:891368}	891368 IMAGE Consortium	AA510714
1605 G	5 021LA1G1	ESTs {IMAGE:891325}	891325 IMAGE Consortium	AA510559
1606 G	7 021LA1G1	ESTs {IMAGE:888571}	888571 IMAGE Consortium	AA498612
1607 G	9 021LA1G1	Mus musculus cGMP phosphodiesterase (PDE9A*1) mRNA, complete cds {IMAGE:889753}	889753 IMAGE Consortium	AA498410
1608 G	11 021LA1G1	ESTs {IMAGE:890699}	890699 IMAGE Consortium	AA510000
1609 A	1 021SA1G2	IMAGE EST {IMAGE:949384}	949384 IMAGE Consortium	AA542065
1610 A	3 021SA1G2	Mouse mRNA for plexin 2, complete cds {IMAGE:922991}	922991 IMAGE Consortium	AA511430
1611 A	5 021SA1G2	IMAGE EST {IMAGE:949641}	949641 IMAGE Consortium	AA537301
1612 A	7 021SA1G2	ESTs {IMAGE:935442}	935442 IMAGE Consortium	AA543368
1613 A	9 021SA1G2	ESTs, Moderately similar to brefeldin A-sensitive Golgi protein LDLC [H.sapiens] {IMAGE:904901}	904901 IMAGE Consortium	AA519035
1614 A	11 021SA1G2	ESTs {IMAGE:584079}	584079 IMAGE Consortium	AA125423
1615 C	1 021SA1G2	IMAGE EST {IMAGE:949035}	949035 IMAGE Consortium	AA543904
1616 C	3 021SA1G2	IMAGE EST {IMAGE:949369}	949369 IMAGE Consortium	AA542069
1617 C	5 021SA1G2	ESTs {IMAGE:922312}	922312 IMAGE Consortium	AA512757
1618 C	7 021SA1G2	ESTs, Highly similar to CAMP-DEPENDENT PROTEIN KINASE TYPE I-ALPHA REGULATORY CHAIN [Homo sapiens] {IMAGE:949663}	949663 IMAGE Consortium	AA537355
1619 C	9 021SA1G2	IMAGE EST {IMAGE:935438}	935438 IMAGE Consortium	AA543366
1620 C	11 021SA1G2	IMAGE EST {IMAGE:948564}	948564 IMAGE Consortium	AA544701
1621 E	1 021SA1G2	IMAGE EST {IMAGE:949810}	949810 IMAGE Consortium	AA537509
1622 E	3 021SA1G2	IMAGE EST {IMAGE:904666}	904666 IMAGE Consortium	AA518801
1623 E	5 021SA1G2	IMAGE EST {IMAGE:948845}	948845 IMAGE Consortium	AA544859
1624 E	7 021SA1G2	IMAGE EST {IMAGE:949807}	949807 IMAGE Consortium	AA537519
1625 E	9 021SA1G2	ESTs, Highly similar to ALDEHYDE DEHYDROGENASE, E3 ISOZYME [Homo sapiens] {IMAGE:658785}	658785 IMAGE Consortium	AA221806
1626 E	11 021SA1G2	IMAGE EST {IMAGE:658839}	658839 IMAGE Consortium	AA217892
1627 G	1 021SA1G2	Mus musculus WW domain binding protein 12 mRNA, partial sequence {IMAGE:670890}	670890 IMAGE Consortium	AA250384
1628 G	3 021SA1G2	ESTs {IMAGE:658863}	658863 IMAGE Consortium	AA217899
1629 G	5 021SA1G2	IMAGE EST {IMAGE:659571}	659571 IMAGE Consortium	AA218162
1630 G	7 021SA1G2	IMAGE EST {IMAGE:659857}	659857 IMAGE Consortium	AA219828
1631 G	9 021SA1G2	ESTs {IMAGE:659662}	659662 IMAGE Consortium	AA221184
1632 G	11 021SA1G2	ESTs {IMAGE:660023}	660023 IMAGE Consortium	AA220755
1633 A	1 021ZA1G3	ESTs {IMAGE:902222}	902222 IMAGE Consortium	AA516813
1634 A	3 021ZA1G3	IMAGE EST {IMAGE:889540}	889540 IMAGE Consortium	AA498704
1635 A	5 021ZA1G3	Retinoblastoma-like 2 {IMAGE:890596}	890596 IMAGE Consortium	AA509751
1636 A	7 021ZA1G3	ESTs, Weakly similar to TROPOMYOSIN, FIBROBLAST ISOFORM 2 [M.musculus] {IMAGE:890486}	890486 IMAGE Consortium	AA509855
1637 A	9 021ZA1G3	ESTs {IMAGE:903905}	903905 IMAGE Consortium	AA521755
1638 A	11 021ZA1G3	IMAGE EST {IMAGE:533561}	533561 IMAGE Consortium	AA105066

1639 C	1 021ZA1G3	IMAGE EST {IMAGE:948846}	948846 IMAGE Consortium	AA544849
1640 C	3 021ZA1G3	ESTs {IMAGE:949116}	949116 IMAGE Consortium	AA544654
1641 C	5 021ZA1G3	ESTs, Highly similar to ACETYL-COENZYME A SYNTHETASE [Escherichia coli] {IMAGE:949423}	949423 IMAGE Consortium	AA537637
1642 C	7 021ZA1G3	X-ray repair complementing defective repair in Chinese hamster {IMAGE:949439}	949439 IMAGE Consortium	AA537645
1643 C	9 021ZA1G3	ESTs {IMAGE:935648}	935648 IMAGE Consortium	AA543973
1644 C	11 021ZA1G3	ESTs, Moderately similar to JAK3 [M.musculus] {IMAGE:949603}	949603 IMAGE Consortium	AA537225
1645 E	1 021ZA1G3	ATPase, Na <sup>+</sup> /K <sup>+</sup> transporting, beta 2 polypeptide {IMAGE:935488}	935488 IMAGE Consortium	AA543167
1646 E	3 021ZA1G3	ESTs {IMAGE:515681}	515681 IMAGE Consortium	AA066021
1647 E	5 021ZA1G3	ESTs {IMAGE:948618}	948618 IMAGE Consortium	AA544789
1648 E	7 021ZA1G3	Mus musculus Btk locus, alpha-D-galactosidase A (Ags), ribosomal protein (L44L), and Bruton's tyrosine kinase (Btk) genes, complete cds {IMAGE:920442}	920442 IMAGE Consortium	AA538508
1649 E	9 021ZA1G3	Tumor necrosis factor induced protein 2 {IMAGE:949472}	949472 IMAGE Consortium	AA537672
1650 E	11 021ZA1G3	ESTs {IMAGE:949784}	949784 IMAGE Consortium	AA537490
1651 G	1 021ZA1G3	IMAGE EST {IMAGE:949532}	949532 IMAGE Consortium	AA537116
1652 G	3 021ZA1G3	ESTs {IMAGE:947809}	947809 IMAGE Consortium	AA542113
1653 G	5 021ZA1G3	Mus musculus MAN2B2 mRNA for alpha-D-mannosidase, complete cds {IMAGE:905640}	905640 IMAGE Consortium	AA529219
1654 G	7 021ZA1G3	PEPTIDYL-PROLYL CIS-TRANS ISOMERASE A {IMAGE:920055}	920055 IMAGE Consortium	AA499926
1655 G	9 021ZA1G3	ESTs {IMAGE:920458}	920458 IMAGE Consortium	AA538527
1656 G	11 021ZA1G3	ESTs {IMAGE:922234}	922234 IMAGE Consortium	AA512287
1657 A	1 0216A1G4	ESTs {IMAGE:658666}	658666 IMAGE Consortium	AA221886
1658 A	3 0216A1G4	ESTs {IMAGE:659103}	659103 IMAGE Consortium	AA217774
1659 A	5 0216A1G4	ESTs, Weakly similar to predicted using Genefinder [C.elegans] {IMAGE:659333}	659333 IMAGE Consortium	AA217887
1660 A	7 0216A1G4	IMAGE EST {IMAGE:659860}	659860 IMAGE Consortium	AA219826
1661 A	9 0216A1G4	Mus musculus mRNA for putative sialyltransferase {IMAGE:660287}	660287 IMAGE Consortium	AA220284
1662 A	11 0216A1G4	ESTs, Moderately similar to R151.6 [C.elegans] {IMAGE:660568}	660568 IMAGE Consortium	AA223028
1663 C	1 0216A1G4	ESTs {IMAGE:671265}	671265 IMAGE Consortium	AA222536
1664 C	3 0216A1G4	IMAGE EST {IMAGE:671748}	671748 IMAGE Consortium	AA242209
1665 C	5 0216A1G4	IMAGE EST {IMAGE:672179}	672179 IMAGE Consortium	AA231351
1666 C	7 0216A1G4	ESTs, Weakly similar to Murine homolog of human ftp-3 [M.musculus] {IMAGE:661089}	661089 IMAGE Consortium	AA222738
1667 C	9 0216A1G4	ESTs {IMAGE:669985}	669985 IMAGE Consortium	AA253856
1668 C	11 0216A1G4	ESTs, Highly similar to HYPOTHETICAL TRP-ASP REPEATS CONTAINING PROTEIN C29E6.01 IN CHROMOSOME I [Schizosaccharomyces pombe]	670050 IMAGE Consortium	AA230588

		{IMAGE:670050}		
1669 E	1	0216A1G4	ESTs {IMAGE:660618}	660618 IMAGE Consortium AA222977
1670 E	3	0216A1G4	EST {IMAGE:660947}	660947 IMAGE Consortium AA253775
1671 E	5	0216A1G4	IMAGE EST {IMAGE:660979}	660979 IMAGE Consortium AA230691
1672 E	7	0216A1G4	IMAGE EST {IMAGE:669831}	669831 IMAGE Consortium AA253750
1673 E	9	0216A1G4	ESTs, Moderately similar to protein kinase C-binding protein RACK7 [H.sapiens] {IMAGE:670201}	670201 IMAGE Consortium AA222144
1674 E	11	0216A1G4	ESTs {IMAGE:677128}	677128 IMAGE Consortium AA212937
1675 G	1	0216A1G4	ESTs {IMAGE:676325}	676325 IMAGE Consortium AA208843
1676 G	3	0216A1G4	ESTs, Highly similar to APOLIPOPROTEIN B-100 PRECURSOR [Homo sapiens] {IMAGE:676410}	676410 IMAGE Consortium AA209065
1677 G	5	0216A1G4	IMAGE EST {IMAGE:670202}	670202 IMAGE Consortium AA222141
1678 G	7	0216A1G4	ESTs {IMAGE:676925}	676925 IMAGE Consortium AA212992
1679 G	9	0216A1G4	Heme oxygenase (decycling) 1 {IMAGE:677499}	677499 IMAGE Consortium AA213167
1680 G	11	0216A1G4	IMAGE EST {IMAGE:677417}	677417 IMAGE Consortium AA212808
1681 A	1	021DA1G5	Mus musculus ATP-dependent RNA helicase mRNA, partial cds {IMAGE:676592}	676592 IMAGE Consortium AA212347
1682 A	3	021DA1G5	Mus musculus mRNA for CtBP1 protein {IMAGE:677093}	677093 IMAGE Consortium AA212717
1683 A	5	021DA1G5	ESTs {IMAGE:676891}	676891 IMAGE Consortium AA213337
1684 A	7	021DA1G5	ESTs {IMAGE:670980}	670980 IMAGE Consortium AA240826
1685 A	9	021DA1G5	ESTs, Highly similar to CLATHRIN HEAVY CHAIN [Rattus norvegicus] {IMAGE:659022}	659022 IMAGE Consortium AA217742
1686 A	11	021DA1G5	NUCLEOSOME ASSEMBLY PROTEIN 1-LIKE 1 {IMAGE:671801}	671801 IMAGE Consortium AA242323
1687 C	1	021DA1G5	ESTs, Highly similar to small acidic protein [H.sapiens] {IMAGE:671789}	671789 IMAGE Consortium AA241756
1688 C	3	021DA1G5	IMAGE EST {IMAGE:672245}	672245 IMAGE Consortium AA231015
1689 C	5	021DA1G5	IMAGE EST {IMAGE:671950}	671950 IMAGE Consortium AA244935
1690 C	7	021DA1G5	IMAGE EST {IMAGE:672427}	672427 IMAGE Consortium AA231034
1691 C	9	021DA1G5	ESTs {IMAGE:672668}	672668 IMAGE Consortium AA231603
1692 C	11	021DA1G5	ESTs, Highly similar to NIFS-LIKE 54.5 KD PROTEIN [Saccharomyces cerevisiae] {IMAGE:676125}	676125 IMAGE Consortium AA209096
1693 E	1	021DA1G5	ESTs, Highly similar to VACUOLAR ATP SYNTHASE SUBUNIT AC45 PRECURSOR [Bos taurus] {IMAGE:671660}	671660 IMAGE Consortium AA241780
1694 E	3	021DA1G5	Yamaguchi sarcoma viral (v-yes-1) oncogene homolog {IMAGE:672263}	672263 IMAGE Consortium AA230995
1695 E	5	021DA1G5	Inter-alpha trypsin inhibitor, heavy chain 2 {IMAGE:676327}	676327 IMAGE Consortium AA208770
1696 E	7	021DA1G5	Glycerol-3-phosphate acyltransferase, mitochondrial {IMAGE:676437}	676437 IMAGE Consortium AA209041
1697 E	9	021DA1G5	IMAGE EST {IMAGE:670136}	670136 IMAGE Consortium AA231045
1698 E	11	021DA1G5	IMAGE EST {IMAGE:677167}	677167 IMAGE Consortium AA213095
1699 G	1	021DA1G5	Mus musculus endothelial monocyte-activating polypeptide I mRNA, complete cds {IMAGE:669969}	669969 IMAGE Consortium AA253928
1700 G	3	021DA1G5	ESTs {IMAGE:670223}	670223 IMAGE Consortium AA221226

1701 G	5 021DA1G5	Proteasome (prosome, macropain) subunit, beta type 7 {IMAGE:677204}	677204 IMAGE Consortium	AA212838
1702 G	7 021DA1G5	IMAGE EST {IMAGE:678648}	678648 IMAGE Consortium	AA238033
1703 G	9 021DA1G5	IMAGE EST {IMAGE:659101}	659101 IMAGE Consortium	AA217773
1704 G	11 021DA1G5	Trophoblast specific protein {IMAGE:659899}	659899 IMAGE Consortium	AA220024
1705 A	1 021KA1G6	IMAGE EST {IMAGE:679181}	679181 IMAGE Consortium	AA245899
1706 A	3 021KA1G6	IMAGE EST {IMAGE:692621}	692621 IMAGE Consortium	AA239789
1707 A	5 021KA1G6	ESTs {IMAGE:692801}	692801 IMAGE Consortium	AA239937
1708 A	7 021KA1G6	CD1d1 antigen {IMAGE:693315}	693315 IMAGE Consortium	AA397125
1709 A	9 021KA1G6	M.musculus mRNA for 6-phosphofructo-2-kinase/fructose-2,6-bisphosphatase, clone 500bC6 {IMAGE:693346}	693346 IMAGE Consortium	AA397024
1710 A	11 021KA1G6	Mus musculus PS-2 mRNA, complete cds {IMAGE:693718}	693718 IMAGE Consortium	AA265273
1711 C	1 021KA1G6	ESTs {IMAGE:693004}	693004 IMAGE Consortium	AA253648
1712 C	3 021KA1G6	Mus musculus DAP-kinase related protein 1 mRNA, partial cds {IMAGE:693231}	693231 IMAGE Consortium	AA396995
1713 C	5 021KA1G6	Mus musculus flavin-containing monooxygenase 5 (FMO5) mRNA, complete cds {IMAGE:680453}	680453 IMAGE Consortium	AA237178
1714 C	7 021KA1G6	Retinoic acid early transcript beta {IMAGE:680861}	680861 IMAGE Consortium	AA250069
1715 C	9 021KA1G6	Complement component 3a receptor 1 {IMAGE:694996}	694996 IMAGE Consortium	AA221119
1716 C	11 021KA1G6	Thymoma viral proto-oncogene {IMAGE:678740}	678740 IMAGE Consortium	AA237366
1717 E	1 021KA1G6	IMAGE EST {IMAGE:680958}	680958 IMAGE Consortium	AA241899
1718 E	3 021KA1G6	IMAGE EST {IMAGE:694332}	694332 IMAGE Consortium	AA221523
1719 E	5 021KA1G6	Mus musculus semaphorin IV isoform b mRNA, complete cds {IMAGE:695117}	695117 IMAGE Consortium	AA216900
1720 E	7 021KA1G6	ESTs {IMAGE:679015}	679015 IMAGE Consortium	AA245808
1721 E	9 021KA1G6	ESTs {IMAGE:692256}	692256 IMAGE Consortium	AA238240
1722 E	11 021KA1G6	ESTs, Weakly similar to breast cancer suppressor candidate 1 [H.sapiens] {IMAGE:679316}	679316 IMAGE Consortium	AA245712
1723 G	1 021KA1G6	ESTs {IMAGE:680871}	680871 IMAGE Consortium	AA250107
1724 G	3 021KA1G6	ESTs, Weakly similar to liver Na+/Pi cotransporter-1 [R.norvegicus] {IMAGE:681706}	681706 IMAGE Consortium	AA245847
1725 G	5 021KA1G6	IMAGE EST {IMAGE:692568}	692568 IMAGE Consortium	AA250662
1726 G	7 021KA1G6	Coagulation factor IX {IMAGE:692718}	692718 IMAGE Consortium	AA239875
1727 G	9 021KA1G6	Solute carrier family 1, member 2 {IMAGE:692852}	692852 IMAGE Consortium	AA238533
1728 G	11 021KA1G6	ESTs, Moderately similar to LETHAL(2) GIANT LARVAE PROTEIN [Drosophila melanogaster] {IMAGE:693234}	693234 IMAGE Consortium	AA397070
1729 A	1 021RA1G7	ESTs, Highly similar to ERGIC-53 PROTEIN PRECURSOR [Homo sapiens] {IMAGE:681835}	681835 IMAGE Consortium	AA244713
1730 A	3 021RA1G7	IMAGE EST {IMAGE:692593}	692593 IMAGE Consortium	AA250689
1731 A	5 021RA1G7	IMAGE EST {IMAGE:692325}	692325 IMAGE Consortium	AA260603
1732 A	7 021RA1G7	ESTs {IMAGE:679665}	679665 IMAGE Consortium	AA245149
1733 A	9 021RA1G7	DOPACHROME TAUTOMERASE PRECURSOR {IMAGE:693365}	693365 IMAGE Consortium	AA397133

1734 A	11 021RA1G7	Mus musculus annexin IV mRNA, complete cds {IMAGE:693045}	693045 IMAGE Consortium	AA397114
1735 C	1 021RA1G7	ESTs {IMAGE:693145}	693145 IMAGE Consortium	AA253680
1736 C	3 021RA1G7	ESTs {IMAGE:693246}	693246 IMAGE Consortium	AA250171
1737 C	5 021RA1G7	Ferritin heavy chain {IMAGE:693166}	693166 IMAGE Consortium	AA250186
1738 C	7 021RA1G7	IMAGE EST {IMAGE:680605}	680605 IMAGE Consortium	AA237260
1739 C	9 021RA1G7	ESTs {IMAGE:680746}	680746 IMAGE Consortium	AA241911
1740 C	11 021RA1G7	ESTs, Weakly similar to POSSIBLE AGMATINASE [Streptomyces clavuligerus] {IMAGE:681564}	681564 IMAGE Consortium	AA237564
1741 E	1 021RA1G7	ESTs, Moderately similar to hypothetical protein [M.musculus] {IMAGE:681010}	681010 IMAGE Consortium	AA250077
1742 E	3 021RA1G7	Mus musculus major histocompatibility locus class II region; Fas-binding protein Daxx (DAXX) gene, partial cds; Bing1 (BING1), tapasin (tapasin), RalGDS-like factor (RLF), KE2 (KE2), BING4 (BING4), beta1, 3-galactosyl transferase {IMAGE:694391}	694391 IMAGE Consortium	AA220114
1743 E	5 021RA1G7	Mus musculus peroxisomal biogenesis factor (Pex11a) mRNA, complete cds {IMAGE:678785}	678785 IMAGE Consortium	AA237383
1744 E	7 021RA1G7	IMAGE EST {IMAGE:681669}	681669 IMAGE Consortium	AA237998
1745 E	9 021RA1G7	ESTs, Highly similar to ALPHA-1-ANTIPROTEINASE PRECURSOR [Rattus norvegicus] {IMAGE:692260}	692260 IMAGE Consortium	AA238242
1746 E	11 021RA1G7	Dipeptidylpeptidase 4 {IMAGE:679896}	679896 IMAGE Consortium	AA237541
1747 G	1 021RA1G7	M.musculus mRNA for alpha-2 antiplasmin {IMAGE:679252}	679252 IMAGE Consortium	AA245727
1748 G	3 021RA1G7	ESTs {IMAGE:692668}	692668 IMAGE Consortium	AA239810
1749 G	5 021RA1G7	ESTs {IMAGE:679931}	679931 IMAGE Consortium	AA237701
1750 G	7 021RA1G7	Heparin cofactor II {IMAGE:693170}	693170 IMAGE Consortium	AA253700
1751 G	9 021RA1G7	ESTs {IMAGE:680250}	680250 IMAGE Consortium	AA237600
1752 G	11 021RA1G7	IMAGE EST {IMAGE:680667}	680667 IMAGE Consortium	AA237331
1753 A	1 021YA1G8	ESTs, Moderately similar to retinoid X receptor interacting protein No. 15 [M.musculus] {IMAGE:679617}	679617 IMAGE Consortium	AA242625
1754 A	3 021YA1G8	ESTs {IMAGE:680318}	680318 IMAGE Consortium	AA237488
1755 A	5 021YA1G8	ESTs, Weakly similar to HYPOTHETICAL 92.1 KD PROTEIN C24H6.03 IN CHROMOSOME I [Schizosaccharomyces pombe] {IMAGE:693584}	693584 IMAGE Consortium	AA266633
1756 A	7 021YA1G8	IMAGE EST {IMAGE:680397}	680397 IMAGE Consortium	AA238208
1757 A	9 021YA1G8	ESTs {IMAGE:681031}	681031 IMAGE Consortium	AA242641
1758 A	11 021YA1G8	Phosphatidylethanolamine N-methyltransferase {IMAGE:680973}	680973 IMAGE Consortium	AA242031
1759 C	1 021YA1G8	ESTs {IMAGE:693341}	693341 IMAGE Consortium	AA397083
1760 C	3 021YA1G8	IMAGE EST {IMAGE:693565}	693565 IMAGE Consortium	AA239277
1761 C	5 021YA1G8	CD68 antigen {IMAGE:694293}	694293 IMAGE Consortium	AA220007
1762 C	7 021YA1G8	ESTs, Weakly similar to tazarotene-induced gene 2 [H.sapiens] {IMAGE:695491}	695491 IMAGE Consortium	AA220505
1763 C	9 021YA1G8	ESTs, Highly similar to INORGANIC PYROPHOSPHATASE	716531 IMAGE Consortium	AA265508

1764 C	11 021YA1G8	[Bos taurus] {IMAGE:716531} GALECTIN-3 {IMAGE:717226}	717226 IMAGE Consortium	AA403841
1765 E	1 021YA1G8	IMAGE EST {IMAGE:717132}	717132 IMAGE Consortium	AA266229
1766 E	3 021YA1G8	Malate dehydrogenase, mitochondrial {IMAGE:717095}	717095 IMAGE Consortium	AA266087
1767 E	5 021YA1G8	ESTs {IMAGE:717873}	717873 IMAGE Consortium	AA249953
1768 E	7 021YA1G8	ESTs {IMAGE:717787}	717787 IMAGE Consortium	AA250312
1769 E	9 021YA1G8	IMAGE EST {IMAGE:697816}	697816 IMAGE Consortium	AA238723
1770 E	11 021YA1G8	IMAGE EST {IMAGE:718542}	718542 IMAGE Consortium	AA266606
1771 G	1 021YA1G8	IMAGE EST {IMAGE:697284}	697284 IMAGE Consortium	AA242075
1772 G	3 021YA1G8	ESTs, Weakly similar to HYPOTHETICAL 36.7 KD PROTEIN C2F7.02C IN CHROMOSOME I [Schizosaccharomyces pombe] {IMAGE:697509}	697509 IMAGE Consortium	AA253923
1773 G	5 021YA1G8	ESTs {IMAGE:718054}	718054 IMAGE Consortium	AA261192
1774 G	7 021YA1G8	M.musculus mRNA for T-cell receptor (helper hybridoma 2B4) {IMAGE:718710}	718710 IMAGE Consortium	AA265714
1775 G	9 021YA1G8	ESTs, Moderately similar to PROBABLE UBIQUITIN CARBOXYL-TERMINAL HYDROLASE [Mus musculus] {IMAGE:718426}	718426 IMAGE Consortium	AA266779
1776 G	11 021YA1G8	Alcohol dehydrogenase 1, complex {IMAGE:695105}	695105 IMAGE Consortium	AA221141
1777 A	1 0215A1G9	Duffy blood group {IMAGE:718084}	718084 IMAGE Consortium	AA266438
1778 A	3 0215A1G9	ESTs {IMAGE:718240}	718240 IMAGE Consortium	AA266367
1779 A	5 0215A1G9	T-COMPLEX PROTEIN 1, ALPHA SUBUNIT B {IMAGE:718103}	718103 IMAGE Consortium	AA266324
1780 A	7 0215A1G9	Kinesin light chain 2 {IMAGE:698302}	698302 IMAGE Consortium	AA239281
1781 A	9 0215A1G9	Serine protease inhibitor 4 {IMAGE:694987}	694987 IMAGE Consortium	AA218279
1782 A	11 0215A1G9	IMAGE EST {IMAGE:716713}	716713 IMAGE Consortium	AA265198
1783 C	1 0215A1G9	Kras oncogene-associated gene {IMAGE:696004}	696004 IMAGE Consortium	AA222521
1784 C	3 0215A1G9	ESTs {IMAGE:716683}	716683 IMAGE Consortium	AA265033
1785 C	5 0215A1G9	ESTs {IMAGE:696489}	696489 IMAGE Consortium	AA239673
1786 C	7 0215A1G9	ESTs, Weakly similar to predicted using Genefinder [C.elegans] {IMAGE:717387}	717387 IMAGE Consortium	AA404192
1787 C	9 0215A1G9	ESTs {IMAGE:697416}	697416 IMAGE Consortium	AA245029
1788 C	11 0215A1G9	ESTs, Moderately similar to Gps2 [H.sapiens] {IMAGE:717710}	717710 IMAGE Consortium	AA250392
1789 E	1 0215A1G9	IMAGE EST {IMAGE:696657}	696657 IMAGE Consortium	AA240258
1790 E	3 0215A1G9	ESTs, Moderately similar to LEYDIG CELL TUMOR 10 KD PROTEIN [Rattus norvegicus] {IMAGE:696473}	696473 IMAGE Consortium	AA239495
1791 E	5 0215A1G9	IMAGE EST {IMAGE:717469}	717469 IMAGE Consortium	AA403436
1792 E	7 0215A1G9	ESTs, Weakly similar to ubiquitin-specific protease [D.melanogaster] {IMAGE:717926}	717926 IMAGE Consortium	AA250238
1793 E	9 0215A1G9	Caudal type homeo box 2 {IMAGE:697874}	697874 IMAGE Consortium	AA238566
1794 E	11 0215A1G9	Mus musculus mRNA for GFRalpha-3, complete cds {IMAGE:697841}	697841 IMAGE Consortium	AA238748

1795 G	1 0215A1G9	Mus musculus CD-1 orphan receptor TAK1 (TAK1) mRNA, complete cds {IMAGE:717471}	717471 IMAGE Consortium	AA403437
1796 G	3 0215A1G9	Xeroderma pigmentosum, complementation group C {IMAGE:717771}	717771 IMAGE Consortium	AA249976
1797 G	5 0215A1G9	IMAGE EST {IMAGE:717911}	717911 IMAGE Consortium	AA250565
1798 G	7 0215A1G9	IMAGE EST {IMAGE:697981}	697981 IMAGE Consortium	AA239015
1799 G	9 0215A1G9	Growth factor independent 1B {IMAGE:718609}	718609 IMAGE Consortium	AA266627
1800 G	11 0215A1G9	IMAGE EST {IMAGE:718663}	718663 IMAGE Consortium	AA265635
1801 A	1 021PA1GA	ESTs, Moderately similar to HETEROGENEOUS NUCLEAR RIBONUCLEOPROTEIN C [Rattus norvegicus] {IMAGE:723554}	723554 IMAGE Consortium	AA260901
1802 A	3 021PA1GA	IMAGE EST {IMAGE:733519}	733519 IMAGE Consortium	AA259874
1803 A	5 021PA1GA	M.musculus mRNA for ubiquitin-conjugating enzyme UbcM3 {IMAGE:733338}	733338 IMAGE Consortium	AA259573
1804 A	7 021PA1GA	ESTs {IMAGE:733903}	733903 IMAGE Consortium	AA269330
1805 A	9 021PA1GA	ESTs {IMAGE:733959}	733959 IMAGE Consortium	AA272750
1806 A	11 021PA1GA	IMAGE EST {IMAGE:719001}	719001 IMAGE Consortium	AA267531
1807 C	1 021PA1GA	IMAGE EST {IMAGE:720754}	720754 IMAGE Consortium	AA266896
1808 C	3 021PA1GA	Interleukin 12b {IMAGE:720765}	720765 IMAGE Consortium	AA267353
1809 C	5 021PA1GA	IMAGE EST {IMAGE:721256}	721256 IMAGE Consortium	AA267016
1810 C	7 021PA1GA	Fibroblast growth factor receptor 1 {IMAGE:734032}	734032 IMAGE Consortium	AA272097
1811 C	9 021PA1GA	IMAGE EST {IMAGE:718739}	718739 IMAGE Consortium	AA265744
1812 C	11 021PA1GA	IMAGE EST {IMAGE:718892}	718892 IMAGE Consortium	AA265828
1813 E	1 021PA1GA	IMAGE EST {IMAGE:720904}	720904 IMAGE Consortium	AA267068
1814 E	3 021PA1GA	IMAGE EST {IMAGE:720885}	720885 IMAGE Consortium	AA267066
1815 E	5 021PA1GA	ESTs, Highly similar to PUTATIVE GLUCOSAMINE-6-PHOSPHATE ISOMERASE [Homo sapiens] {IMAGE:721113}	721113 IMAGE Consortium	AA267196
1816 E	7 021PA1GA	IMAGE EST {IMAGE:733937}	733937 IMAGE Consortium	AA268342
1817 E	9 021PA1GA	ESTs {IMAGE:718955}	718955 IMAGE Consortium	AA265942
1818 E	11 021PA1GA	IMAGE EST {IMAGE:718797}	718797 IMAGE Consortium	AA265633
1819 G	1 021PA1GA	IMAGE EST {IMAGE:720899}	720899 IMAGE Consortium	AA267209
1820 G	3 021PA1GA	ESTs {IMAGE:733642}	733642 IMAGE Consortium	AA259486
1821 G	5 021PA1GA	IMAGE EST {IMAGE:721078}	721078 IMAGE Consortium	AA267610
1822 G	7 021PA1GA	IMAGE EST {IMAGE:734258}	734258 IMAGE Consortium	AA268881
1823 G	9 021PA1GA	IMAGE EST {IMAGE:721674}	721674 IMAGE Consortium	AA260880
1824 G	11 021PA1GA	IMAGE EST {IMAGE:721974}	721974 IMAGE Consortium	AA268404
1825 A	1 021WA1GB	IMAGE EST {IMAGE:717992}	717992 IMAGE Consortium	AA261675
1826 A	3 021WA1GB	IMAGE EST {IMAGE:718515}	718515 IMAGE Consortium	AA266265
1827 A	5 021WA1GB	Cytochrome P450, 2e1, ethanol inducible {IMAGE:718548}	718548 IMAGE Consortium	AA267744
1828 A	7 021WA1GB	IMAGE EST {IMAGE:695508}	695508 IMAGE Consortium	AA220515
1829 A	9 021WA1GB	ESTs {IMAGE:695769}	695769 IMAGE Consortium	AA222221
1830 A	11 021WA1GB	ESTs {IMAGE:696674}	696674 IMAGE Consortium	AA240146
1831 C	1 021WA1GB	ESTs {IMAGE:696794}	696794 IMAGE Consortium	AA240283

1832 C	3	021WA1GB	ESTs {IMAGE:717161}	717161	IMAGE Consortium	AA266579
1833 C	5	021WA1GB	IMAGE EST {IMAGE:697481}	697481	IMAGE Consortium	AA253908
1834 C	7	021WA1GB	IMAGE EST {IMAGE:718018}	718018	IMAGE Consortium	AA261682
1835 C	9	021WA1GB	IMAGE EST {IMAGE:697893}	697893	IMAGE Consortium	AA238765
1836 C	11	021WA1GB	ESTs {IMAGE:699489}	699489	IMAGE Consortium	AA245586
1837 E	1	021WA1GB	IMAGE EST {IMAGE:699340}	699340	IMAGE Consortium	AA245373
1838 E	3	021WA1GB	ESTs, Moderately similar to predicted using Genefinder [C.elegans] {IMAGE:699466}	699466	IMAGE Consortium	AA245578
1839 E	5	021WA1GB	ESTs {IMAGE:721478}	721478	IMAGE Consortium	AA266720
1840 E	7	021WA1GB	ESTs {IMAGE:721944}	721944	IMAGE Consortium	AA267983
1841 E	9	021WA1GB	ESTs, Weakly similar to LIM PROTEIN RIL [Rattus norvegicus] {IMAGE:721812}	721812	IMAGE Consortium	AA267893
1842 E	11	021WA1GB	Serum amyloid P-component {IMAGE:722134}	722134	IMAGE Consortium	AA268587
1843 G	1	021WA1GB	ESTs {IMAGE:733638}	733638	IMAGE Consortium	AA259431
1844 G	3	021WA1GB	Mus musculus L6 antigen mRNA, complete cds {IMAGE:733601}	733601	IMAGE Consortium	AA259414
1845 G	5	021WA1GB	IMAGE EST {IMAGE:733734}	733734	IMAGE Consortium	AA272876
1846 G	7	021WA1GB	Cell division cycle 42 {IMAGE:721518}	721518	IMAGE Consortium	AA266975
1847 G	9	021WA1GB	ESTs {IMAGE:719072}	719072	IMAGE Consortium	AA266014
1848 G	11	021WA1GB	ESTs {IMAGE:719034}	719034	IMAGE Consortium	AA265975
1849 A	1	0213A1GC	Angiotensinogen {IMAGE:733572}	733572	IMAGE Consortium	AA259979
1850 A	3	0213A1GC	IMAGE EST {IMAGE:733438}	733438	IMAGE Consortium	AA259659
1851 A	5	0213A1GC	Nuclear factor, erythroid derived 2, 45 kDa {IMAGE:733953}	733953	IMAGE Consortium	AA272747
1852 A	7	0213A1GC	IMAGE EST {IMAGE:721592}	721592	IMAGE Consortium	AA267450
1853 A	9	0213A1GC	IMAGE EST {IMAGE:718812}	718812	IMAGE Consortium	AA265864
1854 A	11	0213A1GC	IMAGE EST {IMAGE:719062}	719062	IMAGE Consortium	AA265981
1855 C	1	0213A1GC	ESTs {IMAGE:720934}	720934	IMAGE Consortium	AA267077
1856 C	3	0213A1GC	ESTs {IMAGE:721118}	721118	IMAGE Consortium	AA267195
1857 C	5	0213A1GC	ESTs {IMAGE:721417}	721417	IMAGE Consortium	AA267567
1858 C	7	0213A1GC	IMAGE EST {IMAGE:719028}	719028	IMAGE Consortium	AA265973
1859 C	9	0213A1GC	IMAGE EST {IMAGE:719232}	719232	IMAGE Consortium	AA254630
1860 C	11	0213A1GC	ESTs {IMAGE:719457}	719457	IMAGE Consortium	AA261021
1861 E	1	0213A1GC	IMAGE EST {IMAGE:733800}	733800	IMAGE Consortium	AA268030
1862 E	3	0213A1GC	Mus musculus elongation factor 1-beta homolog mRNA, complete cds {IMAGE:733810}	733810	IMAGE Consortium	AA268148
1863 E	5	0213A1GC	ESTs {IMAGE:734084}	734084	IMAGE Consortium	AA272114
1864 E	7	0213A1GC	IMAGE EST {IMAGE:719233}	719233	IMAGE Consortium	AA254633
1865 E	9	0213A1GC	IMAGE EST {IMAGE:721939}	721939	IMAGE Consortium	AA267980
1866 E	11	0213A1GC	TRANSDUCIN-LIKE ENHANCER PROTEIN 3 {IMAGE:722501}	722501	IMAGE Consortium	AA253995
1867 G	1	0213A1GC	IMAGE EST {IMAGE:733726}	733726	IMAGE Consortium	AA268327
1868 G	3	0213A1GC	IMAGE EST {IMAGE:719282}	719282	IMAGE Consortium	AA254672
1869 G	5	0213A1GC	Serine/threonine kinase 2 {IMAGE:722059}	722059	IMAGE Consortium	AA268478



1870 G	7 0213A1GC	ESTs {IMAGE:722238}	722238 IMAGE Consortium	AA254173
1871 G	9 0213A1GC	ESTs, Highly similar to GALACTOKINASE [Escherichia coli] {IMAGE:719937}	719937 IMAGE Consortium	AA255078
1872 G	11 0213A1GC	ESTs {IMAGE:720189}	720189 IMAGE Consortium	AA255028
1873 A	1 021AA1GD	ESTs {IMAGE:719315}	719315 IMAGE Consortium	AA255009
1874 A	3 021AA1GD	IMAGE EST {IMAGE:722262}	722262 IMAGE Consortium	AA254198
1875 A	5 021AA1GD	ESTs {IMAGE:722364}	722364 IMAGE Consortium	AA254110
1876 A	7 021AA1GD	ESTs {IMAGE:722231}	722231 IMAGE Consortium	AA254005
1877 A	9 021AA1GD	ESTs {IMAGE:720380}	720380 IMAGE Consortium	AA261577
1878 A	11 021AA1GD	ESTs, Weakly similar to TIP49 [R.norvegicus] {IMAGE:720870}	720870 IMAGE Consortium	AA267034
1879 C	1 021AA1GD	IMAGE EST {IMAGE:722358}	722358 IMAGE Consortium	AA254256
1880 C	3 021AA1GD	ESTs, Highly similar to PROBABLE UBIQUITIN CARBOXYL-TERMINAL HYDROLASE [Mus musculus] {IMAGE:720155}	720155 IMAGE Consortium	AA254871
1881 C	5 021AA1GD	ESTs {IMAGE:722631}	722631 IMAGE Consortium	AA260701
1882 C	7 021AA1GD	IMAGE EST {IMAGE:723137}	723137 IMAGE Consortium	AA267833
1883 C	9 021AA1GD	ESTs {IMAGE:723311}	723311 IMAGE Consortium	AA267798
1884 C	11 021AA1GD	Ect2 oncogene {IMAGE:721085}	721085 IMAGE Consortium	AA267000
1885 E	1 021AA1GD	ESTs {IMAGE:735376}	735376 IMAGE Consortium	AA269496
1886 E	3 021AA1GD	Mouse testis abundant mRNA sequence {IMAGE:746488}	746488 IMAGE Consortium	AA260173
1887 E	5 021AA1GD	IMAGE EST {IMAGE:747104}	747104 IMAGE Consortium	AA274932
1888 E	7 021AA1GD	ESTs {IMAGE:746872}	746872 IMAGE Consortium	AA274800
1889 E	9 021AA1GD	U2 small nuclear ribonucleoprotein auxiliary factor (U2AF), 35 kDa, related sequence 1 {IMAGE:747294}	747294 IMAGE Consortium	AA274915
1890 E	11 021AA1GD	ESTs {IMAGE:736526}	736526 IMAGE Consortium	AA270527
1891 G	1 021AA1GD	ESTs {IMAGE:735583}	735583 IMAGE Consortium	AA270043
1892 G	3 021AA1GD	ESTs {IMAGE:746597}	746597 IMAGE Consortium	AA268054
1893 G	5 021AA1GD	Mini chromosome maintenance deficient 4 homolog (S. cerevisiae) {IMAGE:746419}	746419 IMAGE Consortium	AA259788
1894 G	7 021AA1GD	ESTs {IMAGE:736011}	736011 IMAGE Consortium	AA269821
1895 G	9 021AA1GD	IMAGE EST {IMAGE:747318}	747318 IMAGE Consortium	AA274945
1896 G	11 021AA1GD	Protein kinase, cAMP dependent regulatory, type I beta {IMAGE:736724}	736724 IMAGE Consortium	AA270948
1897 A	1 021HA1GE	ESTs {IMAGE:734175}	734175 IMAGE Consortium	AA268614
1898 A	3 021HA1GE	IMAGE EST {IMAGE:734816}	734816 IMAGE Consortium	AA259369
1899 A	5 021HA1GE	IMAGE EST {IMAGE:734908}	734908 IMAGE Consortium	AA261708
1900 A	7 021HA1GE	Myoglobin {IMAGE:738009}	738009 IMAGE Consortium	AA271284
1901 A	9 021HA1GE	IMAGE EST {IMAGE:738019}	738019 IMAGE Consortium	AA271530
1902 A	11 021HA1GE	IMAGE EST {IMAGE:746568}	746568 IMAGE Consortium	AA267925
1903 C	1 021HA1GE	ESTs, Weakly similar to homeotic region most like HMPB_DROME: homeotic proboscipedia protein [C.elegans] {IMAGE:747239}	747239 IMAGE Consortium	AA260520
1904 C	3 021HA1GE	Mus musculus peroxisomal acyl-CoA oxidase (muspaox)	747880 IMAGE Consortium	AA261287

1905 C	5 021HA1GE	mRNA, complete cds {IMAGE:747880} IMAGE EST {IMAGE:747939}	747939 IMAGE Consortium	AA260949
1906 C	7 021HA1GE	ESTs, Highly similar to OLIGOSACCHARYL TRANSFERASE STT3 SUBUNIT HOMOLOG [Caenorhabditis elegans] {IMAGE:737445}	737445 IMAGE Consortium	AA277846
1907 C	9 021HA1GE	IMAGE EST {IMAGE:734861}	734861 IMAGE Consortium	AA259400
1908 C	11 021HA1GE	ESTs, Moderately similar to RAF SERINE/THREONINE- PROTEIN KINASE TRANSFORMING PROTEIN [Murine sarcoma virus 3611] {IMAGE:735016}	735016 IMAGE Consortium	AA260158
1909 E	1 021HA1GE	ESTs, Highly similar to autoantigen [H.sapiens] {IMAGE:736611}	736611 IMAGE Consortium	AA270674
1910 E	3 021HA1GE	IMAGE EST {IMAGE:736506}	736506 IMAGE Consortium	AA270729
1911 E	5 021HA1GE	ESTs, Moderately similar to CREB-BINDING PROTEIN [M.musculus] {IMAGE:736783}	736783 IMAGE Consortium	AA271091
1912 E	7 021HA1GE	ESTs, Weakly similar to cDNA EST EMBL:Z14593 comes from this gene [C.elegans] {IMAGE:748059}	748059 IMAGE Consortium	AA275027
1913 E	9 021HA1GE	ESTs {IMAGE:737541}	737541 IMAGE Consortium	AA277366
1914 E	11 021HA1GE	ESTs {IMAGE:737745}	737745 IMAGE Consortium	AA277700
1915 G	1 021HA1GE	IMAGE EST {IMAGE:747279}	747279 IMAGE Consortium	AA274912
1916 G	3 021HA1GE	Mus musculus casein kinase II alpha subunit mRNA, complete cds {IMAGE:747427}	747427 IMAGE Consortium	AA272845
1917 G	5 021HA1GE	ESTs {IMAGE:747503}	747503 IMAGE Consortium	AA268441
1918 G	7 021HA1GE	ESTs {IMAGE:747763}	747763 IMAGE Consortium	AA274840
1919 G	9 021HA1GE	ESTs {IMAGE:737447}	737447 IMAGE Consortium	AA277847
1920 G	11 021HA1GE	ESTs {IMAGE:738252}	738252 IMAGE Consortium	AA271223
1921 A	1 021OA1GF	ESTs, Highly similar to LIVER 60 KD CARBOXYLESTERASE 2 [Oryctolagus cuniculus] {IMAGE:737998}	737998 IMAGE Consortium	AA271522
1922 A	3 021OA1GF	IMAGE EST {IMAGE:735527}	735527 IMAGE Consortium	AA269845
1923 A	5 021OA1GF	IMAGE EST {IMAGE:736101}	736101 IMAGE Consortium	AA270364
1924 A	7 021OA1GF	ESTs {IMAGE:736622}	736622 IMAGE Consortium	AA270672
1925 A	9 021OA1GF	ESTs {IMAGE:737055}	737055 IMAGE Consortium	AA272056
1926 A	11 021OA1GF	IMAGE EST {IMAGE:734362}	734362 IMAGE Consortium	AA268702
1927 C	1 021OA1GF	ESTs, Highly similar to HLA-DR ASSOCIATED PROTEIN I [Homo sapiens] {IMAGE:746994}	746994 IMAGE Consortium	AA260352
1928 C	3 021OA1GF	ESTs, Highly similar to sodium-D-glucose cotransporter [H.sapiens] {IMAGE:747297}	747297 IMAGE Consortium	AA274938
1929 C	5 021OA1GF	ESTs {IMAGE:736718}	736718 IMAGE Consortium	AA270916
1930 C	7 021OA1GF	IMAGE EST {IMAGE:736848}	736848 IMAGE Consortium	AA271272
1931 C	9 021OA1GF	IMAGE EST {IMAGE:747638}	747638 IMAGE Consortium	AA268813
1932 C	11 021OA1GF	ESTs {IMAGE:747862}	747862 IMAGE Consortium	AA261715
1933 E	1 021OA1GF	Mus musculus mRNA for 26S proteasome non-ATPase subunit {IMAGE:747059}	747059 IMAGE Consortium	AA274539
1934 E	3 021OA1GF	IMAGE EST {IMAGE:736046}	736046 IMAGE Consortium	AA270270

1935 E	5 021OA1GF	Mus musculus mRNA for mLGP85/LIMP II, complete cds {IMAGE:747497}	747497 IMAGE Consortium	AA268438
1936 E	7 021OA1GF	Apolipoprotein CII {IMAGE:736992}	736992 IMAGE Consortium	AA271959
1937 E	9 021OA1GF	IMAGE EST {IMAGE:747662}	747662 IMAGE Consortium	AA268786
1938 E	11 021OA1GF	ESTs, Weakly similar to coded for by C. elegans cDNA yk102d10.5 [C.elegans] {IMAGE:734791}	734791 IMAGE Consortium	AA259340
1939 G	1 021OA1GF	E4F transcription factor 1 {IMAGE:736576}	736576 IMAGE Consortium	AA270628
1940 G	3 021OA1GF	MEMBRANE-ASSOCIATED PROTEIN HEM-2 {IMAGE:747583}	747583 IMAGE Consortium	AA270873
1941 G	5 021OA1GF	ESTs {IMAGE:747543}	747543 IMAGE Consortium	AA268977
1942 G	7 021OA1GF	M.musculus mRNA for casein kinase I-alpha {IMAGE:734318}	734318 IMAGE Consortium	AA268671
1943 G	9 021OA1GF	ESTs, Weakly similar to ZINC FINGER PROTEIN HF.12 [Homo sapiens] {IMAGE:737815}	737815 IMAGE Consortium	AA277748
1944 G	11 021OA1GF	Mouse tryptophan-2,3-dioxygenase (TDO) mRNA, complete cds {IMAGE:737586}	737586 IMAGE Consortium	AA277213
1945 A	1 021VA1GG	ESTs {IMAGE:747800}	747800 IMAGE Consortium	AA274528
1946 A	3 021VA1GG	IMAGE EST {IMAGE:737533}	737533 IMAGE Consortium	AA277724
1947 A	5 021VA1GG	ESTs {IMAGE:737519}	737519 IMAGE Consortium	AA277359
1948 A	7 021VA1GG	IMAGE EST {IMAGE:737868}	737868 IMAGE Consortium	AA277159
1949 A	9 021VA1GG	ESTs, Highly similar to FIBRINOGEN GAMMA CHAIN PRECURSOR [Petromyzon marinus] {IMAGE:737780}	737780 IMAGE Consortium	AA277830
1950 A	11 021VA1GG	IMAGE EST {IMAGE:735414}	735414 IMAGE Consortium	AA272825
1951 C	1 021VA1GG	Glutamate cysteine ligase (gamma-glutamylcysteine synthetase), regulatory {IMAGE:750596}	750596 IMAGE Consortium	AA472979
1952 C	3 021VA1GG	ESTs, Moderately similar to Rab8-interacting protein [M.musculus] {IMAGE:750758}	750758 IMAGE Consortium	AA290306
1953 C	5 021VA1GG	ESTs {IMAGE:751146}	751146 IMAGE Consortium	AA288477
1954 C	7 021VA1GG	IMAGE EST {IMAGE:749257}	749257 IMAGE Consortium	AA290443
1955 C	9 021VA1GG	IMAGE EST {IMAGE:749363}	749363 IMAGE Consortium	AA289477
1956 C	11 021VA1GG	IMAGE EST {IMAGE:749449}	749449 IMAGE Consortium	AA289656
1957 E	1 021VA1GG	ESTs, Weakly similar to SPORULATION-SPECIFIC PROTEIN 1 [Saccharomyces cerevisiae] {IMAGE:751108}	751108 IMAGE Consortium	AA290484
1958 E	3 021VA1GG	ESTs {IMAGE:751543}	751543 IMAGE Consortium	AA396044
1959 E	5 021VA1GG	ESTs {IMAGE:751645}	751645 IMAGE Consortium	AA396082
1960 E	7 021VA1GG	ESTs {IMAGE:751969}	751969 IMAGE Consortium	AA396321
1961 E	9 021VA1GG	IMAGE EST {IMAGE:749545}	749545 IMAGE Consortium	AA388682
1962 E	11 021VA1GG	ESTs {IMAGE:749506}	749506 IMAGE Consortium	AA289542
1963 G	1 021VA1GG	Interferon concensus sequence binding protein {IMAGE:751220}	751220 IMAGE Consortium	AA288222
1964 G	3 021VA1GG	IMAGE EST {IMAGE:751693}	751693 IMAGE Consortium	AA396109
1965 G	5 021VA1GG	ESTs {IMAGE:751826}	751826 IMAGE Consortium	AA396152
1966 G	7 021VA1GG	ESTs {IMAGE:751836}	751836 IMAGE Consortium	AA396224
1967 G	9 021VA1GG	ESTs {IMAGE:752474}	752474 IMAGE Consortium	AA289377

1968 G	11 021VA1GG	ESTs {IMAGE:752465}	752465 IMAGE Consortium	AA289517
1969 A	1 0212A1GH	ESTs {IMAGE:751550}	751550 IMAGE Consortium	AA396040
1970 A	3 0212A1GH	E2F transcription factor 1 {IMAGE:751755}	751755 IMAGE Consortium	AA396123
1971 A	5 0212A1GH	ELK3, member of ETS oncogene family {IMAGE:749199}	749199 IMAGE Consortium	AA289606
1972 A	7 0212A1GH	IMAGE EST {IMAGE:752395}	752395 IMAGE Consortium	AA290036
1973 A	9 0212A1GH	ESTs, Highly similar to KINESIN-II 85 KD SUBUNIT [Strongylocentrotus purpuratus] {IMAGE:761595}	761595 IMAGE Consortium	AA272363
1974 A	11 0212A1GH	ESTs {IMAGE:761554}	761554 IMAGE Consortium	AA272148
1975 C	1 0212A1GH	IMAGE EST {IMAGE:751455}	751455 IMAGE Consortium	AA395929
1976 C	3 0212A1GH	RD PROTEIN {IMAGE:749097}	749097 IMAGE Consortium	AA290538
1977 C	5 0212A1GH	ESTs, Highly similar to p20-CGGBP [H.sapiens] {IMAGE:749295}	749295 IMAGE Consortium	AA288550
1978 C	7 0212A1GH	IMAGE EST {IMAGE:749738}	749738 IMAGE Consortium	AA396395
1979 C	9 0212A1GH	IMAGE EST {IMAGE:749579}	749579 IMAGE Consortium	AA288576
1980 C	11 0212A1GH	ESTs {IMAGE:761578}	761578 IMAGE Consortium	AA272321
1981 E	1 0212A1GH	ESTs {IMAGE:749439}	749439 IMAGE Consortium	AA289582
1982 E	3 0212A1GH	IMAGE EST {IMAGE:749451}	749451 IMAGE Consortium	AA289657
1983 E	5 0212A1GH	ESTs {IMAGE:749820}	749820 IMAGE Consortium	AA396410
1984 E	7 0212A1GH	ESTs {IMAGE:750198}	750198 IMAGE Consortium	AA396459
1985 E	9 0212A1GH	ESTs {IMAGE:750537}	750537 IMAGE Consortium	AA472926
1986 E	11 0212A1GH	Mus musculus (10-1) hydroxysteroid sulfotransferase (mSTa2) mRNA, complete cds {IMAGE:748540}	748540 IMAGE Consortium	AA269437
1987 G	1 0212A1GH	IMAGE EST {IMAGE:761596}	761596 IMAGE Consortium	AA272340
1988 G	3 0212A1GH	ESTs, Highly similar to NADH-UBIQUINONE OXIDOREDUCTASE 19 KD SUBUNIT [Bos taurus] {IMAGE:761723}	761723 IMAGE Consortium	AA272175
1989 G	5 0212A1GH	IMAGE EST {IMAGE:762240}	762240 IMAGE Consortium	AA277781
1990 G	7 0212A1GH	ESTs {IMAGE:750498}	750498 IMAGE Consortium	AA472893
1991 G	9 0212A1GH	ESTs {IMAGE:748366}	748366 IMAGE Consortium	AA267934
1992 G	11 0212A1GH	Mus musculus mRNA for prefoldin subunit 2 {IMAGE:748879}	748879 IMAGE Consortium	AA289888
1993 A	1 0219A1GI	Ankyrin 3, epithelial {IMAGE:765848}	765848 IMAGE Consortium	AA274771
1994 A	3 0219A1GI	Mus musculus EXO70 protein (Exo70) mRNA, complete cds {IMAGE:763179}	763179 IMAGE Consortium	AA286651
1995 A	5 0219A1GI	ESTs {IMAGE:763531}	763531 IMAGE Consortium	AA285489
1996 A	7 0219A1GI	ESTs {IMAGE:763871}	763871 IMAGE Consortium	AA285551
1997 A	9 0219A1GI	Sterol O-acyltransferase 1 {IMAGE:764650}	764650 IMAGE Consortium	AA273904
1998 A	11 0219A1GI	Arginosuccinate synthetase 1 {IMAGE:776890}	776890 IMAGE Consortium	AA419665
1999 C	1 0219A1GI	IMAGE EST {IMAGE:763628}	763628 IMAGE Consortium	AA285580
2000 C	3 0219A1GI	ESTs {IMAGE:776238}	776238 IMAGE Consortium	AA276308
2001 C	5 0219A1GI	ESTs {IMAGE:776077}	776077 IMAGE Consortium	AA276036
2002 C	7 0219A1GI	ESTs, Weakly similar to R07G3.8 [C.elegans] {IMAGE:764435}	764435 IMAGE Consortium	AA274203
2003 C	9 0219A1GI	ESTs {IMAGE:765130}	765130 IMAGE Consortium	AA273923

2004 C	11 0219A1GI	IMAGE EST {IMAGE:777290}	777290 IMAGE Consortium	AA276878
2005 E	1 0219A1GI	ESTs {IMAGE:764044}	764044 IMAGE Consortium	AA285972
2006 E	3 0219A1GI	Mouse mRNA for ribosomal protein S12 {IMAGE:776197}	776197 IMAGE Consortium	AA276213
2007 E	5 0219A1GI	IMAGE EST {IMAGE:776013}	776013 IMAGE Consortium	AA286062
2008 E	7 0219A1GI	IMAGE EST {IMAGE:764518}	764518 IMAGE Consortium	AA273668
2009 E	9 0219A1GI	ESTs {IMAGE:765011}	765011 IMAGE Consortium	AA273572
2010 E	11 0219A1GI	IMAGE EST {IMAGE:764975}	764975 IMAGE Consortium	AA273423
2011 G	1 0219A1GI	Mus musculus mRNA for synaptogyrin 2 {IMAGE:775702}	775702 IMAGE Consortium	AA275862
2012 G	3 0219A1GI	IMAGE EST {IMAGE:775951}	775951 IMAGE Consortium	AA276290
2013 G	5 0219A1GI	Interleukin 2 receptor, gamma chain {IMAGE:764497}	764497 IMAGE Consortium	AA274104
2014 G	7 0219A1GI	Epoxide hydrolase 2, cytoplasmic {IMAGE:764542}	764542 IMAGE Consortium	AA274030
2015 G	9 0219A1GI	IMAGE EST {IMAGE:765107}	765107 IMAGE Consortium	AA273917
2016 G	11 0219A1GI	IMAGE EST {IMAGE:765119}	765119 IMAGE Consortium	AA273259
2017 A	1 021GA1GJ	IMAGE EST {IMAGE:762133}	762133 IMAGE Consortium	AA277194
2018 A	3 021GA1GJ	Uroporphyrinogen decarboxylase {IMAGE:748198}	748198 IMAGE Consortium	AA275045
2019 A	5 021GA1GJ	ESTs, Highly similar to GASTRULATION SPECIFIC PROTEIN G12 [Danio rerio] {IMAGE:750654}	750654 IMAGE Consortium	AA290107
2020 A	7 021GA1GJ	Mus musculus erythrocyte membrane protein Rh50 (Rhag) mRNA, complete cds {IMAGE:750909}	750909 IMAGE Consortium	AA289136
2021 A	9 021GA1GJ	ESTs {IMAGE:751020}	751020 IMAGE Consortium	AA289414
2022 A	11 021GA1GJ	ESTs, Moderately similar to dimethylglycine dehydrogenase [R.norvegicus] {IMAGE:748958}	748958 IMAGE Consortium	AA288418
2023 C	1 021GA1GJ	ESTs, Weakly similar to alpha tectorin [M.musculus] {IMAGE:762229}	762229 IMAGE Consortium	AA277779
2024 C	3 021GA1GJ	ESTs {IMAGE:750379}	750379 IMAGE Consortium	AA396765
2025 C	5 021GA1GJ	ESTs, Highly similar to A-kinase anchor protein 95, AKAP95 [R.norvegicus] {IMAGE:750898}	750898 IMAGE Consortium	AA290277
2026 C	7 021GA1GJ	ESTs {IMAGE:748654}	748654 IMAGE Consortium	AA270292
2027 C	9 021GA1GJ	ESTs {IMAGE:751398}	751398 IMAGE Consortium	AA395957
2028 C	11 021GA1GJ	IMAGE EST {IMAGE:751716}	751716 IMAGE Consortium	AA396116
2029 E	1 021GA1GJ	IMAGE EST {IMAGE:750780}	750780 IMAGE Consortium	AA290312
2030 E	3 021GA1GJ	Mus musculus orphan nuclear hormone receptor isoform mCAR2 mRNA, complete cds {IMAGE:748487}	748487 IMAGE Consortium	AA272757
2031 E	5 021GA1GJ	ESTs {IMAGE:749048}	749048 IMAGE Consortium	AA288990
2032 E	7 021GA1GJ	ESTs {IMAGE:751477}	751477 IMAGE Consortium	AA395996
2033 E	9 021GA1GJ	IMAGE EST {IMAGE:749164}	749164 IMAGE Consortium	AA289547
2034 E	11 021GA1GJ	ESTs, Moderately similar to GOLIATH PROTEIN [Drosophila melanogaster] {IMAGE:752174}	752174 IMAGE Consortium	AA288977
2035 G	1 021GA1GJ	ESTs {IMAGE:751152}	751152 IMAGE Consortium	AA290441
2036 G	3 021GA1GJ	IMAGE EST {IMAGE:751217}	751217 IMAGE Consortium	AA288224
2037 G	5 021GA1GJ	Mus musculus mRNA for acyl-CoA thioesterase {IMAGE:748816}	748816 IMAGE Consortium	AA288170
2038 G	7 021GA1GJ	ESTs, Highly similar to p53 tumor suppressor-binding protein	749161 IMAGE Consortium	AA289551

		1 [H.sapiens] {IMAGE:749161}		
2039 G	9	021GA1GJ IMAGE EST {IMAGE:749428}	749428	IMAGE Consortium AA289571
2040 G	11	021GA1GJ IMAGE EST {IMAGE:749291}	749291	IMAGE Consortium AA288548
2041 A	1	021NA1GK ESTs, Moderately similar to SYNUCLEIN [Bos taurus] {IMAGE:763358}	763358	IMAGE Consortium AA272342
2042 A	3	021NA1GK Polycystic kidney disease 2 {IMAGE:775726}	775726	IMAGE Consortium AA275686
2043 A	5	021NA1GK IMAGE EST {IMAGE:775983}	775983	IMAGE Consortium AA286053
2044 A	7	021NA1GK IMAGE EST {IMAGE:776181}	776181	IMAGE Consortium AA276424
2045 A	9	021NA1GK IMAGE EST {IMAGE:764663}	764663	IMAGE Consortium AA273386
2046 A	11	021NA1GK IMAGE EST {IMAGE:765216}	765216	IMAGE Consortium AA273281
2047 C	1	021NA1GK Mouse mRNA for AF1q, complete cds {IMAGE:775751}	775751	IMAGE Consortium AA275865
2048 C	3	021NA1GK Major urinary protein 2 {IMAGE:776007}	776007	IMAGE Consortium AA285921
2049 C	5	021NA1GK ESTs, Moderately similar to JAK3 [M.musculus] {IMAGE:776301}	776301	IMAGE Consortium AA276492
2050 C	7	021NA1GK IMAGE EST {IMAGE:765073}	765073	IMAGE Consortium AA274182
2051 C	9	021NA1GK ESTs {IMAGE:765552}	765552	IMAGE Consortium AA275118
2052 C	11	021NA1GK IMAGE EST {IMAGE:765272}	765272	IMAGE Consortium AA273311
2053 E	1	021NA1GK IMAGE EST {IMAGE:764204}	764204	IMAGE Consortium AA285697
2054 E	3	021NA1GK ESTs, Highly similar to HYPOTHETICAL 47.9 KD PROTEIN B0303.3 IN CHROMOSOME III [Caenorhabditis elegans] {IMAGE:776448}	776448	IMAGE Consortium AA286024
2055 E	5	021NA1GK Mus musculus mRNA for DBY RNA helicase {IMAGE:765123}	765123	IMAGE Consortium AA273927
2056 E	7	021NA1GK M.musculus mRNA for selenoprotein P {IMAGE:777018}	777018	IMAGE Consortium AA276440
2057 E	9	021NA1GK Interferon regulatory factor 1 {IMAGE:777393}	777393	IMAGE Consortium AA276697
2058 E	11	021NA1GK Trefoil factor 3, intestinal {IMAGE:762404}	762404	IMAGE Consortium AA273366
2059 G	1	021NA1GK ESTs, Weakly similar to HYPOTHETICAL 22.6 KD PROTEIN F46C5.8 IN CHROMOSOME II [C.elegans] {IMAGE:765111}	765111	IMAGE Consortium AA273919
2060 G	3	021NA1GK Synaptonemal complex protein 3 {IMAGE:777467}	777467	IMAGE Consortium AA276721
2061 G	5	021NA1GK IMAGE EST {IMAGE:762476}	762476	IMAGE Consortium AA273673
2062 G	7	021NA1GK IMAGE EST {IMAGE:766102}	766102	IMAGE Consortium AA275581
2063 G	9	021NA1GK IMAGE EST {IMAGE:766447}	766447	IMAGE Consortium AA278027
2064 G	11	021NA1GK Glutathione-S-transferase, alpha 3 {IMAGE:766582}	766582	IMAGE Consortium AA274682
2065 A	1	021UA1GL ESTs {IMAGE:777204}	777204	IMAGE Consortium AA276769
2066 A	3	021UA1GL IMAGE EST {IMAGE:777379}	777379	IMAGE Consortium AA277040
2067 A	5	021UA1GL ESTs {IMAGE:765727}	765727	IMAGE Consortium AA277084
2068 A	7	021UA1GL Glycerol kinase {IMAGE:766704}	766704	IMAGE Consortium AA275564
2069 A	9	021UA1GL ESTs {IMAGE:766567}	766567	IMAGE Consortium AA274729
2070 A	11	021UA1GL IMAGE EST {IMAGE:763352}	763352	IMAGE Consortium AA272458
2071 C	1	021UA1GL IMAGE EST {IMAGE:777101}	777101	IMAGE Consortium AA276365
2072 C	3	021UA1GL ESTs {IMAGE:777403}	777403	IMAGE Consortium AA276579
2073 C	5	021UA1GL Mus musculus myotubularin (Mtm1) mRNA, complete cds {IMAGE:765871}	765871	IMAGE Consortium AA277909

2074 C	7 021UA1GL	ESTs, Moderately similar to HYPOTHETICAL 36.7 KD PROTEIN AH6.2 IN CHROMOSOME II [Caenorhabditis elegans] {IMAGE:766441}	766441 IMAGE Consortium	AA278024
2075 C	9 021UA1GL	Hydroxysteroid dehydrogenase-4, delta<5>-3-beta {IMAGE:766591}	766591 IMAGE Consortium	AA274685
2076 C	11 021UA1GL	ESTs {IMAGE:775800}	775800 IMAGE Consortium	AA275957
2077 E	1 021UA1GL	IMAGE EST {IMAGE:764996}	764996 IMAGE Consortium	AA273425
2078 E	3 021UA1GL	ESTs, Highly similar to P63 protein [H.sapiens] {IMAGE:765338}	765338 IMAGE Consortium	AA275163
2079 E	5 021UA1GL	ESTs {IMAGE:777475}	777475 IMAGE Consortium	AA276537
2080 E	7 021UA1GL	ESTs, Weakly similar to HYPOTHETICAL 23.6 KD PROTEIN IN YUH1-URA8 INTERGENIC REGION [S.cerevisiae] {IMAGE:766498}	766498 IMAGE Consortium	AA275696
2081 E	9 021UA1GL	Mus musculus smooth muscle LIM protein (Crp2/SmLim) mRNA, complete cds {IMAGE:763424}	763424 IMAGE Consortium	AA272260
2082 E	11 021UA1GL	ESTs {IMAGE:775603}	775603 IMAGE Consortium	AA275781
2083 G	1 021UA1GL	Small inducible cytokine A12 {IMAGE:777824}	777824 IMAGE Consortium	AA286393
2084 G	3 021UA1GL	ESTs {IMAGE:777592}	777592 IMAGE Consortium	AA276996
2085 G	5 021UA1GL	ESTs, Highly similar to putative 2-component regulator, interaction with sigma 54 [E.coli] {IMAGE:777574}	777574 IMAGE Consortium	AA286546
2086 G	7 021UA1GL	IMAGE EST {IMAGE:818256}	818256 IMAGE Consortium	AA435376
2087 G	9 021UA1GL	ESTs {IMAGE:818419}	818419 IMAGE Consortium	AA434863
2088 G	11 021UA1GL	ESTs {IMAGE:818756}	818756 IMAGE Consortium	AA466665
2089 A	1 0211A1GM	IMAGE EST {IMAGE:809288}	809288 IMAGE Consortium	AA435038
2090 A	3 0211A1GM	Catenin alpha 1 {IMAGE:818508}	818508 IMAGE Consortium	AA434955
2091 A	5 0211A1GM	ESTs, Highly similar to pM5 [H.sapiens] {IMAGE:818915}	818915 IMAGE Consortium	AA434908
2092 A	7 0211A1GM	Sialyltransferase 8 (alpha-2, 8-sialyltransferase) A {IMAGE:820155}	820155 IMAGE Consortium	AA423143
2093 A	9 0211A1GM	IMAGE EST {IMAGE:820335}	820335 IMAGE Consortium	AA423199
2094 A	11 0211A1GM	Growth arrest specific 2 {IMAGE:820540}	820540 IMAGE Consortium	AA423395
2095 C	1 0211A1GM	Secreted frizzled-related sequence protein 3 {IMAGE:803404}	803404 IMAGE Consortium	AA473070
2096 C	3 0211A1GM	IMAGE EST {IMAGE:803388}	803388 IMAGE Consortium	AA473034
2097 C	5 0211A1GM	IMAGE EST {IMAGE:820359}	820359 IMAGE Consortium	AA423279
2098 C	7 0211A1GM	ESTs {IMAGE:807727}	807727 IMAGE Consortium	AA432892
2099 C	9 0211A1GM	IMAGE EST {IMAGE:807816}	807816 IMAGE Consortium	AA432627
2100 C	11 0211A1GM	ESTs {IMAGE:777655}	777655 IMAGE Consortium	AA276873
2101 E	1 0211A1GM	ESTs {IMAGE:806727}	806727 IMAGE Consortium	AA404079
2102 E	3 0211A1GM	Transcription factor 1 {IMAGE:831701}	831701 IMAGE Consortium	AA467043
2103 E	5 0211A1GM	IMAGE EST {IMAGE:831683}	831683 IMAGE Consortium	AA467030
2104 E	7 0211A1GM	Mus musculus histone H2A.Z (H2A.Z) mRNA, complete cds {IMAGE:831643}	831643 IMAGE Consortium	AA466087
2105 E	9 0211A1GM	ESTs {IMAGE:808213}	808213 IMAGE Consortium	AA433077
2106 E	11 0211A1GM	IMAGE EST {IMAGE:808948}	808948 IMAGE Consortium	AA467710

2107 G	1	0211A1GM	Mus musculus IER5 (Ier5) mRNA, complete cds {IMAGE:808000}	808000	IMAGE Consortium	AA432934
2108 G	3	0211A1GM	ESTs {IMAGE:777651}	777651	IMAGE Consortium	AA276871
2109 G	5	0211A1GM	IMAGE EST {IMAGE:777519}	777519	IMAGE Consortium	AA276746
2110 G	7	0211A1GM	IMAGE EST {IMAGE:777789}	777789	IMAGE Consortium	AA286489
2111 G	9	0211A1GM	ESTs {IMAGE:817939}	817939	IMAGE Consortium	AA466525
2112 G	11	0211A1GM	ESTs, Highly similar to PROTEIN TRANSPORT PROTEIN SEC23 [Saccharomyces cerevisiae] {IMAGE:790571}	790571	IMAGE Consortium	AA290021
2113 A	1	0218A1GN	IMAGE EST {IMAGE:809016}	809016	IMAGE Consortium	AA466026
2114 A	3	0218A1GN	Mus musculus 9ORF binding protein 1 (9BP-1) mRNA, partial cds {IMAGE:809179}	809179	IMAGE Consortium	AA434696
2115 A	5	0218A1GN	ESTs {IMAGE:818324}	818324	IMAGE Consortium	AA435117
2116 A	7	0218A1GN	Discs, large homolog 1 (Drosophila) {IMAGE:818613}	818613	IMAGE Consortium	AA444322
2117 A	9	0218A1GN	Mus musculus Cdk4 and Cdk6 inhibitor p18 protein mRNA, complete cds {IMAGE:820303}	820303	IMAGE Consortium	AA423252
2118 A	11	0218A1GN	Mus musculus E2F-like transcriptional repressor protein mRNA, complete cds {IMAGE:806692}	806692	IMAGE Consortium	AA403531
2119 C	1	0218A1GN	IMAGE EST {IMAGE:818694}	818694	IMAGE Consortium	AA466569
2120 C	3	0218A1GN	IMAGE EST {IMAGE:820112}	820112	IMAGE Consortium	AA423089
2121 C	5	0218A1GN	IMAGE EST {IMAGE:820334}	820334	IMAGE Consortium	AA423204
2122 C	7	0218A1GN	ESTs, Moderately similar to HYPOTHETICAL 167.8 KD PROTEIN CCE1-CAP1 INTERGENIC REGION [Saccharomyces cerevisiae] {IMAGE:820435}	820435	IMAGE Consortium	AA423325
2123 C	9	0218A1GN	ESTs, Highly similar to GLYCOPROTEIN 25L PRECURSOR [Canis familiaris] {IMAGE:831744}	831744	IMAGE Consortium	AA444943
2124 C	11	0218A1GN	ESTs, Weakly similar to Similarity with the drosophila homeotic protein bithoraxoid [C.elegans] {IMAGE:808086}	808086	IMAGE Consortium	AA444362
2125 E	1	0218A1GN	ESTs {IMAGE:807018}	807018	IMAGE Consortium	AA404083
2126 E	3	0218A1GN	ESTs {IMAGE:807449}	807449	IMAGE Consortium	AA412825
2127 E	5	0218A1GN	ESTs {IMAGE:777166}	777166	IMAGE Consortium	AA276381
2128 E	7	0218A1GN	ESTs {IMAGE:777797}	777797	IMAGE Consortium	AA286383
2129 E	9	0218A1GN	ESTs {IMAGE:777539}	777539	IMAGE Consortium	AA276752
2130 E	11	0218A1GN	ESTs {IMAGE:777738}	777738	IMAGE Consortium	AA286274
2131 G	1	0218A1GN	IMAGE EST {IMAGE:777821}	777821	IMAGE Consortium	AA286399
2132 G	3	0218A1GN	ESTs, Moderately similar to NUCLEAR PORE COMPLEX PROTEIN NUP107 [R.norvegicus] {IMAGE:777779}	777779	IMAGE Consortium	AA286352
2133 G	5	0218A1GN	IMAGE EST {IMAGE:777858}	777858	IMAGE Consortium	AA286515
2134 G	7	0218A1GN	ESTs {IMAGE:818435}	818435	IMAGE Consortium	AA434737
2135 G	9	0218A1GN	IMAGE EST {IMAGE:803353}	803353	IMAGE Consortium	AA473022
2136 G	11	0218A1GN	CCAAT/enhancer binding protein (C/EBP), delta {IMAGE:820376}	820376	IMAGE Consortium	AA423439
2137 A	1	021FA1GO	Moloney leukemia virus 10 {IMAGE:808819}	808819	IMAGE Consortium	AA467518
2138 A	3	021FA1GO	IMAGE EST {IMAGE:809105}	809105	IMAGE Consortium	AA435436
2139 A	5	021FA1GO	IMAGE EST {IMAGE:817942}	817942	IMAGE Consortium	AA466515



2140 A	7 021FA1GO	IMAGE EST {IMAGE:818883}	818883 IMAGE Consortium	AA434889
2141 A	9 021FA1GO	IMAGE EST {IMAGE:820158}	820158 IMAGE Consortium	AA423116
2142 A	11 021FA1GO	Pyruvate dehydrogenase E1alpha subunit {IMAGE:820409}	820409 IMAGE Consortium	AA423301
2143 C	1 021FA1GO			
2144 C	3 021FA1GO			
2145 C	5 021FA1GO			
2146 C	7 021FA1GO			
2147 C	9 021FA1GO			
2148 C	11 021FA1GO			
2149 E	1 021FA1GO			
2150 E	3 021FA1GO			
2151 E	5 021FA1GO			
2152 E	7 021FA1GO			
2153 E	9 021FA1GO			
2154 E	11 021FA1GO			
2155 G	1 021FA1GO			
2156 G	3 021FA1GO			
2157 G	5 021FA1GO			
2158 G	7 021FA1GO			
2159 G	9 021FA1GO			
2160 G	11 021FA1GO			
2161 A	1 021MA1GP	Mus musculus FLI-LRR associated protein-1 mRNA, complete cds {IMAGE:532888}	532888 IMAGE Consortium	AA068950
2162 A	3 021MA1GP	Tetratricopeptide repeat domain {IMAGE:524364}	524364 IMAGE Consortium	AA065714
2163 A	5 021MA1GP	ESTs, Moderately similar to KIAA0281 [H.sapiens] {IMAGE:535388}	535388 IMAGE Consortium	AA087542
2164 A	7 021MA1GP	IMAGE EST {IMAGE:405994}	405994 IMAGE Consortium	W83771
2165 A	9 021MA1GP	ESTs, Moderately similar to LIPOIC ACID SYNTHETASE PRECURSOR [Saccharomyces cerevisiae] {IMAGE:525104}	525104 IMAGE Consortium	AA097093
2166 A	11 021MA1GP	Mus musculus cdc2/CDC28-like protein kinase 3 (Clk3) mRNA, complete cds {IMAGE:536695}	536695 IMAGE Consortium	AA080032
2167 C	1 021MA1GP	IMAGE EST {IMAGE:536577}	536577 IMAGE Consortium	AA073904
2168 C	3 021MA1GP	ESTs, Weakly similar to cell cycle progression restoration 8 protein [H.sapiens] {IMAGE:532961}	532961 IMAGE Consortium	AA068968
2169 C	5 021MA1GP	IMAGE EST {IMAGE:522979}	522979 IMAGE Consortium	AA067215
2170 C	7 021MA1GP	ESTs {IMAGE:524078}	524078 IMAGE Consortium	AA066128
2171 C	9 021MA1GP	IMAGE EST {IMAGE:524086}	524086 IMAGE Consortium	AA066250
2172 C	11 021MA1GP	ESTs, Weakly similar to Rab8-interacting protein [M.musculus] {IMAGE:524435}	524435 IMAGE Consortium	AA068568
2173 E	1 021MA1GP	ESTs, Weakly similar to predicted using Genefinder [C.elegans] {IMAGE:524459}	524459 IMAGE Consortium	AA068575
2174 E	3 021MA1GP	Procollagen, type I, alpha 1 {IMAGE:536306}	536306 IMAGE Consortium	AA073604
2175 E	5 021MA1GP	IMAGE EST {IMAGE:536030}	536030 IMAGE Consortium	AA073247

2176 E	7	021MA1GP	NAD(P)H menadione oxidoreductase 1, dioxin inducible {IMAGE:532186}	532186	IMAGE Consortium	AA068375
2177 E	9	021MA1GP	ESTs, Weakly similar to KIAA0005 [H.sapiens] {IMAGE:532659}	532659	IMAGE Consortium	AA068436
2178 E	11	021MA1GP	IMAGE EST {IMAGE:522544}	522544	IMAGE Consortium	AA067977
2179 G	1	021MA1GP	Caspase 1 {IMAGE:550766}	550766	IMAGE Consortium	AA098139
2180 G	3	021MA1GP	ESTs, Highly similar to KERATIN, TYPE II MICROFIBRILLAR, COMPONENT 7C [Ovis aries] {IMAGE:522485}	522485	IMAGE Consortium	AA067967
2181 G	5	021MA1GP	Calumenin {IMAGE:523088}	523088	IMAGE Consortium	AA067258
2182 G	7	021MA1GP	IMAGE EST {IMAGE:523761}	523761	IMAGE Consortium	AA067131
2183 G	9	021MA1GP	IMAGE EST {IMAGE:524392}	524392	IMAGE Consortium	AA067937
2184 G	11	021MA1GP	IMAGE EST {IMAGE:535625}	535625	IMAGE Consortium	AA110866
2185 A	1	021TA1GQ	ESTs {IMAGE:524638}	524638	IMAGE Consortium	AA068486
2186 A	3	021TA1GQ	IMAGE EST {IMAGE:550649}	550649	IMAGE Consortium	AA097946
2187 A	5	021TA1GQ	Stromal cell derived factor 1 {IMAGE:533003}	533003	IMAGE Consortium	AA068750
2188 A	7	021TA1GQ	IMAGE EST {IMAGE:523018}	523018	IMAGE Consortium	AA066425
2189 A	9	021TA1GQ	IMAGE EST {IMAGE:534006}	534006	IMAGE Consortium	AA071829
2190 A	11	021TA1GQ	IMAGE EST {IMAGE:535288}	535288	IMAGE Consortium	AA087768
2191 C	1	021TA1GQ	Tyrosine 3-monooxygenase/tryptophan 5-monooxygenase activatioprotein, epsilon polypeptide {IMAGE:536619}	536619	IMAGE Consortium	AA080011
2192 C	3	021TA1GQ	ESTs {IMAGE:536634}	536634	IMAGE Consortium	AA080156
2193 C	5	021TA1GQ	ESTs {IMAGE:550700}	550700	IMAGE Consortium	AA097957
2194 C	7	021TA1GQ	IMAGE EST {IMAGE:523463}	523463	IMAGE Consortium	AA080272
2195 C	9	021TA1GQ	Alkaline phosphatase 2, liver {IMAGE:535409}	535409	IMAGE Consortium	AA087559
2196 C	11	021TA1GQ	Ribosomal protein L7 {IMAGE:524442}	524442	IMAGE Consortium	AA067910
2197 E	1	021TA1GQ	Amyloid beta (A4) precursor protein {IMAGE:535652}	535652	IMAGE Consortium	AA110872
2198 E	3	021TA1GQ	ESTs {IMAGE:550680}	550680	IMAGE Consortium	AA098210
2199 E	5	021TA1GQ	ESTs, Highly similar to LEUCOCYTE ANTIGEN CD97 PRECURSOR [Homo sapiens] {IMAGE:551243}	551243	IMAGE Consortium	AA080529
2200 E	7	021TA1GQ	ESTs {IMAGE:522549}	522549	IMAGE Consortium	AA067999
2201 E	9	021TA1GQ	ESTs, Highly similar to PHOSPHOENOLPYRUVATE CARBOXYKINASE, CYTOSOLIC [Rattus norvegicus] {IMAGE:523400}	523400	IMAGE Consortium	AA080172
2202 E	11	021TA1GQ	Fas-associated factor 1 {IMAGE:534132}	534132	IMAGE Consortium	AA072606
2203 G	1	021TA1GQ	Mus musculus Btk locus, alpha-D-galactosidase A (Ags), ribosomal protein (L44L), and Bruton's tyrosine kinase (Btk) genes, complete cds {IMAGE:437469}	437469	IMAGE Consortium	AA003549
2204 G	3	021TA1GQ	IMAGE EST {IMAGE:656934}	656934	IMAGE Consortium	AA240365
2205 G	5	021TA1GQ	IMAGE EST {IMAGE:661317}	661317	IMAGE Consortium	AA207613
2206 G	7	021TA1GQ	ESTs {IMAGE:651532}	651532	IMAGE Consortium	AA212455
2207 G	9	021TA1GQ	ESTs {IMAGE:656984}	656984	IMAGE Consortium	AA240645
2208 G	11	021TA1GQ	ESTs {IMAGE:651293}	651293	IMAGE Consortium	AA213360

2209 A	1 0210A1GR	IMAGE EST {IMAGE:662634}	662634 IMAGE Consortium	AA208066
2210 A	3 0210A1GR	IMAGE EST {IMAGE:652259}	652259 IMAGE Consortium	AA215107
2211 A	5 0210A1GR	ESTs, Highly similar to 80 KD NUCLEAR CAP BINDING PROTEIN [H.sapiens] {IMAGE:633370}	633370 IMAGE Consortium	AA184580
2212 A	7 0210A1GR	Mus musculus CDK-activating kinase assembly factor p36/MAT1, complete cds {IMAGE:651834}	651834 IMAGE Consortium	AA212256
2213 A	9 0210A1GR	Nucleoside phosphorylase {IMAGE:607469}	607469 IMAGE Consortium	AA172867
2214 A	11 0210A1GR	IMAGE EST {IMAGE:656204}	656204 IMAGE Consortium	AA230500
2215 C	1 0210A1GR	ESTs, Moderately similar to CLEAVAGE STIMULATION FACTOR, 50 KD SUBUNIT [Homo sapiens] {IMAGE:661491}	661491 IMAGE Consortium	AA207759
2216 C	3 0210A1GR	IMAGE EST {IMAGE:732889}	732889 IMAGE Consortium	AA286159
2217 C	5 0210A1GR	IMAGE EST {IMAGE:918177}	918177 IMAGE Consortium	AA492651
2218 C	7 0210A1GR	IMAGE EST {IMAGE:656325}	656325 IMAGE Consortium	AA241251
2219 C	9 0210A1GR	IMAGE EST {IMAGE:732290}	732290 IMAGE Consortium	AA276554
2220 C	11 0210A1GR	Mus musculus protein phosphatase 2A B'alpha3 regulatory subunit mRNA, partial cds {IMAGE:652191}	652191 IMAGE Consortium	AA215001
2221 E	1 0210A1GR	IMAGE EST {IMAGE:633284}	633284 IMAGE Consortium	AA177264
2222 E	3 0210A1GR	ESTs, Highly similar to FRUCTOSE-1,6-BISPHOSPHATASE [Rattus norvegicus] {IMAGE:776124}	776124 IMAGE Consortium	AA276043
2223 E	5 0210A1GR	IMAGE EST {IMAGE:633838}	633838 IMAGE Consortium	AA199051
2224 E	7 0210A1GR	ESTs {IMAGE:661821}	661821 IMAGE Consortium	AA208426
2225 E	9 0210A1GR	IMAGE EST {IMAGE:651825}	651825 IMAGE Consortium	AA212204
2226 E	11 0210A1GR	SPERM MITOCHONDRIAL CAPSULE SELENOPROTEIN {IMAGE:917403}	917403 IMAGE Consortium	AA497767
2227 G	1 0210A1GR	Esterase 1 {IMAGE:733022}	733022 IMAGE Consortium	AA403730
2228 G	3 0210A1GR	ESTs, Moderately similar to DOLICHYL-PHOSPHATE BETA-GLUCOSYLTRANSFERASE [Saccharomyces cerevisiae] {IMAGE:657077}	657077 IMAGE Consortium	AA241340
2229 G	5 0210A1GR	IMAGE EST {IMAGE:662072}	662072 IMAGE Consortium	AA207292
2230 G	7 0210A1GR	ESTs, Weakly similar to p40 [H.sapiens] {IMAGE:732999}	732999 IMAGE Consortium	AA403723
2231 G	9 0210A1GR	IMAGE EST {IMAGE:656601}	656601 IMAGE Consortium	AA240420
2232 G	11 0210A1GR			
2233 A	1 021Q9974	Control: Ratio 1:3 (Cy3:Cy5)	Ratio_A	Controls
2234 A	3 021Q9974	Control: Ratio 3:1 (Cy3:Cy5)	Ratio_B	Controls
2235 A	5 021Q9974	Control: Ratio 1:10 (Cy3:Cy5)	Ratio_C	Controls
2236 A	7 021Q9974	Control: Ratio 10:1 (Cy3:Cy5)	Ratio_D	Controls
2237 A	9 021Q9974	Control: Ratio 1:25 (Cy3:Cy5)	Ratio_E	Controls
2238 A	11 021Q9974	Control: Ratio 25:1 (Cy3:Cy5)	Ratio_F	Controls
2239 C	1 021Q9974	Control: Sensitivity 2pg	Sensitivity_A	Controls
2240 C	3 021Q9974	Control: Sensitivity 20pg	Sensitivity_B	Controls
2241 C	5 021Q9974	Control: Sensitivity 200pg	Sensitivity_C	Controls
2242 C	7 021Q9974	Control: Sensitivity 2000pg	Sensitivity_D	Controls
2243 C	9 021Q9974	Control: Buffer Only	Buffer_Only_Control	Controls

2244 C	11	021Q9974	Control: Buffer Only	Buffer_Only_Control	Controls		
2245 E	1	021Q9974	Control: Internal	Internal_Control_A	Controls		
2246 E	3	021Q9974	Control: Buffer Only	Buffer_Only_Control	Controls		
2247 E	5	021Q9974	Control: Internal	Internal_Control_B	Controls		
2248 E	7	021Q9974	Control: Buffer Only	Buffer_Only_Control	Controls		
2249 E	9	021Q9974	Control: Internal	Internal_Control_C	Controls		
2250 E	11	021Q9974	Control: Internal	Internal_Control_D	Controls		
2251 G	1	021Q9974	Control: Complex Target (Mus musculus)	RNA_Quality_MM	Controls		
2252 G	3	021Q9974	Control: Complex Target (Mus musculus)	RNA_Quality_MM	Controls		
2253 G	5	021Q9974	Control: Complex Target (Mus musculus)	RNA_Quality_MM	Controls		
2254 G	7	021Q9974	Control: Housekeeping (Alpha Tubulin)	Housekeeping_B	Controls		
2255 G	9	021Q9974	Control: Housekeeping (Ribosomal S9)	Housekeeping_A	Controls		
2256 G	11	021Q9974	Control: Housekeeping (23kD HBP)	Housekeeping_C	Controls		
2257 A	2	2159971	Internal_Control_E	Internal_Control_E	Controls		
2258 A	4	2159971	Internal_Control_F	Internal_Control_F	Controls		
2259 A	6	2159971	Internal_Control_G	Internal_Control_G	Controls		
2260 A	8	2159971	Internal_Control_H	Internal_Control_H	Controls		
2261 A	10	2159971	Internal_Control_I	Internal_Control_I	Controls		
2262 A	12	2159971	Internal_Control_J	Internal_Control_J	Controls		
2263 C	2	2159971	Internal_Control_K	Internal_Control_K	Controls		
2264 C	4	2159971	Internal_Control_L	Internal_Control_L	Controls		
2265 C	6	2159971	Internal_Control_M	Internal_Control_M	Controls		
2266 C	8	2159971	Internal_Control_N	Internal_Control_N	Controls		
2267 C	10	2159971	Internal_Control_O	Internal_Control_O	Controls		
2268 C	12	2159971	Internal_Control_P	Internal_Control_P	Controls		
2269 E	2	2159971	Internal_Control_Q	Internal_Control_Q	Controls		
2270 E	4	2159971	Internal_Control_R	Internal_Control_R	Controls		
2271 E	6	2159971	Internal_Control_S	Internal_Control_S	Controls		
2272 E	8	2159971	Internal_Control_T	Internal_Control_T	Controls		
2273 E	10	2159971	Internal_Control_U	Internal_Control_U	Controls		
2274 E	12	2159971	Internal_Control_V	Internal_Control_V	Controls		
2275 G	2	2159971	Internal_Control_W	Internal_Control_W	Controls		
2276 G	4	2159971	Internal_Control_X	Internal_Control_X	Controls		
2277 G	6	2159971	Internal_Control_Y	Internal_Control_Y	Controls		
2278 G	8	2159971	Internal_Control_Z	Internal_Control_Z	Controls		
2279 G	10	2159971	Internal_Control_AA	Internal_Control_AA	Controls		
2280 G	12	2159971	Internal_Control_AB	Internal_Control_AB	Controls		
2281 A	2	021MA1E8	IMAGE EST {IMAGE:933183}	933183	IMAGE Consortium	AA538218	
2282 A	4	021MA1E8	IMAGE EST {IMAGE:934063}	934063	IMAGE Consortium	AA544565	
2283 A	6	021MA1E8	ESTs {IMAGE:934635}	934635	IMAGE Consortium	AA536838	
2284 A	8	021MA1E8	ESTs {IMAGE:833601}	833601	IMAGE Consortium	AA432967	
2285 A	10	021MA1E8	IMAGE EST {IMAGE:834840}	834840	IMAGE Consortium	AA467180	

2286 A	12 021MA1E8	IMAGE EST {IMAGE:834712}	834712 IMAGE Consortium	AA467053
2287 C	2 021MA1E8	ESTs, Highly similar to PUTATIVE ASPARAGINYL-TRNA SYNTHETASE DED81 [ <i>Saccharomyces cerevisiae</i> ] {IMAGE:805842}	805842 IMAGE Consortium	AA399854
2288 C	4 021MA1E8	MutS homolog 2 ( <i>E. coli</i> ) {IMAGE:945359}	945359 IMAGE Consortium	AA546673
2289 C	6 021MA1E8	ESTs {IMAGE:834934}	834934 IMAGE Consortium	AA467267
2290 C	8 021MA1E8	IMAGE EST {IMAGE:894372}	894372 IMAGE Consortium	AA517431
2291 C	10 021MA1E8	IMAGE EST {IMAGE:761178}	761178 IMAGE Consortium	AA387817
2292 C	12 021MA1E8	IMAGE EST {IMAGE:761302}	761302 IMAGE Consortium	AA388323
2293 E	2 021MA1E8	Sine oculis-related homeobox 3, ( <i>Drosophila</i> ) {IMAGE:761326}	761326 IMAGE Consortium	AA388136
2294 E	4 021MA1E8	ESTs, Weakly similar to protein kinase [ <i>M.musculus</i> ] {IMAGE:933770}	933770 IMAGE Consortium	AA545690
2295 E	6 021MA1E8	ESTs {IMAGE:933852}	933852 IMAGE Consortium	AA544325
2296 E	8 021MA1E8	Guanine nucleotide binding protein, related sequence 1 {IMAGE:945010}	945010 IMAGE Consortium	AA537232
2297 E	10 021MA1E8	ESTs, Highly similar to UDP-GLUCOSE 4-EPIMERASE [ <i>Rattus norvegicus</i> ] {IMAGE:760889}	760889 IMAGE Consortium	AA386807
2298 E	12 021MA1E8	IMAGE EST {IMAGE:761213}	761213 IMAGE Consortium	AA387625
2299 G	2 021MA1E8	IMAGE EST {IMAGE:932999}	932999 IMAGE Consortium	AA538199
2300 G	4 021MA1E8	ESTs, Moderately similar to EPIDERMAL GROWTH FACTOR RECEPTOR SUBSTRATE SUBSTRATE 15 [ <i>Homo sapiens</i> ] {IMAGE:804453}	804453 IMAGE Consortium	AA474757
2301 G	6 021MA1E8	ESTs {IMAGE:805496}	805496 IMAGE Consortium	AA474600
2302 G	8 021MA1E8	ESTs, Weakly similar to predicted using Genefinder [ <i>C.elegans</i> ] {IMAGE:945059}	945059 IMAGE Consortium	AA537323
2303 G	10 021MA1E8	ESTs {IMAGE:834262}	834262 IMAGE Consortium	AA413159
2304 G	12 021MA1E8	ESTs, Moderately similar to The ha1438 gene product is related to a C728 protein encoded in <i>S.cerevisiae</i> chromosome II. [ <i>H.sapiens</i> ] {IMAGE:804482}	804482 IMAGE Consortium	AA474342
2305 A	2 021TA1E9	<i>Mus musculus</i> aromatic-L-amino-acid decarboxylase mRNA, complete cds {IMAGE:804722}	804722 IMAGE Consortium	AA474389
2306 A	4 021TA1E9	IMAGE EST {IMAGE:805121}	805121 IMAGE Consortium	AA397275
2307 A	6 021TA1E9	ESTs, Weakly similar to unknown [ <i>S.cerevisiae</i> ] {IMAGE:805547}	805547 IMAGE Consortium	AA397252
2308 A	8 021TA1E9	ESTs {IMAGE:944942}	944942 IMAGE Consortium	AA537095
2309 A	10 021TA1E9	ESTs, Moderately similar to p53BP2 [ <i>M.musculus</i> ] {IMAGE:834947}	834947 IMAGE Consortium	AA467287
2310 A	12 021TA1E9	<i>M.musculus</i> mRNA for myosin I heavy chain {IMAGE:761018}	761018 IMAGE Consortium	AA387258
2311 C	2 021TA1E9	IMAGE EST {IMAGE:893933}	893933 IMAGE Consortium	AA517145
2312 C	4 021TA1E9	ESTs {IMAGE:761066}	761066 IMAGE Consortium	AA387558
2313 C	6 021TA1E9	Est2 repressor factor {IMAGE:894472}	894472 IMAGE Consortium	AA517533
2314 C	8 021TA1E9	ESTs, Weakly similar to GOLIATH PROTEIN [ <i>Drosophila melanogaster</i> ] {IMAGE:804680}	804680 IMAGE Consortium	AA474374

2315 C	10 021TA1E9	Sjogren syndrome antigen B {IMAGE:933844}	933844 IMAGE Consortium	AA544321
2316 C	12 021TA1E9	ESTs, Highly similar to UTP--GLUCOSE-1-PHOSPHATE URIDYLTRANSFERASE [Homo sapiens] {IMAGE:805218}	805218 IMAGE Consortium	AA473123
2317 E	2 021TA1E9	ESTs {IMAGE:805386}	805386 IMAGE Consortium	AA473205
2318 E	4 021TA1E9	ESTs {IMAGE:833814}	833814 IMAGE Consortium	AA412914
2319 E	6 021TA1E9	IMAGE EST {IMAGE:834644}	834644 IMAGE Consortium	AA467549
2320 E	8 021TA1E9	M.musculus es64 mRNA {IMAGE:760928}	760928 IMAGE Consortium	AA386704
2321 E	10 021TA1E9	IMAGE EST {IMAGE:761264}	761264 IMAGE Consortium	AA388221
2322 E	12 021TA1E9	ESTs {IMAGE:761038}	761038 IMAGE Consortium	AA387579
2323 G	2 021TA1E9	ESTs, Weakly similar to ZINC FINGER PROTEIN HF.12 [Homo sapiens] {IMAGE:761019}	761019 IMAGE Consortium	AA387262
2324 G	4 021TA1E9	IMAGE EST {IMAGE:761062}	761062 IMAGE Consortium	AA387586
2325 G	6 021TA1E9	IMAGE EST {IMAGE:775083}	775083 IMAGE Consortium	AA387050
2326 G	8 021TA1E9	ESTs, Highly similar to TENSIN [Gallus gallus] {IMAGE:774810}	774810 IMAGE Consortium	AA386710
2327 G	10 021TA1E9	IMAGE EST {IMAGE:775235}	775235 IMAGE Consortium	AA387210
2328 G	12 021TA1E9	IMAGE EST {IMAGE:774996}	774996 IMAGE Consortium	AA387314
2329 A	2 021DA1EA	Mus musculus copper transport protein Atox1 (ATOX1) mRNA, complete cds {IMAGE:791264}	791264 IMAGE Consortium	AA416325
2330 A	4 021DA1EA	IMAGE EST {IMAGE:791058}	791058 IMAGE Consortium	AA415240
2331 A	6 021DA1EA	IMAGE EST {IMAGE:791618}	791618 IMAGE Consortium	AA437580
2332 A	8 021DA1EA	ESTs {IMAGE:835536}	835536 IMAGE Consortium	AA422515
2333 A	10 021DA1EA	IMAGE EST {IMAGE:792610}	792610 IMAGE Consortium	AA415228
2334 A	12 021DA1EA	ESTs {IMAGE:779143}	779143 IMAGE Consortium	AA467571
2335 C	2 021DA1EA	IMAGE EST {IMAGE:804331}	804331 IMAGE Consortium	AA474849
2336 C	4 021DA1EA	Growth differentiation factor 9 {IMAGE:835757}	835757 IMAGE Consortium	AA422377
2337 C	6 021DA1EA	IMAGE EST {IMAGE:792735}	792735 IMAGE Consortium	AA416438
2338 C	8 021DA1EA	ESTs, Moderately similar to HYPOTHETICAL 49.7 KD PROTEIN IN GIN2-STE3 INTERGENIC REGION [Saccharomyces cerevisiae] {IMAGE:779257}	779257 IMAGE Consortium	AA445775
2339 C	10 021DA1EA	IMAGE EST {IMAGE:793494}	793494 IMAGE Consortium	AA415915
2340 C	12 021DA1EA	IMAGE EST {IMAGE:780127}	780127 IMAGE Consortium	AA414780
2341 E	2 021DA1EA	IMAGE EST {IMAGE:803684}	803684 IMAGE Consortium	AA422830
2342 E	4 021DA1EA	ESTs, Weakly similar to Zic2 protein [M.musculus] {IMAGE:791455}	791455 IMAGE Consortium	AA416308
2343 E	6 021DA1EA	Bcl2-like 10 {IMAGE:836487}	836487 IMAGE Consortium	AA426964
2344 E	8 021DA1EA	ESTs {IMAGE:792886}	792886 IMAGE Consortium	AA415370
2345 E	10 021DA1EA	Mus musculus capping protein alpha 1 subunit mRNA, partial cds {IMAGE:779754}	779754 IMAGE Consortium	AA414612
2346 E	12 021DA1EA	IMAGE EST {IMAGE:791503}	791503 IMAGE Consortium	AA437432
2347 G	2 021DA1EA	GUANINE NUCLEOTIDE-BINDING PROTEIN, ALPHA-13 SUBUNIT {IMAGE:779611}	779611 IMAGE Consortium	AA466926
2348 G	4 021DA1EA	ESTs, Highly similar to torsinA [H.sapiens] {IMAGE:803613}	803613 IMAGE Consortium	AA422787

2349 G	6 021DA1EA	IMAGE EST {IMAGE:833472}	833472 IMAGE Consortium	AA465966
2350 G	8 021DA1EA	IMAGE EST {IMAGE:833346}	833346 IMAGE Consortium	AA432472
2351 G	10 021DA1EA	IMAGE EST {IMAGE:792987}	792987 IMAGE Consortium	AA415475
2352 G	12 021DA1EA	IMAGE EST {IMAGE:959839}	959839 IMAGE Consortium	AA547022
2353 A	2 021KA1EB	IMAGE EST {IMAGE:775495}	775495 IMAGE Consortium	AA388274
2354 A	4 021KA1EB	ESTs {IMAGE:775484}	775484 IMAGE Consortium	AA387243
2355 A	6 021KA1EB	ESTs, Highly similar to Gene product with similarity to Ubiquitin binding enzyme [H.sapiens] {IMAGE:790140}	790140 IMAGE Consortium	AA388099
2356 A	8 021KA1EB	IMAGE EST {IMAGE:774806}	774806 IMAGE Consortium	AA386744
2357 A	10 021KA1EB	ESTs {IMAGE:775538}	775538 IMAGE Consortium	AA388304
2358 A	12 021KA1EB	IMAGE EST {IMAGE:775407}	775407 IMAGE Consortium	AA387340
2359 C	2 021KA1EB	IMAGE EST {IMAGE:774808}	774808 IMAGE Consortium	AA386709
2360 C	4 021KA1EB	ESTs, Weakly similar to GRAVIN [Homo sapiens] {IMAGE:775253}	775253 IMAGE Consortium	AA387076
2361 C	6 021KA1EB	Mus musculus phosphomannomutase Sec53p homolog mRNA, complete cds {IMAGE:775067}	775067 IMAGE Consortium	AA387369
2362 C	8 021KA1EB	IMAGE EST {IMAGE:789694}	789694 IMAGE Consortium	AA387537
2363 C	10 021KA1EB	IMAGE EST {IMAGE:775121}	775121 IMAGE Consortium	AA386889
2364 C	12 021KA1EB	ESTs, Weakly similar to ZK856.11 [C.elegans] {IMAGE:778381}	778381 IMAGE Consortium	AA413555
2365 E	2 021KA1EB	IMAGE EST {IMAGE:779744}	779744 IMAGE Consortium	AA414028
2366 E	4 021KA1EB	IMAGE EST {IMAGE:779797}	779797 IMAGE Consortium	AA414720
2367 E	6 021KA1EB	IMAGE EST {IMAGE:803920}	803920 IMAGE Consortium	AA435261
2368 E	8 021KA1EB	ESTs {IMAGE:791765}	791765 IMAGE Consortium	AA414642
2369 E	10 021KA1EB	ESTs, Weakly similar to HYPOTHETICAL 40.4 KD PROTEIN R06F6.5 IN CHROMOSOME II [Caenorhabditis elegans] {IMAGE:792870}	792870 IMAGE Consortium	AA415519
2370 E	12 021KA1EB	ESTs {IMAGE:779847}	779847 IMAGE Consortium	AA414733
2371 G	2 021KA1EB	ESTs, Weakly similar to butyrophilin [H.sapiens] {IMAGE:833259}	833259 IMAGE Consortium	AA465838
2372 G	4 021KA1EB	ESTs {IMAGE:833351}	833351 IMAGE Consortium	AA432477
2373 G	6 021KA1EB	ESTs {IMAGE:793218}	793218 IMAGE Consortium	AA437891
2374 G	8 021KA1EB	ESTs, Weakly similar to HEM45 [H.sapiens] {IMAGE:793347}	793347 IMAGE Consortium	AA438241
2375 G	10 021KA1EB	ESTs, Highly similar to RSP5 PROTEIN [Saccharomyces cerevisiae] {IMAGE:779451}	779451 IMAGE Consortium	AA414211
2376 G	12 021KA1EB	IMAGE EST {IMAGE:779824}	779824 IMAGE Consortium	AA414788
2377 A	2 021RA1EC	Protein kinase, cAMP dependent, catalytic, beta {IMAGE:961171}	961171 IMAGE Consortium	AA547428
2378 A	4 021RA1EC	T-cell lymphoma breakpoint 1 {IMAGE:961083}	961083 IMAGE Consortium	AA547208
2379 A	6 021RA1EC	Catenin beta {IMAGE:313426}	313426 IMAGE Consortium	W10023
2380 A	8 021RA1EC	ESTs {IMAGE:318000}	318000 IMAGE Consortium	W11605
2381 A	10 021RA1EC	Fumarylacetoacetate hydrolase {IMAGE:314442}	314442 IMAGE Consortium	W12834
2382 A	12 021RA1EC	Syntrophin, acidic 1 {IMAGE:318957}	318957 IMAGE Consortium	W34029

2383 C	2 021RA1EC	ESTs {IMAGE:317344}	317344 IMAGE Consortium	W34013
2384 C	4 021RA1EC	Myosin heavy chain, skeletal muscle, perinatal {IMAGE:317476}	317476 IMAGE Consortium	W13528
2385 C	6 021RA1EC	Mitsugumin 29 {IMAGE:317610}	317610 IMAGE Consortium	W13114
2386 C	8 021RA1EC	ESTs, Weakly similar to PROBABLE LEUCYL-TRNA SYNTHETASE [C.elegans] {IMAGE:318157}	318157 IMAGE Consortium	W11665
2387 C	10 021RA1EC	Coiled-coil like protein 1 {IMAGE:314850}	314850 IMAGE Consortium	W12503
2388 C	12 021RA1EC	ESTs {IMAGE:329367}	329367 IMAGE Consortium	W42101
2389 E	2 021RA1EC	ESTs {IMAGE:317778}	317778 IMAGE Consortium	W13043
2390 E	4 021RA1EC	ESTs {IMAGE:314479}	314479 IMAGE Consortium	W10632
2391 E	6 021RA1EC	ESTs {IMAGE:318874}	318874 IMAGE Consortium	W34179
2392 E	8 021RA1EC	ESTs, Moderately similar to HYPOTHETICAL 9.3 KD PROTEIN ZK652.1 IN CHROMOSOME III [Caenorhabditis elegans] {IMAGE:315581}	315581 IMAGE Consortium	W09867
2393 E	10 021RA1EC	IMAGE EST {IMAGE:315936}	315936 IMAGE Consortium	W08989
2394 E	12 021RA1EC	ESTs, Highly similar to COP-COATED VESICLE MEMBRANE PROTEIN P24 PRECURSOR [Cricetulus griseus] {IMAGE:316187}	316187 IMAGE Consortium	W14393
2395 G	2 021RA1EC	ESTs, Highly similar to 2-OXOGLUTARATE DEHYDROGENASE E1 COMPONENT PRECURSOR [Homo sapiens] {IMAGE:329728}	329728 IMAGE Consortium	W13320
2396 G	4 021RA1EC	IMAGE EST {IMAGE:330241}	330241 IMAGE Consortium	W13950
2397 G	6 021RA1EC	IMAGE EST {IMAGE:330224}	330224 IMAGE Consortium	W13933
2398 G	8 021RA1EC	ESTs {IMAGE:330613}	330613 IMAGE Consortium	W14186
2399 G	10 021RA1EC	Mus musculus FK506 binding protein (Fkbp63) mRNA, complete cds {IMAGE:317218}	317218 IMAGE Consortium	W13698
2400 G	12 021RA1EC	IMAGE EST {IMAGE:313529}	313529 IMAGE Consortium	W10596
2401 A	2 021YA1ED	ESTs {IMAGE:330661}	330661 IMAGE Consortium	W14247
2402 A	4 021YA1ED	ESTs, Moderately similar to KIAA0182 [H.sapiens] {IMAGE:317338}	317338 IMAGE Consortium	W13229
2403 A	6 021YA1ED	Dynamin {IMAGE:317587}	317587 IMAGE Consortium	W13111
2404 A	8 021YA1ED	ESTs {IMAGE:314576}	314576 IMAGE Consortium	W10852
2405 A	10 021YA1ED	ESTs {IMAGE:318938}	318938 IMAGE Consortium	W34387
2406 A	12 021YA1ED	IMAGE EST {IMAGE:329563}	329563 IMAGE Consortium	W42223
2407 C	2 021YA1ED	ESTs {IMAGE:317863}	317863 IMAGE Consortium	W12913
2408 C	4 021YA1ED	ESTs {IMAGE:314979}	314979 IMAGE Consortium	W12546
2409 C	6 021YA1ED	IMAGE EST {IMAGE:329372}	329372 IMAGE Consortium	W41977
2410 C	8 021YA1ED	Mus musculus secreted carbonic anhydrase isozyme VI precursor, mRNA, complete cds {IMAGE:329940}	329940 IMAGE Consortium	W11772
2411 C	10 021YA1ED	ESTs, Highly similar to kidney-specific transport protein [M.musculus] {IMAGE:330194}	330194 IMAGE Consortium	W11794
2412 C	12 021YA1ED	ESTs, Highly similar to RAS-RELATED PROTEIN RAB-9 [Canis familiaris] {IMAGE:316415}	316415 IMAGE Consortium	W13154
2413 E	2 021YA1ED	Mouse mRNA for preprotachykinin B, complete cds	330108 IMAGE Consortium	W11882



2414 E	4 021YA1ED	{IMAGE:330108} ESTs, Highly similar to DERMATOPONTIN [Bos taurus] {IMAGE:330218}	330218 IMAGE Consortium	W13931
2415 E	6 021YA1ED	ESTs {IMAGE:330595}	330595 IMAGE Consortium	W14214
2416 E	8 021YA1ED	Tight junction protein 1 {IMAGE:317808}	317808 IMAGE Consortium	W13290
2417 E	10 021YA1ED	ESTs {IMAGE:313629}	313629 IMAGE Consortium	W10658
2418 E	12 021YA1ED	ESTs {IMAGE:314377}	314377 IMAGE Consortium	W10683
2419 G	2 021YA1ED	ESTs {IMAGE:318200}	318200 IMAGE Consortium	W11492
2420 G	4 021YA1ED	Hexokinase 1 {IMAGE:318642}	318642 IMAGE Consortium	W11571
2421 G	6 021YA1ED	ESTs, Weakly similar to SPECTRIN BETA CHAIN, ERYTHROCYTE [M.musculus] {IMAGE:330183}	330183 IMAGE Consortium	W11792
2422 G	8 021YA1ED	ESTs {IMAGE:330583}	330583 IMAGE Consortium	W15074
2423 G	10 021YA1ED	ESTs, Highly similar to CATHEPSIN [Bos taurus] {IMAGE:330668}	330668 IMAGE Consortium	W14289
2424 G	12 021YA1ED	ESTs {IMAGE:317537}	317537 IMAGE Consortium	W33877
2425 A	2 0215A1EE	ESTs {IMAGE:334268}	334268 IMAGE Consortium	W36984
2426 A	4 0215A1EE	ESTs, Weakly similar to cDNA EST EMBL:D71020 comes from this gene [C.elegans] {IMAGE:338105}	338105 IMAGE Consortium	W29374
2427 A	6 0215A1EE	IMAGE EST {IMAGE:331424}	331424 IMAGE Consortium	W14960
2428 A	8 0215A1EE	IMAGE EST {IMAGE:331681}	331681 IMAGE Consortium	W14332
2429 A	10 0215A1EE	IMAGE EST {IMAGE:331768}	331768 IMAGE Consortium	W08090
2430 A	12 0215A1EE	Histocompatibility 2, complement component factor B {IMAGE:336159}	336159 IMAGE Consortium	W18121
2431 C	2 0215A1EE	IMAGE EST {IMAGE:331731}	331731 IMAGE Consortium	W14454
2432 C	4 0215A1EE	ESTs, Weakly similar to Similarity to Sheep transthyretin precursor [C.elegans] {IMAGE:331792}	331792 IMAGE Consortium	W08097
2433 C	6 0215A1EE	Untitled {IMAGE:332480}	332480 IMAGE Consortium	W08488
2434 C	8 0215A1EE	IMAGE EST {IMAGE:333042}	333042 IMAGE Consortium	W15636
2435 C	10 0215A1EE	ESTs, Highly similar to phosphoethanolamine cytidyltransferase [H.sapiens] {IMAGE:333669}	333669 IMAGE Consortium	W17420
2436 C	12 0215A1EE	ESTs, Moderately similar to FKHR protein [H.sapiens] {IMAGE:334077}	334077 IMAGE Consortium	W36356
2437 E	2 0215A1EE	ESTs {IMAGE:333418}	333418 IMAGE Consortium	W16354
2438 E	4 0215A1EE	IMAGE EST {IMAGE:333914}	333914 IMAGE Consortium	W16053
2439 E	6 0215A1EE	ESTs, Weakly similar to KIAA0116 [H.sapiens] {IMAGE:337565}	337565 IMAGE Consortium	W29299
2440 E	8 0215A1EE	IMAGE EST {IMAGE:334584}	334584 IMAGE Consortium	W18341
2441 E	10 0215A1EE	IMAGE EST {IMAGE:335018}	335018 IMAGE Consortium	W17914
2442 E	12 0215A1EE	IMAGE EST {IMAGE:331335}	331335 IMAGE Consortium	W14692
2443 G	2 0215A1EE	ESTs, Highly similar to GLUTAMINASE, LIVER ISOFORM PRECURSOR [Rattus norvegicus] {IMAGE:337170}	337170 IMAGE Consortium	W20907
2444 G	4 0215A1EE	ESTs {IMAGE:334514}	334514 IMAGE Consortium	W36597
2445 G	6 0215A1EE	ESTs {IMAGE:348362}	348362 IMAGE Consortium	W33467

2446 G	8 0215A1EE	Mus musculus mRNA for monoglyceride lipase {IMAGE:335220}	335220 IMAGE Consortium	W17582
2447 G	10 0215A1EE	M.musculus mRNA for Ulip protein {IMAGE:335572}	335572 IMAGE Consortium	W18828
2448 G	12 0215A1EE	Mus musculus uterine-specific proline-rich acidic protein mRNA, complete cds {IMAGE:336053}	336053 IMAGE Consortium	W17866
2449 A	2 021CA1EF	Esterase 1 {IMAGE:330447}	330447 IMAGE Consortium	W14805
2450 A	4 021CA1EF	Sphingomyelin phosphodiesterase 1, acid lysosomal {IMAGE:330863}	330863 IMAGE Consortium	W14928
2451 A	6 021CA1EF	IMAGE EST {IMAGE:313686}	313686 IMAGE Consortium	W11055
2452 A	8 021CA1EF	M.musculus mRNA for LPS-binding protein {IMAGE:314215}	314215 IMAGE Consortium	W09930
2453 A	10 021CA1EF	IMAGE EST {IMAGE:314645}	314645 IMAGE Consortium	W10915
2454 A	12 021CA1EF	ESTs {IMAGE:314394}	314394 IMAGE Consortium	W12827
2455 C	2 021CA1EF	IMAGE EST {IMAGE:336458}	336458 IMAGE Consortium	W18362
2456 C	4 021CA1EF	Mouse mRNA for fructose 1,6-bisphosphatase, complete cds {IMAGE:336727}	336727 IMAGE Consortium	W36831
2457 C	6 021CA1EF	PROTEIN PHOSPHATASE 2C ALPHA ISOFORM {IMAGE:337407}	337407 IMAGE Consortium	W18642
2458 C	8 021CA1EF	Procollagen, type VI, alpha 1 {IMAGE:334132}	334132 IMAGE Consortium	W16221
2459 C	10 021CA1EF	Mus musculus fetuin (Ahsg) gene, complete cds {IMAGE:334195}	334195 IMAGE Consortium	W36917
2460 C	12 021CA1EF	ESTs, Weakly similar to apolipoprotein A-IV precursor [R.norvegicus] {IMAGE:338005}	338005 IMAGE Consortium	W18951
2461 E	2 021CA1EF	ESTs {IMAGE:333486}	333486 IMAGE Consortium	AA052356
2462 E	4 021CA1EF	IMAGE EST {IMAGE:333307}	333307 IMAGE Consortium	W15872
2463 E	6 021CA1EF	IMAGE EST {IMAGE:333952}	333952 IMAGE Consortium	W18167
2464 E	8 021CA1EF	ESTs, Weakly similar to cDNA EST EMBL:D75506 comes from this gene [C.elegans] {IMAGE:334182}	334182 IMAGE Consortium	W16247
2465 E	10 021CA1EF	IMAGE EST {IMAGE:334291}	334291 IMAGE Consortium	W36995
2466 E	12 021CA1EF	Mus musculus FLI-LRR associated protein-1 mRNA, complete cds {IMAGE:348137}	348137 IMAGE Consortium	W30242
2467 G	2 021CA1EF	IMAGE EST {IMAGE:348006}	348006 IMAGE Consortium	W29778
2468 G	4 021CA1EF	ESTs, Highly similar to CHLORINE CHANNEL PROTEIN P64 [Bos taurus] {IMAGE:348115}	348115 IMAGE Consortium	W30612
2469 G	6 021CA1EF	ESTs {IMAGE:335187}	335187 IMAGE Consortium	W17598
2470 G	8 021CA1EF	ESTs {IMAGE:335133}	335133 IMAGE Consortium	W36959
2471 G	10 021CA1EF	ESTs, Moderately similar to WDNM1 PROTEIN [Rattus norvegicus] {IMAGE:331847}	331847 IMAGE Consortium	W08116
2472 G	12 021CA1EF	Replication factor C, 140 kDa {IMAGE:336605}	336605 IMAGE Consortium	W36532
2473 A	2 021JA1EG	ESTs {IMAGE:334383}	334383 IMAGE Consortium	W36545
2474 A	4 021JA1EG	Cathelin-like protein {IMAGE:334755}	334755 IMAGE Consortium	W17771
2475 A	6 021JA1EG	IMAGE EST {IMAGE:348459}	348459 IMAGE Consortium	W35799
2476 A	8 021JA1EG	IMAGE EST {IMAGE:335197}	335197 IMAGE Consortium	W17603
2477 A	10 021JA1EG	IMAGE EST {IMAGE:335864}	335864 IMAGE Consortium	W34148
2478 A	12 021JA1EG	ESTs {IMAGE:336173}	336173 IMAGE Consortium	W17938

2479 C	2 021JA1EG	IMAGE EST {IMAGE:423574}	423574 IMAGE Consortium	W98791
2480 C	4 021JA1EG	IMAGE EST {IMAGE:335413}	335413 IMAGE Consortium	W17593
2481 C	6 021JA1EG	ESTs {IMAGE:335983}	335983 IMAGE Consortium	W18015
2482 C	8 021JA1EG	IMAGE EST {IMAGE:332327}	332327 IMAGE Consortium	W08452
2483 C	10 021JA1EG	IMAGE EST {IMAGE:332532}	332532 IMAGE Consortium	W08555
2484 C	12 021JA1EG	IMAGE EST {IMAGE:332975}	332975 IMAGE Consortium	W08784
2485 E	2 021JA1EG	ESTs {IMAGE:331690}	331690 IMAGE Consortium	W14184
2486 E	4 021JA1EG	ESTs, Highly similar to CREATINE KINASE, SARCOMERIC MITOCHONDRIAL PRECURSOR [Rattus norvegicus] {IMAGE:336085}	336085 IMAGE Consortium	W18057
2487 E	6 021JA1EG	ESTs, Weakly similar to D2045.9 [C.elegans] {IMAGE:336362}	336362 IMAGE Consortium	W36470
2488 E	8 021JA1EG	Phenylalkylamine Ca <sup>2+</sup> antagonist (emopamil) binding protein {IMAGE:332556}	332556 IMAGE Consortium	W08348
2489 E	10 021JA1EG	ESTs, Weakly similar to RIL protein [M.musculus] {IMAGE:336726}	336726 IMAGE Consortium	W36825
2490 E	12 021JA1EG	ESTs, Weakly similar to weakly similar to myosin heavy chain [C.elegans] {IMAGE:337212}	337212 IMAGE Consortium	W36864
2491 G	2 021JA1EG	ESTs, Highly similar to ERPROT 213-21 [H.sapiens] {IMAGE:353957}	353957 IMAGE Consortium	W44112
2492 G	4 021JA1EG	IMAGE EST {IMAGE:354595}	354595 IMAGE Consortium	W46020
2493 G	6 021JA1EG	ESTs {IMAGE:352044}	352044 IMAGE Consortium	W33414
2494 G	8 021JA1EG	IMAGE EST {IMAGE:352723}	352723 IMAGE Consortium	W35880
2495 G	10 021JA1EG	ESTs {IMAGE:353537}	353537 IMAGE Consortium	W41638
2496 G	12 021JA1EG	IMAGE EST {IMAGE:350670}	350670 IMAGE Consortium	W42233
2497 A	2 021QA1EH	IMAGE EST {IMAGE:352488}	352488 IMAGE Consortium	W33741
2498 A	4 021QA1EH	M.musculus trop2 gene {IMAGE:352938}	352938 IMAGE Consortium	W34921
2499 A	6 021QA1EH	Glutamine fructose-6-phosphate transaminase {IMAGE:353215}	353215 IMAGE Consortium	W41682
2500 A	8 021QA1EH	ESTs, Highly similar to STATHMIN-LIKE PROTEIN XB3 [Xenopus laevis] {IMAGE:350652}	350652 IMAGE Consortium	W42241
2501 A	10 021QA1EH	IMAGE EST {IMAGE:351204}	351204 IMAGE Consortium	W41229
2502 A	12 021QA1EH	Mus musculus cell death activator CIDE-A (Cide-a) mRNA, complete cds {IMAGE:351557}	351557 IMAGE Consortium	W41358
2503 C	2 021QA1EH	IMAGE EST {IMAGE:352302}	352302 IMAGE Consortium	W33661
2504 C	4 021QA1EH	IMAGE EST {IMAGE:352574}	352574 IMAGE Consortium	W34404
2505 C	6 021QA1EH	ESTs {IMAGE:353059}	353059 IMAGE Consortium	W34786
2506 C	8 021QA1EH	ESTs {IMAGE:353229}	353229 IMAGE Consortium	W41821
2507 C	10 021QA1EH	M.musculus mRNA for ribosomal protein S8 {IMAGE:354601}	354601 IMAGE Consortium	W46090
2508 C	12 021QA1EH	IMAGE EST {IMAGE:353024}	353024 IMAGE Consortium	W36002
2509 E	2 021QA1EH	ESTs, Moderately similar to predicted using Genefinder [C.elegans] {IMAGE:350400}	350400 IMAGE Consortium	W34995
2510 E	4 021QA1EH	ESTs {IMAGE:353206}	353206 IMAGE Consortium	W41811
2511 E	6 021QA1EH	Early development regulator {IMAGE:350848}	350848 IMAGE Consortium	W40777

2512 E	8 021QA1EH	TYRO3 protein tyrosine kinase 3 {IMAGE:351163}	351163 IMAGE Consortium	W41438
2513 E	10 021QA1EH	ESTs, Highly similar to P80-COILIN [Homo sapiens] {IMAGE:351869}	351869 IMAGE Consortium	W40627
2514 E	12 021QA1EH	EST {IMAGE:352780}	352780 IMAGE Consortium	W34657
2515 G	2 021QA1EH	Mus musculus alpha-synuclein mRNA, complete cds {IMAGE:353366}	353366 IMAGE Consortium	W41663
2516 G	4 021QA1EH	POU domain, class 2, transcription factor 3 {IMAGE:350731}	350731 IMAGE Consortium	W44060
2517 G	6 021QA1EH	ESTs, Weakly similar to TRANSLATION INITIATION FACTOR EIF-2B-GAMMA SUBUNIT [Saccharomyces cerevisiae] {IMAGE:354186}	354186 IMAGE Consortium	W44081
2518 G	8 021QA1EH	Cytochrome P450, 2e1, ethanol inducible {IMAGE:354629}	354629 IMAGE Consortium	W46098
2519 G	10 021QA1EH	ESTs {IMAGE:354377}	354377 IMAGE Consortium	W46106
2520 G	12 021QA1EH	IMAGE EST {IMAGE:352837}	352837 IMAGE Consortium	W34678
2521 A	2 021XA1EI	Mus musculus myosin binding protein H (MyBP-H) mRNA, complete cds {IMAGE:374030}	374030 IMAGE Consortium	W59642
2522 A	4 021XA1EI	IMAGE EST {IMAGE:372679}	372679 IMAGE Consortium	W62668
2523 A	6 021XA1EI	ESTs {IMAGE:367584}	367584 IMAGE Consortium	W54562
2524 A	8 021XA1EI	ESTs {IMAGE:373169}	373169 IMAGE Consortium	W57031
2525 A	10 021XA1EI	Transforming growth factor, beta 2 {IMAGE:367780}	367780 IMAGE Consortium	W53962
2526 A	12 021XA1EI	IMAGE EST {IMAGE:374103}	374103 IMAGE Consortium	W59087
2527 C	2 021XA1EI	IMAGE EST {IMAGE:367436}	367436 IMAGE Consortium	W54284
2528 C	4 021XA1EI	IMAGE EST {IMAGE:373631}	373631 IMAGE Consortium	W64529
2529 C	6 021XA1EI	ESTs, Weakly similar to ORF YGR200c [S.cerevisiae] {IMAGE:374247}	374247 IMAGE Consortium	W59202
2530 C	8 021XA1EI	ESTs, Highly similar to RSP5 PROTEIN [Saccharomyces cerevisiae] {IMAGE:354910}	354910 IMAGE Consortium	W45913
2531 C	10 021XA1EI	Heat shock protein, 84 kDa 1 {IMAGE:356196}	356196 IMAGE Consortium	W50759
2532 C	12 021XA1EI	IMAGE EST {IMAGE:372913}	372913 IMAGE Consortium	W59516
2533 E	2 021XA1EI	IMAGE EST {IMAGE:367748}	367748 IMAGE Consortium	W54729
2534 E	4 021XA1EI	Cyclic nucleotide phosphodiesterase 1 {IMAGE:374248}	374248 IMAGE Consortium	W63987
2535 E	6 021XA1EI	IMAGE EST {IMAGE:374372}	374372 IMAGE Consortium	W65234
2536 E	8 021XA1EI	ESTs {IMAGE:374747}	374747 IMAGE Consortium	W59462
2537 E	10 021XA1EI	ESTs {IMAGE:385142}	385142 IMAGE Consortium	W64729
2538 E	12 021XA1EI	IMAGE EST {IMAGE:388537}	388537 IMAGE Consortium	W70986
2539 G	2 021XA1EI	ESTs, Weakly similar to PROTEIN E(SEV)2B [Drosophila melanogaster] {IMAGE:386996}	386996 IMAGE Consortium	W65915
2540 G	4 021XA1EI	IMAGE EST {IMAGE:388264}	388264 IMAGE Consortium	W66628
2541 G	6 021XA1EI	Cadherin 5 {IMAGE:385761}	385761 IMAGE Consortium	W62248
2542 G	8 021XA1EI	ESTs {IMAGE:385744}	385744 IMAGE Consortium	W64241
2543 G	10 021XA1EI	IMAGE EST {IMAGE:389920}	389920 IMAGE Consortium	W71555
2544 G	12 021XA1EI	IMAGE EST {IMAGE:374349}	374349 IMAGE Consortium	W65230
2545 A	2 0214A1EJ	IMAGE EST {IMAGE:351056}	351056 IMAGE Consortium	W40879
2546 A	4 0214A1EJ	IMAGE EST {IMAGE:354451}	354451 IMAGE Consortium	W45764

2547 A	6 0214A1EJ	IMAGE EST {IMAGE:355223}	355223 IMAGE Consortium	W47923
2548 A	8 0214A1EJ	ESTs {IMAGE:372421}	372421 IMAGE Consortium	W53621
2549 A	10 0214A1EJ	Mus musculus tbc1 mRNA, complete cds {IMAGE:367636}	367636 IMAGE Consortium	W54803
2550 A	12 0214A1EJ	ESTs, Moderately similar to MANNOSE-1-PHOSPHATE GUANYLTRANSFERASE [ <i>Saccharomyces cerevisiae</i> ] {IMAGE:373922}	373922 IMAGE Consortium	W59400
2551 C	2 0214A1EJ	ESTs, Highly similar to transcription factor NF-AT 45K chain [ <i>H.sapiens</i> ] {IMAGE:368164}	368164 IMAGE Consortium	W54089
2552 C	4 0214A1EJ	ESTs {IMAGE:355209}	355209 IMAGE Consortium	W47752
2553 C	6 0214A1EJ	IMAGE EST {IMAGE:355952}	355952 IMAGE Consortium	W50628
2554 C	8 0214A1EJ	IMAGE EST {IMAGE:372940}	372940 IMAGE Consortium	W59578
2555 C	10 0214A1EJ	ESTs, Highly similar to TUBULIN GAMMA CHAIN [ <i>Homo sapiens</i> ] {IMAGE:367382}	367382 IMAGE Consortium	W50677
2556 C	12 0214A1EJ	ESTs {IMAGE:368260}	368260 IMAGE Consortium	W54143
2557 E	2 0214A1EJ	FOLATE RECEPTOR BETA PRECURSOR {IMAGE:374190}	374190 IMAGE Consortium	W59165
2558 E	4 0214A1EJ	IMAGE EST {IMAGE:354965}	354965 IMAGE Consortium	W46001
2559 E	6 0214A1EJ	IMAGE EST {IMAGE:355442}	355442 IMAGE Consortium	W48318
2560 E	8 0214A1EJ	IMAGE EST {IMAGE:355265}	355265 IMAGE Consortium	W47857
2561 E	10 0214A1EJ	IMAGE EST {IMAGE:367138}	367138 IMAGE Consortium	W54048
2562 E	12 0214A1EJ	ESTs, Highly similar to KIAA0156 gene product is related to <i>Xenopus</i> nucleolin. [ <i>H.sapiens</i> ] {IMAGE:367601}	367601 IMAGE Consortium	W54471
2563 G	2 0214A1EJ	IMAGE EST {IMAGE:374267}	374267 IMAGE Consortium	W59215
2564 G	4 0214A1EJ	Chondroitin sulfate proteoglycan 2 {IMAGE:355990}	355990 IMAGE Consortium	W49048
2565 G	6 0214A1EJ	ESTs {IMAGE:373000}	373000 IMAGE Consortium	W58977
2566 G	8 0214A1EJ	ESTs {IMAGE:367890}	367890 IMAGE Consortium	W54486
2567 G	10 0214A1EJ	Cytochrome P450, 2e1, ethanol inducible {IMAGE:368382}	368382 IMAGE Consortium	W55715
2568 G	12 0214A1EJ	<i>M. musculus</i> mRNA for MAP kinase-activated protein kinase 2 {IMAGE:354859}	354859 IMAGE Consortium	W45833
2569 A	2 021BA1EK	ESTs {IMAGE:390171}	390171 IMAGE Consortium	W70705
2570 A	4 021BA1EK	ESTs, Highly similar to suppressin [ <i>R.norvegicus</i> ] {IMAGE:390207}	390207 IMAGE Consortium	W71604
2571 A	6 021BA1EK	IMAGE EST {IMAGE:374688}	374688 IMAGE Consortium	W59438
2572 A	8 021BA1EK	ESTs {IMAGE:387237}	387237 IMAGE Consortium	W65557
2573 A	10 021BA1EK	IMAGE EST {IMAGE:387762}	387762 IMAGE Consortium	W66779
2574 A	12 021BA1EK	ESTs, Highly similar to nuclear transcriptional repressor Mph1 [ <i>M.musculus</i> ] {IMAGE:388233}	388233 IMAGE Consortium	W67089
2575 C	2 021BA1EK	<i>Mus musculus</i> major histocompatibility locus class II region; Fas-binding protein Daxx (DAXX) gene, partial cds; Bing1 (BING1), tapasin (tapasin), RalGDS-like factor (RLF), KE2 (KE2), BING4 (BING4), beta1, 3-galactosyl transferase {IMAGE:387000}	387000 IMAGE Consortium	W65519
2576 C	4 021BA1EK	ESTs, Weakly similar to CALCIUM/CALMODULIN-DEPENDENT PROTEIN KINASE TYPE IV CATALYTIC CHAIN [ <i>M.musculus</i> ] {IMAGE:374727}	374727 IMAGE Consortium	W59265

2577 C	6 021BA1EK	IMAGE EST {IMAGE:387722}	387722 IMAGE Consortium	W70495
2578 C	8 021BA1EK	IMAGE EST {IMAGE:388050}	388050 IMAGE Consortium	W66889
2579 C	10 021BA1EK	Mus musculus mRNA for leptin receptor gene-related protein {IMAGE:385417}	385417 IMAGE Consortium	W62925
2580 C	12 021BA1EK	IMAGE EST {IMAGE:388798}	388798 IMAGE Consortium	W71286
2581 E	2 021BA1EK	ESTs {IMAGE:388410}	388410 IMAGE Consortium	W66710
2582 E	4 021BA1EK	IMAGE EST {IMAGE:385429}	385429 IMAGE Consortium	W62931
2583 E	6 021BA1EK	ESTs {IMAGE:388919}	388919 IMAGE Consortium	W67027
2584 E	8 021BA1EK	IMAGE EST {IMAGE:389827}	389827 IMAGE Consortium	W71523
2585 E	10 021BA1EK	ESTs, Highly similar to T-KININOGEN II PRECURSOR [Rattus norvegicus] {IMAGE:390390}	390390 IMAGE Consortium	W70823
2586 E	12 021BA1EK	IMAGE EST {IMAGE:387002}	387002 IMAGE Consortium	W65520
2587 G	2 021BA1EK	IMAGE EST {IMAGE:386820}	386820 IMAGE Consortium	W64463
2588 G	4 021BA1EK	IMAGE EST {IMAGE:374631}	374631 IMAGE Consortium	W59393
2589 G	6 021BA1EK	ESTs {IMAGE:387179}	387179 IMAGE Consortium	W65930
2590 G	8 021BA1EK	IMAGE EST {IMAGE:387922}	387922 IMAGE Consortium	W66800
2591 G	10 021BA1EK	IMAGE EST {IMAGE:388308}	388308 IMAGE Consortium	W70876
2592 G	12 021BA1EK	ESTs {IMAGE:385467}	385467 IMAGE Consortium	W61416
2593 A	2 021IA1EL	ESTs, Highly similar to INTERFERON-RELATED PROTEIN PC4 [Mus musculus] {IMAGE:387482}	387482 IMAGE Consortium	W65790
2594 A	4 021IA1EL	IMAGE EST {IMAGE:387946}	387946 IMAGE Consortium	W66752
2595 A	6 021IA1EL	Mouse mRNA for KIFC1, complete cds {IMAGE:388296}	388296 IMAGE Consortium	W70871
2596 A	8 021IA1EL	IMAGE EST {IMAGE:385314}	385314 IMAGE Consortium	W63814
2597 A	10 021IA1EL	IMAGE EST {IMAGE:388619}	388619 IMAGE Consortium	W66727
2598 A	12 021IA1EL	ESTs {IMAGE:385435}	385435 IMAGE Consortium	W64144
2599 C	2 021IA1EL	ESTs {IMAGE:385387}	385387 IMAGE Consortium	W74897
2600 C	4 021IA1EL	IMAGE EST {IMAGE:388691}	388691 IMAGE Consortium	W71048
2601 C	6 021IA1EL	IMAGE EST {IMAGE:385988}	385988 IMAGE Consortium	W64277
2602 C	8 021IA1EL	ESTs, Highly similar to SEC14L [H.sapiens] {IMAGE:390192}	390192 IMAGE Consortium	W70708
2603 C	10 021IA1EL	Cytochrome P450, 2e1, ethanol inducible {IMAGE:390178}	390178 IMAGE Consortium	W71365
2604 C	12 021IA1EL	Distal-less homeobox 5 {IMAGE:387148}	387148 IMAGE Consortium	W65729
2605 E	2 021IA1EL	IMAGE EST {IMAGE:375009}	375009 IMAGE Consortium	W62018
2606 E	4 021IA1EL	IMAGE EST {IMAGE:387925}	387925 IMAGE Consortium	W66749
2607 E	6 021IA1EL	IMAGE EST {IMAGE:388310}	388310 IMAGE Consortium	W70877
2608 E	8 021IA1EL	ESTs {IMAGE:388644}	388644 IMAGE Consortium	W66730
2609 E	10 021IA1EL	ESTs, Moderately similar to THYROID RECEPTOR INTERACTING PROTEIN 7 [H.sapiens] {IMAGE:385581}	385581 IMAGE Consortium	W64196
2610 E	12 021IA1EL	ESTs, Highly similar to FRUCTOSE-BISPHOSPHATE ALDOLASE C [Homo sapiens] {IMAGE:390313}	390313 IMAGE Consortium	W70792
2611 G	2 021IA1EL	ESTs, Highly similar to TRANSLOCON-ASSOCIATED PROTEIN, GAMMA SUBUNIT [Rattus norvegicus] {IMAGE:404526}	404526 IMAGE Consortium	W83038
2612 G	4 021IA1EL	M.musculus mRNA for splicing factor U2AF (65 kD)	405962 IMAGE Consortium	W83291

		{IMAGE:405962}		
2613 G	6 021IA1EL	IMAGE EST {IMAGE:402504}	402504 IMAGE Consortium	W80096
2614 G	8 021IA1EL	Cellular retinoic acid binding protein II {IMAGE:403035}	403035 IMAGE Consortium	W81912
2615 G	10 021IA1EL	Matrix metalloproteinase 2 {IMAGE:402738}	402738 IMAGE Consortium	W80177
2616 G	12 021IA1EL	Nucleosome assembly protein 1 like 4 {IMAGE:403311}	403311 IMAGE Consortium	W81759
2617 A	2 021PA1EM	40S RIBOSOMAL PROTEIN SA {IMAGE:400741}	400741 IMAGE Consortium	W75686
2618 A	4 021PA1EM	IMAGE EST {IMAGE:400841}	400841 IMAGE Consortium	W76784
2619 A	6 021PA1EM	ESTs {IMAGE:401221}	401221 IMAGE Consortium	W97933
2620 A	8 021PA1EM	Nucleosome assembly protein 1 like 4 {IMAGE:401788}	401788 IMAGE Consortium	W78317
2621 A	10 021PA1EM	Benzodiazepine receptor, peripheral {IMAGE:404615}	404615 IMAGE Consortium	W82946
2622 A	12 021PA1EM	IMAGE EST {IMAGE:402544}	402544 IMAGE Consortium	W80263
2623 C	2 021PA1EM	ESTs, Moderately similar to NECDIN [Mus musculus] {IMAGE:401980}	401980 IMAGE Consortium	W77507
2624 C	4 021PA1EM	ESTs, Weakly similar to ZK792.1 [C.elegans] {IMAGE:403104}	403104 IMAGE Consortium	W82005
2625 C	6 021PA1EM	ESTs, Highly similar to NADH-UBIQUINONE OXIDOREDUCTASE B8 SUBUNIT [Bos taurus] {IMAGE:400312}	400312 IMAGE Consortium	W77476
2626 C	8 021PA1EM	IMAGE EST {IMAGE:403695}	403695 IMAGE Consortium	W82286
2627 C	10 021PA1EM	ESTs {IMAGE:404044}	404044 IMAGE Consortium	W82416
2628 C	12 021PA1EM	Thymoma viral proto-oncogene 2 {IMAGE:404199}	404199 IMAGE Consortium	W82557
2629 E	2 021PA1EM	IMAGE EST {IMAGE:403624}	403624 IMAGE Consortium	W82122
2630 E	4 021PA1EM	ESTs {IMAGE:404140}	404140 IMAGE Consortium	W82514
2631 E	6 021PA1EM	Even skipped homeotic gene 2 homolog {IMAGE:401372}	401372 IMAGE Consortium	AA000473
2632 E	8 021PA1EM	Crystallin, gamma E {IMAGE:401766}	401766 IMAGE Consortium	W78478
2633 E	10 021PA1EM	ESTs, Highly similar to PROTEIN PHOSPHATASE INHIBITOR 1 [Rattus norvegicus] {IMAGE:402053}	402053 IMAGE Consortium	W75893
2634 E	12 021PA1EM	ESTs {IMAGE:402843}	402843 IMAGE Consortium	W80245
2635 G	2 021PA1EM	Procollagen, type XV {IMAGE:406344}	406344 IMAGE Consortium	W83331
2636 G	4 021PA1EM	ESTs, Moderately similar to KIAA0330 [H.sapiens] {IMAGE:406188}	406188 IMAGE Consortium	W83838
2637 G	6 021PA1EM	IMAGE EST {IMAGE:406728}	406728 IMAGE Consortium	W83737
2638 G	8 021PA1EM	IMAGE EST {IMAGE:419419}	419419 IMAGE Consortium	W88186
2639 G	10 021PA1EM	ESTs {IMAGE:408115}	408115 IMAGE Consortium	W85333
2640 G	12 021PA1EM	ESTs {IMAGE:408634}	408634 IMAGE Consortium	W85646
2641 A	2 021WA1EN	IMAGE EST {IMAGE:406106}	406106 IMAGE Consortium	W84047
2642 A	4 021WA1EN	IMAGE EST {IMAGE:402700}	402700 IMAGE Consortium	W80060
2643 A	6 021WA1EN	ESTs {IMAGE:403252}	403252 IMAGE Consortium	W81803
2644 A	8 021WA1EN	ESTs {IMAGE:400370}	400370 IMAGE Consortium	W77133
2645 A	10 021WA1EN	Kinesin heavy chain member 4 {IMAGE:400198}	400198 IMAGE Consortium	W77625
2646 A	12 021WA1EN	ESTs {IMAGE:403607}	403607 IMAGE Consortium	W82121
2647 C	2 021WA1EN	IMAGE EST {IMAGE:403313}	403313 IMAGE Consortium	W81838
2648 C	4 021WA1EN	Erythroid Kruppel-like factor {IMAGE:400416}	400416 IMAGE Consortium	W77245

2649 C	6 021WA1EN	ESTs, Highly similar to DNAJ PROTEIN HOMOLOG 1 [Homo sapiens] {IMAGE:400617}	400617 IMAGE Consortium	W75670
2650 C	8 021WA1EN	IMAGE EST {IMAGE:404002}	404002 IMAGE Consortium	W82777
2651 C	10 021WA1EN	IMAGE EST {IMAGE:401856}	401856 IMAGE Consortium	W82389
2652 C	12 021WA1EN	ESTs, Weakly similar to HYPOTHETICAL 20.1 KD PROTEIN IN HSCA 5'REGION [Escherichia coli] {IMAGE:404564}	404564 IMAGE Consortium	W82888
2653 E	2 021WA1EN	ESTs {IMAGE:401809}	401809 IMAGE Consortium	W78456
2654 E	4 021WA1EN	ESTs, Moderately similar to similar to ATPases associated with various cellular activities [C.elegans] {IMAGE:401821}	401821 IMAGE Consortium	W78557
2655 E	6 021WA1EN	ESTs {IMAGE:401976}	401976 IMAGE Consortium	W77505
2656 E	8 021WA1EN	ESTs {IMAGE:402894}	402894 IMAGE Consortium	W80156
2657 E	10 021WA1EN	ESTs {IMAGE:403185}	403185 IMAGE Consortium	W81735
2658 E	12 021WA1EN	ESTs, Moderately similar to SIGNAL RECOGNITION PARTICLE 19 KD PROTEIN [Homo sapiens] {IMAGE:400681}	400681 IMAGE Consortium	W75853
2659 G	2 021WA1EN	ESTs, Moderately similar to CASEIN KINASE I, GAMMA ISOFORM [Bos taurus] {IMAGE:402551}	402551 IMAGE Consortium	W80270
2660 G	4 021WA1EN	ESTs, Highly similar to HYPOTHETICAL 39.7 KD PROTEIN C34E10.2 IN CHROMOSOME III [Caenorhabditis elegans] {IMAGE:403005}	403005 IMAGE Consortium	W81857
2661 G	6 021WA1EN	IMAGE EST {IMAGE:400113}	400113 IMAGE Consortium	W77375
2662 G	8 021WA1EN	IMAGE EST {IMAGE:400525}	400525 IMAGE Consortium	W77508
2663 G	10 021WA1EN	ESTs {IMAGE:400864}	400864 IMAGE Consortium	W76785
2664 G	12 021WA1EN	IMAGE EST {IMAGE:401764}	401764 IMAGE Consortium	W78477
2665 A	2 0213A1EO	IMAGE EST {IMAGE:419265}	419265 IMAGE Consortium	AA052196
2666 A	4 0213A1EO	IMAGE EST {IMAGE:419420}	419420 IMAGE Consortium	W88183
2667 A	6 0213A1EO	IMAGE EST {IMAGE:407691}	407691 IMAGE Consortium	W84939
2668 A	8 0213A1EO	Dlk1-like homolog (Drosophila) {IMAGE:419848}	419848 IMAGE Consortium	W91219
2669 A	10 0213A1EO	Erythrocyte protein band 7.2 {IMAGE:419756}	419756 IMAGE Consortium	W89549
2670 A	12 0213A1EO	ESTs {IMAGE:420029}	420029 IMAGE Consortium	W91010
2671 C	2 0213A1EO	IMAGE EST {IMAGE:419553}	419553 IMAGE Consortium	W88173
2672 C	4 0213A1EO	IMAGE EST {IMAGE:419516}	419516 IMAGE Consortium	AA052074
2673 C	6 0213A1EO	IMAGE EST {IMAGE:419788}	419788 IMAGE Consortium	W89385
2674 C	8 0213A1EO	IMAGE EST {IMAGE:419896}	419896 IMAGE Consortium	W89614
2675 C	10 0213A1EO	IMAGE EST {IMAGE:419900}	419900 IMAGE Consortium	W89615
2676 C	12 0213A1EO	IMAGE EST {IMAGE:408996}	408996 IMAGE Consortium	W87195
2677 E	2 0213A1EO	Interferon dependent positive acting transcription factor 3 gamma {IMAGE:407116}	407116 IMAGE Consortium	W84302
2678 E	4 0213A1EO	ESTs {IMAGE:407274}	407274 IMAGE Consortium	W85166
2679 E	6 0213A1EO	ESTs {IMAGE:407997}	407997 IMAGE Consortium	W85441
2680 E	8 0213A1EO	IMAGE EST {IMAGE:408381}	408381 IMAGE Consortium	W85374
2681 E	10 0213A1EO	U1 AND U2 SMALL NUCLEAR RIBONUCLEOPROTEIN E {IMAGE:408693}	408693 IMAGE Consortium	W85296



2682 E	12 0213A1EO	IMAGE EST {IMAGE:409068}	409068 IMAGE Consortium	W87147
2683 G	2 0213A1EO	ESTs {IMAGE:419387}	419387 IMAGE Consortium	W88044
2684 G	4 0213A1EO	ESTs {IMAGE:419301}	419301 IMAGE Consortium	W89770
2685 G	6 0213A1EO	ESTs, Moderately similar to TFE3 transcription factor [H.sapiens] {IMAGE:419821}	419821 IMAGE Consortium	W91192
2686 G	8 0213A1EO	Mus musculus mRNA for translation initiation factor eIF2 gamma X {IMAGE:419950}	419950 IMAGE Consortium	W89599
2687 G	10 0213A1EO	EST {IMAGE:420204}	420204 IMAGE Consortium	W89845
2688 G	12 0213A1EO	Tuberous sclerosis 2 {IMAGE:420304}	420304 IMAGE Consortium	W89921
2689 A	2 021AA1EP	3'-phosphoadenosine 5'-phosphosulfate synthase 1 {IMAGE:419421}	419421 IMAGE Consortium	W88187
2690 A	4 021AA1EP	ESTs {IMAGE:407816}	407816 IMAGE Consortium	W85641
2691 A	6 021AA1EP	IMAGE EST {IMAGE:419705}	419705 IMAGE Consortium	W89538
2692 A	8 021AA1EP	ESTs, Weakly similar to SULFITE OXIDASE PRECURSOR [Rattus norvegicus] {IMAGE:419711}	419711 IMAGE Consortium	W89541
2693 A	10 021AA1EP	Mouse chromosome 6 BAC-284H12 (Research Genetics mouse BAC library) complete sequence {IMAGE:420054}	420054 IMAGE Consortium	W91014
2694 A	12 021AA1EP	IMAGE EST {IMAGE:420085}	420085 IMAGE Consortium	W89796
2695 C	2 021AA1EP	Nidogen {IMAGE:419437}	419437 IMAGE Consortium	W88036
2696 C	4 021AA1EP	ESTs {IMAGE:419493}	419493 IMAGE Consortium	AA052162
2697 C	6 021AA1EP	IMAGE EST {IMAGE:419774}	419774 IMAGE Consortium	W91176
2698 C	8 021AA1EP	IMAGE EST {IMAGE:408673}	408673 IMAGE Consortium	W85238
2699 C	10 021AA1EP	ESTs, Weakly similar to coded for by C. elegans cDNA yk30b3.5 [C.elegans] {IMAGE:409057}	409057 IMAGE Consortium	W87071
2700 C	12 021AA1EP	IMAGE EST {IMAGE:418706}	418706 IMAGE Consortium	W99246
2701 E	2 021AA1EP	ESTs {IMAGE:419398}	419398 IMAGE Consortium	W88079
2702 E	4 021AA1EP	ESTs {IMAGE:407702}	407702 IMAGE Consortium	W85245
2703 E	6 021AA1EP	IMAGE EST {IMAGE:408110}	408110 IMAGE Consortium	W85338
2704 E	8 021AA1EP	ESTs {IMAGE:408986}	408986 IMAGE Consortium	W87030
2705 E	10 021AA1EP	IMAGE EST {IMAGE:408879}	408879 IMAGE Consortium	W85621
2706 E	12 021AA1EP	IMAGE EST {IMAGE:420602}	420602 IMAGE Consortium	W89427
2707 G	2 021AA1EP	IMAGE EST {IMAGE:407492}	407492 IMAGE Consortium	W89976
2708 G	4 021AA1EP	ESTs {IMAGE:419846}	419846 IMAGE Consortium	W91197
2709 G	6 021AA1EP	ESTs {IMAGE:408626}	408626 IMAGE Consortium	W85644
2710 G	8 021AA1EP	ESTs, Moderately similar to 26 S protease subunit 5b {IMAGE:418563}	418563 IMAGE Consortium	W99037
2711 G	10 021AA1EP	IMAGE EST {IMAGE:420411}	420411 IMAGE Consortium	AA052723
2712 G	12 021AA1EP	Transcription factor 12 {IMAGE:405907}	405907 IMAGE Consortium	W83524
2713 A	2 021HA1EQ	ESTs, Highly similar to MVP1 PROTEIN [Saccharomyces cerevisiae] {IMAGE:423727}	423727 IMAGE Consortium	W98550
2714 A	4 021HA1EQ	ESTs {IMAGE:422123}	422123 IMAGE Consortium	W97364
2715 A	6 021HA1EQ	ESTs, Weakly similar to ORF YOL080c [S.cerevisiae] {IMAGE:422205}	422205 IMAGE Consortium	W97502

2716 A	8 021HA1EQ	IMAGE EST {IMAGE:422743}	422743 IMAGE Consortium	W97858
2717 A	10 021HA1EQ	ESTs {IMAGE:422797}	422797 IMAGE Consortium	W96990
2718 A	12 021HA1EQ	ESTs {IMAGE:423366}	423366 IMAGE Consortium	W98846
2719 C	2 021HA1EQ	IMAGE EST {IMAGE:423107}	423107 IMAGE Consortium	W97863
2720 C	4 021HA1EQ	Interleukin 11 {IMAGE:423087}	423087 IMAGE Consortium	W98051
2721 C	6 021HA1EQ	IMAGE EST {IMAGE:423140}	423140 IMAGE Consortium	W97265
2722 C	8 021HA1EQ	IMAGE EST {IMAGE:422094}	422094 IMAGE Consortium	W97346
2723 C	10 021HA1EQ	RD PROTEIN {IMAGE:421962}	421962 IMAGE Consortium	W97403
2724 C	12 021HA1EQ	ESTs, Highly similar to symplekin [H.sapiens] {IMAGE:422363}	422363 IMAGE Consortium	W97544
2725 E	2 021HA1EQ	ESTs {IMAGE:422220}	422220 IMAGE Consortium	W97484
2726 E	4 021HA1EQ	EST {IMAGE:421017}	421017 IMAGE Consortium	W90956
2727 E	6 021HA1EQ	IMAGE EST {IMAGE:422442}	422442 IMAGE Consortium	W97646
2728 E	8 021HA1EQ	IMAGE EST {IMAGE:423002}	423002 IMAGE Consortium	W97964
2729 E	10 021HA1EQ	ESTs {IMAGE:422983}	422983 IMAGE Consortium	W97959
2730 E	12 021HA1EQ	IMAGE EST {IMAGE:422823}	422823 IMAGE Consortium	W97181
2731 G	2 021HA1EQ	ESTs {IMAGE:422413}	422413 IMAGE Consortium	W97658
2732 G	4 021HA1EQ	ESTs {IMAGE:421385}	421385 IMAGE Consortium	W97046
2733 G	6 021HA1EQ	IMAGE EST {IMAGE:421776}	421776 IMAGE Consortium	W98948
2734 G	8 021HA1EQ	Mus musculus methyl-CpG binding protein MBD2 (Mbd2) mRNA, complete cds {IMAGE:421592}	421592 IMAGE Consortium	W97165
2735 G	10 021HA1EQ	EST {IMAGE:421524}	421524 IMAGE Consortium	W97155
2736 G	12 021HA1EQ	Dlk1-like homolog (Drosophila) {IMAGE:421995}	421995 IMAGE Consortium	W96838
2737 A	2 021OA1ER	IMAGE EST {IMAGE:409001}	409001 IMAGE Consortium	W87227
2738 A	4 021OA1ER	IMAGE EST {IMAGE:420020}	420020 IMAGE Consortium	W90879
2739 A	6 021OA1ER	ESTs {IMAGE:420562}	420562 IMAGE Consortium	AA052144
2740 A	8 021OA1ER	Mouse mRNA for alpha-adaptin (C) {IMAGE:406330}	406330 IMAGE Consortium	W83182
2741 A	10 021OA1ER	IMAGE EST {IMAGE:406222}	406222 IMAGE Consortium	W84062
2742 A	12 021OA1ER	Cryptochrome 1 (photolyase-like) {IMAGE:406761}	406761 IMAGE Consortium	W83558
2743 C	2 021OA1ER	IMAGE EST {IMAGE:420178}	420178 IMAGE Consortium	W89832
2744 C	4 021OA1ER	HOMEBOX PROTEIN HOX-B8 {IMAGE:418660}	418660 IMAGE Consortium	W99235
2745 C	6 021OA1ER	ESTs {IMAGE:418690}	418690 IMAGE Consortium	W96925
2746 C	8 021OA1ER	IMAGE EST {IMAGE:418551}	418551 IMAGE Consortium	W99122
2747 C	10 021OA1ER	NAD-DEPENDENT METHYLENETETRAHYDROFOLATE DEHYDROGENASE {IMAGE:406031}	406031 IMAGE Consortium	W84014
2748 C	12 021OA1ER	ESTs {IMAGE:419185}	419185 IMAGE Consortium	W87955
2749 E	2 021OA1ER	IMAGE EST {IMAGE:420468}	420468 IMAGE Consortium	W90847
2750 E	4 021OA1ER	Putative phosphatase {IMAGE:421061}	421061 IMAGE Consortium	W91228
2751 E	6 021OA1ER	Solute carrier family 4 (anion exchanger), member 1 {IMAGE:422440}	422440 IMAGE Consortium	W97645
2752 E	8 021OA1ER	ESTs {IMAGE:421216}	421216 IMAGE Consortium	W98988
2753 E	10 021OA1ER	IMAGE EST {IMAGE:422854}	422854 IMAGE Consortium	W97947
2754 E	12 021OA1ER	ESTs {IMAGE:423126}	423126 IMAGE Consortium	W97867

2755 G	2 021OA1ER	IMAGE EST {IMAGE:421717}	421717 IMAGE Consortium	W98949
2756 G	4 021OA1ER	ESTs, Weakly similar to PROBABLE PROTEIN DISULFIDE ISOMERASE ER-60 PRECURSOR [M.musculus] {IMAGE:421530}	421530 IMAGE Consortium	W97172
2757 G	6 021OA1ER	Mus musculus (clone 41510) BAD protein mRNA, complete cds {IMAGE:423721}	423721 IMAGE Consortium	W98547
2758 G	8 021OA1ER	IMAGE EST {IMAGE:420709}	420709 IMAGE Consortium	W89713
2759 G	10 021OA1ER	IMAGE EST {IMAGE:420475}	420475 IMAGE Consortium	W89664
2760 G	12 021OA1ER	IMAGE EST {IMAGE:422306}	422306 IMAGE Consortium	W97679
2761 A	2 021VA1ES	ESTs {IMAGE:421716}	421716 IMAGE Consortium	W98964
2762 A	4 021VA1ES	ESTs {IMAGE:421719}	421719 IMAGE Consortium	W98950
2763 A	6 021VA1ES	ESTs {IMAGE:421629}	421629 IMAGE Consortium	W98927
2764 A	8 021VA1ES	IMAGE EST {IMAGE:422163}	422163 IMAGE Consortium	W97385
2765 A	10 021VA1ES	Calpactin I heavy chain {IMAGE:420429}	420429 IMAGE Consortium	W89518
2766 A	12 021VA1ES	ESTs, Highly similar to RNA helicase HEL117 [R.norvegicus] {IMAGE:421991}	421991 IMAGE Consortium	W97204
2767 C	2 021VA1ES	ESTs, Highly similar to ALPHA-ACTININ, SMOOTH MUSCLE ISOFORM [Gallus gallus] {IMAGE:423596}	423596 IMAGE Consortium	W98655
2768 C	4 021VA1ES	IMAGE EST {IMAGE:421060}	421060 IMAGE Consortium	AA052065
2769 C	6 021VA1ES	Protein kinase C, delta {IMAGE:421002}	421002 IMAGE Consortium	W91539
2770 C	8 021VA1ES	IMAGE EST {IMAGE:422325}	422325 IMAGE Consortium	W97588
2771 C	10 021VA1ES	ESTs {IMAGE:421287}	421287 IMAGE Consortium	W90924
2772 C	12 021VA1ES	IMAGE EST {IMAGE:421672}	421672 IMAGE Consortium	W98666
2773 E	2 021VA1ES	M.musculus mRNA homologous to S. cerevisiae RAD54 {IMAGE:425419}	425419 IMAGE Consortium	AA000163
2774 E	4 021VA1ES	IMAGE EST {IMAGE:423884}	423884 IMAGE Consortium	AA052753
2775 E	6 021VA1ES	ESTs {IMAGE:425777}	425777 IMAGE Consortium	AA000799
2776 E	8 021VA1ES	IMAGE EST {IMAGE:425830}	425830 IMAGE Consortium	AA000811
2777 E	10 021VA1ES	IMAGE EST {IMAGE:426378}	426378 IMAGE Consortium	AA002886
2778 E	12 021VA1ES	IMAGE EST {IMAGE:425005}	425005 IMAGE Consortium	W99994
2779 G	2 021VA1ES	ESTs {IMAGE:426445}	426445 IMAGE Consortium	AA002301
2780 G	4 021VA1ES	IMAGE EST {IMAGE:424994}	424994 IMAGE Consortium	W99918
2781 G	6 021VA1ES	IMAGE EST {IMAGE:425053}	425053 IMAGE Consortium	AA000009
2782 G	8 021VA1ES	ESTs {IMAGE:426741}	426741 IMAGE Consortium	AA002733
2783 G	10 021VA1ES	IMAGE EST {IMAGE:425478}	425478 IMAGE Consortium	AA000209
2784 G	12 021VA1ES	EST {IMAGE:425553}	425553 IMAGE Consortium	AA000119
2785 A	2 021A1ET	ESTs {IMAGE:426493}	426493 IMAGE Consortium	AA002322
2786 A	4 021A1ET	Mus musculus mRNA for heterogeneous nuclear ribonucleoprotein H {IMAGE:426163}	426163 IMAGE Consortium	AA003085
2787 A	6 021A1ET	ESTs, Highly similar to KINESIN-II 85 KD SUBUNIT [Strongylocentrotus purpuratus] {IMAGE:426837}	426837 IMAGE Consortium	AA003205
2788 A	8 021A1ET	IMAGE EST {IMAGE:427115}	427115 IMAGE Consortium	AA003384
2789 A	10 021A1ET	IMAGE EST {IMAGE:425677}	425677 IMAGE Consortium	AA000310

2790 A	12 0212A1ET	IMAGE EST {IMAGE:425635}	425635 IMAGE Consortium	AA000249
2791 C	2 0212A1ET	EST {IMAGE:426646}	426646 IMAGE Consortium	AA002743
2792 C	4 0212A1ET	ESTs, Weakly similar to ACTIN POLYMERIZATION INHIBITOR [Gallus gallus] {IMAGE:426965}	426965 IMAGE Consortium	AA003272
2793 C	6 0212A1ET	ESTs, Highly similar to DIAMINE ACETYLTRANSFERASE [Mus musculus] {IMAGE:425721}	425721 IMAGE Consortium	AA000322
2794 C	8 0212A1ET	ESTs {IMAGE:425415}	425415 IMAGE Consortium	AA000136
2795 C	10 0212A1ET	ESTs {IMAGE:425636}	425636 IMAGE Consortium	AA000246
2796 C	12 0212A1ET	ESTs, Highly similar to GLYCOGEN SYNTHASE KINASE-3 ALPHA [Rattus norvegicus] {IMAGE:423863}	423863 IMAGE Consortium	W91360
2797 E	2 0212A1ET	IMAGE EST {IMAGE:427259}	427259 IMAGE Consortium	AA003047
2798 E	4 0212A1ET	IMAGE EST {IMAGE:427028}	427028 IMAGE Consortium	AA003283
2799 E	6 0212A1ET	Mus musculus mRNA for synaptotagumin III, complete cds {IMAGE:427562}	427562 IMAGE Consortium	AA002524
2800 E	8 0212A1ET	IMAGE EST {IMAGE:425857}	425857 IMAGE Consortium	AA000775
2801 E	10 0212A1ET	ESTs {IMAGE:424341}	424341 IMAGE Consortium	W98480
2802 E	12 0212A1ET	Lymphocyte antigen 6 complex, locus C {IMAGE:425855}	425855 IMAGE Consortium	AA000712
2803 G	2 0212A1ET	IMAGE EST {IMAGE:423915}	423915 IMAGE Consortium	W91463
2804 G	4 0212A1ET	Anti-oxidant protein 1 {IMAGE:423832}	423832 IMAGE Consortium	W91307
2805 G	6 0212A1ET	Cytochrome P450, 2e1, ethanol inducible {IMAGE:424393}	424393 IMAGE Consortium	W98316
2806 G	8 0212A1ET	IMAGE EST {IMAGE:424329}	424329 IMAGE Consortium	W98402
2807 G	10 0212A1ET	IMAGE EST {IMAGE:425804}	425804 IMAGE Consortium	AA000804
2808 G	12 0212A1ET	ESTs {IMAGE:426178}	426178 IMAGE Consortium	AA003062
2809 A	2 0219A1EU	ESTs, Moderately similar to HYPOTHETICAL 92.1 KD PROTEIN C24H6.03 IN CHROMOSOME I [Schizosaccharomyces pombe] {IMAGE:437413}	437413 IMAGE Consortium	AA003633
2810 A	4 0219A1EU	ESTs, Highly similar to LYSYL-TRNA SYNTHETASE [Cricetulus longicaudatus] {IMAGE:439548}	439548 IMAGE Consortium	AA004176
2811 A	6 0219A1EU	Aminolevulinic acid synthase 2, erythroid {IMAGE:439199}	439199 IMAGE Consortium	AA003951
2812 A	8 0219A1EU	IMAGE EST {IMAGE:439783}	439783 IMAGE Consortium	AA008478
2813 A	10 0219A1EU	IMAGE EST {IMAGE:439811}	439811 IMAGE Consortium	AA008593
2814 A	12 0219A1EU	ESTs, Highly similar to ENDOTHELIAL ACTIN-BINDING PROTEIN [Homo sapiens] {IMAGE:427360}	427360 IMAGE Consortium	AA003323
2815 C	2 0219A1EU	ESTs {IMAGE:439411}	439411 IMAGE Consortium	AA004111
2816 C	4 0219A1EU	IMAGE EST {IMAGE:439439}	439439 IMAGE Consortium	AA004146
2817 C	6 0219A1EU	IMAGE EST {IMAGE:439732}	439732 IMAGE Consortium	AA014379
2818 C	8 0219A1EU	IMAGE EST {IMAGE:438578}	438578 IMAGE Consortium	AA050963
2819 C	10 0219A1EU	ESTs {IMAGE:427480}	427480 IMAGE Consortium	AA002452
2820 C	12 0219A1EU	Ephrin B1 {IMAGE:427319}	427319 IMAGE Consortium	AA003297
2821 E	2 0219A1EU	IMAGE EST {IMAGE:439535}	439535 IMAGE Consortium	AA008543
2822 E	4 0219A1EU	Myeloid/lymphoid or mixed-lineage leukemia {IMAGE:438257}	438257 IMAGE Consortium	AA008412
2823 E	6 0219A1EU	IMAGE EST {IMAGE:439620}	439620 IMAGE Consortium	AA008686

2824 E	8 0219A1EU	IMAGE EST {IMAGE:438592}	438592 IMAGE Consortium	AA050931
2825 E	10 0219A1EU	ESTs, Weakly similar to F01F1.11 [C.elegans] {IMAGE:438550}	438550 IMAGE Consortium	AA051051
2826 E	12 0219A1EU	IMAGE EST {IMAGE:438914}	438914 IMAGE Consortium	AA007763
2827 G	2 0219A1EU	IMAGE EST {IMAGE:438483}	438483 IMAGE Consortium	AA008256
2828 G	4 0219A1EU	ESTs, Highly similar to HYPOTHETICAL 94.2 KD PROTEIN C38D4.5 IN CHROMOSOME III [Caenorhabditis elegans] {IMAGE:427601}	427601 IMAGE Consortium	AA002692
2829 G	6 0219A1EU	ESTs {IMAGE:437040}	437040 IMAGE Consortium	AA002821
2830 G	8 0219A1EU	IMAGE EST {IMAGE:438938}	438938 IMAGE Consortium	AA007808
2831 G	10 0219A1EU	IMAGE EST {IMAGE:437336}	437336 IMAGE Consortium	AA003698
2832 G	12 0219A1EU	IMAGE EST {IMAGE:439402}	439402 IMAGE Consortium	AA004061
2833 A	2 021GA1EV	EST {IMAGE:426025}	426025 IMAGE Consortium	AA000768
2834 A	4 021GA1EV	IMAGE EST {IMAGE:424462}	424462 IMAGE Consortium	W98444
2835 A	6 021GA1EV	IMAGE EST {IMAGE:424618}	424618 IMAGE Consortium	AA000563
2836 A	8 021GA1EV	ESTs {IMAGE:425025}	425025 IMAGE Consortium	AA000002
2837 A	10 021GA1EV	Annexin V {IMAGE:426546}	426546 IMAGE Consortium	AA002439
2838 A	12 021GA1EV	ESTs {IMAGE:427062}	427062 IMAGE Consortium	AA003247
2839 C	2 021GA1EV	IMAGE EST {IMAGE:425049}	425049 IMAGE Consortium	AA000008
2840 C	4 021GA1EV	ESTs {IMAGE:425279}	425279 IMAGE Consortium	W99891
2841 C	6 021GA1EV	ESTs, Highly similar to HYPOTHETICAL TRP-ASP REPEATS CONTAINING PROTEIN IN SMY2-RPS101 INTERGENIC REGION [Saccharomyces cerevisiae] {IMAGE:425452}	425452 IMAGE Consortium	AA000181
2842 C	8 021GA1EV	EST {IMAGE:426934}	426934 IMAGE Consortium	AA003308
2843 C	10 021GA1EV	ESTs {IMAGE:425834}	425834 IMAGE Consortium	AA000466
2844 C	12 021GA1EV	ESTs {IMAGE:424559}	424559 IMAGE Consortium	W98278
2845 E	2 021GA1EV	ESTs, Highly similar to pescadillo [H.sapiens] {IMAGE:427222}	427222 IMAGE Consortium	AA003101
2846 E	4 021GA1EV	ESTs {IMAGE:427469}	427469 IMAGE Consortium	AA002332
2847 E	6 021GA1EV	ESTs {IMAGE:426033}	426033 IMAGE Consortium	AA002836
2848 E	8 021GA1EV	IMAGE EST {IMAGE:425800}	425800 IMAGE Consortium	AA000802
2849 E	10 021GA1EV	Protein kinase C substrate 80K-H {IMAGE:424686}	424686 IMAGE Consortium	AA000837
2850 E	12 021GA1EV	ESTs, Highly similar to GLUCOSE-6-PHOSPHATASE [Homo sapiens] {IMAGE:424968}	424968 IMAGE Consortium	W99912
2851 G	2 021GA1EV	NEURONAL PROTEIN 3.1 {IMAGE:425866}	425866 IMAGE Consortium	AA000945
2852 G	4 021GA1EV	ESTs {IMAGE:424848}	424848 IMAGE Consortium	W98118
2853 G	6 021GA1EV	IMAGE EST {IMAGE:426492}	426492 IMAGE Consortium	AA002317
2854 G	8 021GA1EV	Mus musculus beta5A integrin mRNA, complete cds {IMAGE:426454}	426454 IMAGE Consortium	AA002481
2855 G	10 021GA1EV	ESTs {IMAGE:425297}	425297 IMAGE Consortium	W99850
2856 G	12 021GA1EV	IMAGE EST {IMAGE:425544}	425544 IMAGE Consortium	AA000328
2857 A	2 021NA1EW	65 KD YES-ASSOCIATED PROTEIN {IMAGE:436977}	436977 IMAGE Consortium	AA002763

2858 A	4	021NA1EW	IMAGE EST {IMAGE:436935}	436935	IMAGE Consortium	AA002625
2859 A	6	021NA1EW	IMAGE EST {IMAGE:437480}	437480	IMAGE Consortium	AA003791
2860 A	8	021NA1EW	IMAGE EST {IMAGE:437537}	437537	IMAGE Consortium	AA003578
2861 A	10	021NA1EW	Zinc finger protein 144 {IMAGE:439450}	439450	IMAGE Consortium	AA004082
2862 A	12	021NA1EW	IMAGE EST {IMAGE:439262}	439262	IMAGE Consortium	AA003885
2863 C	2	021NA1EW	ESTs, Highly similar to SIGNAL RECOGNITION PARTICLE 68 KD PROTEIN [Canis familiaris] {IMAGE:437257}	437257	IMAGE Consortium	AA003646
2864 C	4	021NA1EW	ESTs {IMAGE:437601}	437601	IMAGE Consortium	AA003728
2865 C	6	021NA1EW	IMAGE EST {IMAGE:437633}	437633	IMAGE Consortium	AA003786
2866 C	8	021NA1EW	ESTs {IMAGE:439211}	439211	IMAGE Consortium	AA003898
2867 C	10	021NA1EW	M.musculus mRNA for novel stromal cell protein {IMAGE:439215}	439215	IMAGE Consortium	AA003900
2868 C	12	021NA1EW	IMAGE EST {IMAGE:439460}	439460	IMAGE Consortium	AA004149
2869 E	2	021NA1EW	Membrane protein, palmitoylated (55 kDa) {IMAGE:437290}	437290	IMAGE Consortium	AA003740
2870 E	4	021NA1EW	Protein kinase C substrate 80K-H {IMAGE:439239}	439239	IMAGE Consortium	AA003924
2871 E	6	021NA1EW	ESTs, Weakly similar to similar to pyridoxamine 5'-phosphate oxidase [C.elegans] {IMAGE:438286}	438286	IMAGE Consortium	AA008419
2872 E	8	021NA1EW	IMAGE EST {IMAGE:438604}	438604	IMAGE Consortium	AA050979
2873 E	10	021NA1EW	IMAGE EST {IMAGE:427596}	427596	IMAGE Consortium	AA002544
2874 E	12	021NA1EW	IMAGE EST {IMAGE:436994}	436994	IMAGE Consortium	AA002767
2875 G	2	021NA1EW	IMAGE EST {IMAGE:439737}	439737	IMAGE Consortium	AA014511
2876 G	4	021NA1EW	IMAGE EST {IMAGE:439861}	439861	IMAGE Consortium	AA008503
2877 G	6	021NA1EW	RAB24, member RAS oncogene family {IMAGE:438755}	438755	IMAGE Consortium	AA051083
2878 G	8	021NA1EW	IMAGE EST {IMAGE:438792}	438792	IMAGE Consortium	AA008317
2879 G	10	021NA1EW	ESTs, Highly similar to N-methyl-D-aspartate receptor glutamate-binding chain [R.norvegicus] {IMAGE:439108}	439108	IMAGE Consortium	AA003823
2880 G	12	021NA1EW	IMAGE EST {IMAGE:437602}	437602	IMAGE Consortium	AA003725
2881 A	2	021UA1EX	IMAGE EST {IMAGE:437220}	437220	IMAGE Consortium	AA003536
2882 A	4	021UA1EX	IMAGE EST {IMAGE:437130}	437130	IMAGE Consortium	AA003408
2883 A	6	021UA1EX	IMAGE EST {IMAGE:437626}	437626	IMAGE Consortium	AA003731
2884 A	8	021UA1EX	ESTs {IMAGE:437301}	437301	IMAGE Consortium	AA003680
2885 A	10	021UA1EX	ESTs, Weakly similar to EF-7 [M.musculus] {IMAGE:437743}	437743	IMAGE Consortium	AA007870
2886 A	12	021UA1EX	IMAGE EST {IMAGE:439479}	439479	IMAGE Consortium	AA004179
2887 C	2	021UA1EX	IMAGE EST {IMAGE:439236}	439236	IMAGE Consortium	AA003915
2888 C	4	021UA1EX	Lymphocyte protein tyrosine kinase {IMAGE:438125}	438125	IMAGE Consortium	AA008016
2889 C	6	021UA1EX	ESTs {IMAGE:438372}	438372	IMAGE Consortium	AA008157
2890 C	8	021UA1EX	IMAGE EST {IMAGE:438376}	438376	IMAGE Consortium	AA008228
2891 C	10	021UA1EX	IMAGE EST {IMAGE:439763}	439763	IMAGE Consortium	AA008579
2892 C	12	021UA1EX	IMAGE EST {IMAGE:439814}	439814	IMAGE Consortium	AA008591
2893 E	2	021UA1EX	IMAGE EST {IMAGE:440290}	440290	IMAGE Consortium	AA016693
2894 E	4	021UA1EX	IMAGE EST {IMAGE:441754}	441754	IMAGE Consortium	AA013818
2895 E	6	021UA1EX	ESTs {IMAGE:440534}	440534	IMAGE Consortium	AA014595
2896 E	8	021UA1EX	ESTs {IMAGE:441992}	441992	IMAGE Consortium	AA014730

2897 E	10 021UA1EX	IMAGE EST {IMAGE:440819}	440819 IMAGE Consortium	AA009302
2898 E	12 021UA1EX	IMAGE EST {IMAGE:439647}	439647 IMAGE Consortium	AA008708
2899 G	2 021UA1EX	E26 avian leukemia oncogene 2, 3' domain {IMAGE:440564}	440564 IMAGE Consortium	AA014761
2900 G	4 021UA1EX	ESTs, Highly similar to ELONGATION FACTOR 1-DELTA [Homo sapiens] {IMAGE:442585}	442585 IMAGE Consortium	AA015365
2901 G	6 021UA1EX	IMAGE EST {IMAGE:441176}	441176 IMAGE Consortium	AA009059
2902 G	8 021UA1EX	IMAGE EST {IMAGE:441276}	441276 IMAGE Consortium	AA011790
2903 G	10 021UA1EX	IMAGE EST {IMAGE:441521}	441521 IMAGE Consortium	AA013896
2904 G	12 021UA1EX	IMAGE EST {IMAGE:442403}	442403 IMAGE Consortium	AA015351
2905 A	2 0211A1EY	IMAGE EST {IMAGE:442805}	442805 IMAGE Consortium	AA035872
2906 A	4 0211A1EY	EST {IMAGE:442793}	442793 IMAGE Consortium	AA036486
2907 A	6 0211A1EY	Mus musculus MAST205 protein kinase mRNA, complete cds {IMAGE:444956}	444956 IMAGE Consortium	AA016762
2908 A	8 0211A1EY	ESTs {IMAGE:445505}	445505 IMAGE Consortium	AA018013
2909 A	10 0211A1EY	ESTs, Moderately similar to KERATIN, TYPE II CYTOSKELETAL 5 [Homo sapiens] {IMAGE:443870}	443870 IMAGE Consortium	AA014127
2910 A	12 0211A1EY	ESTs, Weakly similar to ORF YGL231c [S.cerevisiae] {IMAGE:442681}	442681 IMAGE Consortium	AA015149
2911 C	2 0211A1EY	ESTs {IMAGE:442508}	442508 IMAGE Consortium	AA015295
2912 C	4 0211A1EY	IMAGE EST {IMAGE:442979}	442979 IMAGE Consortium	AA014174
2913 C	6 0211A1EY	ESTs {IMAGE:442842}	442842 IMAGE Consortium	AA014194
2914 C	8 0211A1EY	ESTs {IMAGE:443408}	443408 IMAGE Consortium	AA013981
2915 C	10 0211A1EY	ESTs {IMAGE:443441}	443441 IMAGE Consortium	AA014031
2916 C	12 0211A1EY	IMAGE EST {IMAGE:445358}	445358 IMAGE Consortium	AA017876
2917 E	2 0211A1EY	IMAGE EST {IMAGE:444283}	444283 IMAGE Consortium	AA015180
2918 E	4 0211A1EY	BRAIN PROTEIN D3 {IMAGE:444794}	444794 IMAGE Consortium	AA016422
2919 E	6 0211A1EY	IMAGE EST {IMAGE:445088}	445088 IMAGE Consortium	AA017737
2920 E	8 0211A1EY	IMAGE EST {IMAGE:443711}	443711 IMAGE Consortium	AA013757
2921 E	10 0211A1EY	Abelson murine leukemia oncogene {IMAGE:444107}	444107 IMAGE Consortium	AA014945
2922 E	12 0211A1EY	Homeo box, msh-like 2 {IMAGE:444842}	444842 IMAGE Consortium	AA016730
2923 G	2 0211A1EY	EST {IMAGE:444360}	444360 IMAGE Consortium	AA016889
2924 G	4 0211A1EY	Mus musculus S100A3 calcium binding protein mRNA, complete cds {IMAGE:442707}	442707 IMAGE Consortium	AA015155
2925 G	6 0211A1EY	ESTs {IMAGE:442964}	442964 IMAGE Consortium	AA014141
2926 G	8 0211A1EY	ESTs {IMAGE:445567}	445567 IMAGE Consortium	AA017848
2927 G	10 0211A1EY	ESTs, Weakly similar to PROBABLE UBIQUITIN CARBOXYL-TERMINAL HYDROLASE [M.musculus] {IMAGE:445527}	445527 IMAGE Consortium	AA017921
2928 G	12 0211A1EY	IMAGE EST {IMAGE:445988}	445988 IMAGE Consortium	AA020051
2929 A	2 0218A1EZ	IMAGE EST {IMAGE:441331}	441331 IMAGE Consortium	AA009097
2930 A	4 0218A1EZ	IMAGE EST {IMAGE:441873}	441873 IMAGE Consortium	AA013573
2931 A	6 0218A1EZ	IMAGE EST {IMAGE:439649}	439649 IMAGE Consortium	AA008497
2932 A	8 0218A1EZ	Mus musculus BM28 homolog mRNA, complete cds	441229 IMAGE Consortium	AA011839

		{IMAGE:441229}		
2933 A	10 0218A1EZ	IMAGE EST {IMAGE:440152}	440152 IMAGE Consortium	AA016631
2934 A	12 0218A1EZ	Mus musculus homer-2a (homer-2a) mRNA, complete cds {IMAGE:441857}	441857 IMAGE Consortium	AA013888
2935 C	2 0218A1EZ	ESTs {IMAGE:441853}	441853 IMAGE Consortium	AA014063
2936 C	4 0218A1EZ	IMAGE EST {IMAGE:442111}	442111 IMAGE Consortium	AA013551
2937 C	6 0218A1EZ	ESTs, Moderately similar to probable ataxia-telangiectasia group D protein [H.sapiens] {IMAGE:441290}	441290 IMAGE Consortium	AA009191
2938 C	8 0218A1EZ	Cell division cycle 25A {IMAGE:441394}	441394 IMAGE Consortium	AA008925
2939 C	10 0218A1EZ	IMAGE EST {IMAGE:441350}	441350 IMAGE Consortium	AA009269
2940 C	12 0218A1EZ	ESTs, Moderately similar to BUTYROPHILIN PRECURSOR [Bos taurus] {IMAGE:440322}	440322 IMAGE Consortium	AA011754
2941 E	2 0218A1EZ	Mus musculus 14-3-3 protein sigma mRNA, complete cds {IMAGE:440755}	440755 IMAGE Consortium	AA009229
2942 E	4 0218A1EZ	ESTs {IMAGE:442433}	442433 IMAGE Consortium	AA015547
2943 E	6 0218A1EZ	ESTs, Weakly similar to ZINC FINGER PROTEIN ZFP-27 [Mus musculus] {IMAGE:444796}	444796 IMAGE Consortium	AA016423
2944 E	8 0218A1EZ	ESTs, Moderately similar to TFIIIC2 subunit [H.sapiens] {IMAGE:442894}	442894 IMAGE Consortium	AA014209
2945 E	10 0218A1EZ	IMAGE EST {IMAGE:444955}	444955 IMAGE Consortium	AA016765
2946 E	12 0218A1EZ	IMAGE EST {IMAGE:443516}	443516 IMAGE Consortium	AA013571
2947 G	2 0218A1EZ	Reticulocalbin {IMAGE:444388}	444388 IMAGE Consortium	AA016913
2948 G	4 0218A1EZ	ESTs, Highly similar to ERYTHROID KRUEPPEL-LIKE TRANSCRIPTION FACTOR [Mus musculus] {IMAGE:444844}	444844 IMAGE Consortium	AA016731
2949 G	6 0218A1EZ	ESTs, Highly similar to TRANSCRIPTIONAL REGULATORY PROTEIN RPD3 [Saccharomyces cerevisiae] {IMAGE:444964}	444964 IMAGE Consortium	AA017742
2950 G	8 0218A1EZ	IMAGE EST {IMAGE:445236}	445236 IMAGE Consortium	AA017756
2951 G	10 0218A1EZ	Mouse procollagen type V alpha 2 (Col5a-2) mRNA, complete cds {IMAGE:445075}	445075 IMAGE Consortium	AA023549
2952 G	12 0218A1EZ	ESTs, Highly similar to GLYCOLIPID TRANSFER PROTEIN [Sus scrofa] {IMAGE:445946}	445946 IMAGE Consortium	AA020219
2953 A	2 0218A1F0	EST {IMAGE:455559}	455559 IMAGE Consortium	AA020485
2954 A	4 0218A1F0	ESTs {IMAGE:457270}	457270 IMAGE Consortium	AA023648
2955 A	6 0218A1F0	IMAGE EST {IMAGE:455866}	455866 IMAGE Consortium	AA023135
2956 A	8 0218A1F0	Janus kinase 3 {IMAGE:457114}	457114 IMAGE Consortium	AA023670
2957 A	10 0218A1F0	ESTs, Moderately similar to COP-COATED VESICLE MEMBRANE PROTEIN P24 PRECURSOR [Cricetulus griseus] {IMAGE:455457}	455457 IMAGE Consortium	AA020489
2958 A	12 0218A1F0	ESTs {IMAGE:457510}	457510 IMAGE Consortium	AA021895
2959 C	2 0218A1F0	Notch gene homolog 4, (Drosophila) {IMAGE:457150}	457150 IMAGE Consortium	AA022302
2960 C	4 0218A1F0	Carcinoembryonic antigen 2 {IMAGE:455869}	455869 IMAGE Consortium	AA023140
2961 C	6 0218A1F0	ESTs, Weakly similar to PIAS3 [M.musculus]	458298 IMAGE Consortium	AA024336



		{IMAGE:458298}		
2962 C	8 0218A1F0	ESTs {IMAGE:458513}	458513 IMAGE Consortium	AA023949
2963 C	10 0218A1F0	Mus musculus mRNA for heme-binding protein, complete cds {IMAGE:457198}	457198 IMAGE Consortium	AA022130
2964 C	12 0218A1F0	IMAGE EST {IMAGE:457478}	457478 IMAGE Consortium	AA021671
2965 E	2 0218A1F0	ESTs {IMAGE:457541}	457541 IMAGE Consortium	AA021940
2966 E	4 0218A1F0	ESTs, Weakly similar to brain carcinoembryonic antigen [M.musculus] {IMAGE:456237}	456237 IMAGE Consortium	AA023268
2967 E	6 0218A1F0	IMAGE EST {IMAGE:455366}	455366 IMAGE Consortium	AA020425
2968 E	8 0218A1F0	ESTs, Highly similar to HETEROGENEOUS NUCLEAR RIBONUCLEOPROTEIN 27C [Drosophila melanogaster] {IMAGE:457976}	457976 IMAGE Consortium	AA030214
2969 E	10 0218A1F0	IMAGE EST {IMAGE:457840}	457840 IMAGE Consortium	AA030159
2970 E	12 0218A1F0	ESTs {IMAGE:456261}	456261 IMAGE Consortium	AA023276
2971 G	2 0218A1F0	Mus musculus SMAD3 (mSmad3) mRNA, complete cds {IMAGE:456861}	456861 IMAGE Consortium	AA023641
2972 G	4 0218A1F0	IMAGE EST {IMAGE:455576}	455576 IMAGE Consortium	AA020532
2973 G	6 0218A1F0	ESTs, Highly similar to ERYTHROCYTE ADDUCIN BETA SUBUNIT [Homo sapiens] {IMAGE:457503}	457503 IMAGE Consortium	AA021816
2974 G	8 0218A1F0	IMAGE EST {IMAGE:457532}	457532 IMAGE Consortium	AA021790
2975 G	10 0218A1F0	ESTs {IMAGE:456064}	456064 IMAGE Consortium	AA023101
2976 G	12 0218A1F0	ESTs {IMAGE:456249}	456249 IMAGE Consortium	AA023245
2977 A	2 021FA1F1	IMAGE EST {IMAGE:456204}	456204 IMAGE Consortium	AA023221
2978 A	4 021FA1F1	IMAGE EST {IMAGE:463556}	463556 IMAGE Consortium	AA028638
2979 A	6 021FA1F1	IMAGE EST {IMAGE:458830}	458830 IMAGE Consortium	AA024036
2980 A	8 021FA1F1	IMAGE EST {IMAGE:459509}	459509 IMAGE Consortium	AA027451
2981 A	10 021FA1F1	ESTs {IMAGE:459380}	459380 IMAGE Consortium	AA027459
2982 A	12 021FA1F1	ESTs, Highly similar to CALTRACTIN [Homo sapiens] {IMAGE:459822}	459822 IMAGE Consortium	AA027609
2983 C	2 021FA1F1	ESTs, Highly similar to TRYPSINOGEN [Bos taurus] {IMAGE:464085}	464085 IMAGE Consortium	AA028356
2984 C	4 021FA1F1	IMAGE EST {IMAGE:464358}	464358 IMAGE Consortium	AA030205
2985 C	6 021FA1F1	IMAGE EST {IMAGE:464391}	464391 IMAGE Consortium	AA030506
2986 C	8 021FA1F1	IMAGE EST {IMAGE:464712}	464712 IMAGE Consortium	AA030061
2987 C	10 021FA1F1	IMAGE EST {IMAGE:464575}	464575 IMAGE Consortium	AA030534
2988 C	12 021FA1F1	IMAGE EST {IMAGE:463464}	463464 IMAGE Consortium	AA027728
2989 E	2 021FA1F1	ESTs {IMAGE:463200}	463200 IMAGE Consortium	AA030361
2990 E	4 021FA1F1	IMAGE EST {IMAGE:463157}	463157 IMAGE Consortium	AA030609
2991 E	6 021FA1F1	IMAGE EST {IMAGE:463257}	463257 IMAGE Consortium	AA030598
2992 E	8 021FA1F1	IMAGE EST {IMAGE:463024}	463024 IMAGE Consortium	AA030579
2993 E	10 021FA1F1	IMAGE EST {IMAGE:463050}	463050 IMAGE Consortium	AA030476
2994 E	12 021FA1F1	IMAGE EST {IMAGE:463567}	463567 IMAGE Consortium	AA028675
2995 G	2 021FA1F1	IMAGE EST {IMAGE:464800}	464800 IMAGE Consortium	AA030796

2996 G	4 021FA1F1	IMAGE EST {IMAGE:463457}	463457 IMAGE Consortium	AA030650
2997 G	6 021FA1F1	IMAGE EST {IMAGE:463956}	463956 IMAGE Consortium	AA028411
2998 G	8 021FA1F1	IMAGE EST {IMAGE:459377}	459377 IMAGE Consortium	AA027478
2999 G	10 021FA1F1	IMAGE EST {IMAGE:459538}	459538 IMAGE Consortium	AA027510
3000 G	12 021FA1F1	IMAGE EST {IMAGE:463249}	463249 IMAGE Consortium	AA030377
3001 A	2 021MA1F2	IMAGE EST {IMAGE:459254}	459254 IMAGE Consortium	AA030253
3002 A	4 021MA1F2	GATA-binding protein 2 {IMAGE:459700}	459700 IMAGE Consortium	AA027569
3003 A	6 021MA1F2	IMAGE EST {IMAGE:464265}	464265 IMAGE Consortium	AA032929
3004 A	8 021MA1F2	IMAGE EST {IMAGE:464177}	464177 IMAGE Consortium	AA028591
3005 A	10 021MA1F2	Mus musculus mRNA for Bloom's syndrome protein {IMAGE:464447}	464447 IMAGE Consortium	AA030433
3006 A	12 021MA1F2	Insulin-like growth factor 2 {IMAGE:464598}	464598 IMAGE Consortium	AA030546
3007 C	2 021MA1F2	Mouse mRNA for mCDC46 protein, complete cds {IMAGE:466047}	466047 IMAGE Consortium	AA031056
3008 C	4 021MA1F2	ESTs {IMAGE:468050}	468050 IMAGE Consortium	AA015312
3009 C	6 021MA1F2	IMAGE EST {IMAGE:468270}	468270 IMAGE Consortium	AA014357
3010 C	8 021MA1F2	Mus musculus synapsin Ib (SynI) mRNA, complete cds {IMAGE:466295}	466295 IMAGE Consortium	AA033398
3011 C	10 021MA1F2	IMAGE EST {IMAGE:466810}	466810 IMAGE Consortium	AA031072
3012 C	12 021MA1F2	ESTs {IMAGE:465348}	465348 IMAGE Consortium	AA032414
3013 E	2 021MA1F2	IMAGE EST {IMAGE:466099}	466099 IMAGE Consortium	AA033340
3014 E	4 021MA1F2	Mus musculus ubiquinone biosynthesis protein coq7 mRNA, complete cds {IMAGE:466542}	466542 IMAGE Consortium	AA030846
3015 E	6 021MA1F2	IMAGE EST {IMAGE:464974}	464974 IMAGE Consortium	AA032581
3016 E	8 021MA1F2	ESTs, Moderately similar to NEURONAL PROTEIN [Felis catus] {IMAGE:465653}	465653 IMAGE Consortium	AA033211
3017 E	10 021MA1F2	IMAGE EST {IMAGE:467194}	467194 IMAGE Consortium	AA034788
3018 E	12 021MA1F2	IMAGE EST {IMAGE:467455}	467455 IMAGE Consortium	AA034710
3019 G	2 021MA1F2	Mouse procollagen type V alpha 2 (Col5a-2) mRNA, complete cds {IMAGE:467107}	467107 IMAGE Consortium	AA034564
3020 G	4 021MA1F2	IMAGE EST {IMAGE:467207}	467207 IMAGE Consortium	AA034847
3021 G	6 021MA1F2	IMAGE EST {IMAGE:467480}	467480 IMAGE Consortium	AA036602
3022 G	8 021MA1F2	IMAGE EST {IMAGE:467785}	467785 IMAGE Consortium	AA036495
3023 G	10 021MA1F2	IMAGE EST {IMAGE:468319}	468319 IMAGE Consortium	AA014368
3024 G	12 021MA1F2	IMAGE EST {IMAGE:466411}	466411 IMAGE Consortium	AA034766
3025 A	2 021TA1F3	Mus musculus mRNA for neutral sphingomyelinase {IMAGE:464204}	464204 IMAGE Consortium	AA028477
3026 A	4 021TA1F3	ESTs {IMAGE:464619}	464619 IMAGE Consortium	AA033008
3027 A	6 021TA1F3	ESTs, Weakly similar to Ydr299wp [S.cerevisiae] {IMAGE:463028}	463028 IMAGE Consortium	AA030581
3028 A	8 021TA1F3	IMAGE EST {IMAGE:465002}	465002 IMAGE Consortium	AA032443
3029 A	10 021TA1F3	ESTs {IMAGE:465175}	465175 IMAGE Consortium	AA030752
3030 A	12 021TA1F3	IMAGE EST {IMAGE:458948}	458948 IMAGE Consortium	AA024256

3031 C	2 021TA1F3	ESTs {IMAGE:459143}	459143 IMAGE Consortium	AA024085
3032 C	4 021TA1F3	Insulin-like growth factor 2 {IMAGE:463135}	463135 IMAGE Consortium	AA030603
3033 C	6 021TA1F3	IMAGE EST {IMAGE:463145}	463145 IMAGE Consortium	AA030129
3034 C	8 021TA1F3	IMAGE EST {IMAGE:464635}	464635 IMAGE Consortium	AA030556
3035 C	10 021TA1F3	Male enhanced antigen 1 {IMAGE:463746}	463746 IMAGE Consortium	AA028786
3036 C	12 021TA1F3	ESTs {IMAGE:458944}	458944 IMAGE Consortium	AA024255
3037 E	2 021TA1F3	IMAGE EST {IMAGE:464803}	464803 IMAGE Consortium	AA030833
3038 E	4 021TA1F3	ESTs {IMAGE:465218}	465218 IMAGE Consortium	AA032559
3039 E	6 021TA1F3	Farnesyltransferase, CAAX box, alpha {IMAGE:465152}	465152 IMAGE Consortium	AA032725
3040 E	8 021TA1F3	ESTs, Moderately similar to E1B 19K/Bcl-2-interacting protein Nip3 [H.sapiens] {IMAGE:464020}	464020 IMAGE Consortium	AA028342
3041 E	10 021TA1F3	Eph receptor A2 {IMAGE:459170}	459170 IMAGE Consortium	AA024123
3042 E	12 021TA1F3	Keratin complex 1, acidic, gene 19 {IMAGE:464060}	464060 IMAGE Consortium	AA028346
3043 G	2 021TA1F3	Tissue specific transplantation antigen P91A {IMAGE:463757}	463757 IMAGE Consortium	AA028215
3044 G	4 021TA1F3	ESTs, Moderately similar to MULTIDRUG RESISTANCE-ASSOCIATED PROTEIN 1 [Homo sapiens] {IMAGE:459410}	459410 IMAGE Consortium	AA027420
3045 G	6 021TA1F3	ESTs {IMAGE:464084}	464084 IMAGE Consortium	AA028352
3046 G	8 021TA1F3	IMAGE EST {IMAGE:459661}	459661 IMAGE Consortium	AA027559
3047 G	10 021TA1F3	Chaperonin subunit 4 (delta) {IMAGE:459668}	459668 IMAGE Consortium	AA027583
3048 G	12 021TA1F3	IMAGE EST {IMAGE:463231}	463231 IMAGE Consortium	AA030374
3049 A	2 0210A1F4	ESTs, Highly similar to peroxin 12 [H.sapiens] {IMAGE:466120}	466120 IMAGE Consortium	AA033343
3050 A	4 0210A1F4	IMAGE EST {IMAGE:466730}	466730 IMAGE Consortium	AA031057
3051 A	6 0210A1F4	IMAGE EST {IMAGE:466807}	466807 IMAGE Consortium	AA031077
3052 A	8 0210A1F4	IMAGE EST {IMAGE:465481}	465481 IMAGE Consortium	AA032754
3053 A	10 0210A1F4	ESTs, Highly similar to CART PROTEIN PRECURSOR [Rattus norvegicus] {IMAGE:466874}	466874 IMAGE Consortium	AA031207
3054 A	12 0210A1F4	IMAGE EST {IMAGE:465620}	465620 IMAGE Consortium	AA032760
3055 C	2 0210A1F4	Signal sequence receptor, delta {IMAGE:466568}	466568 IMAGE Consortium	AA030906
3056 C	4 0210A1F4	ESTs {IMAGE:466624}	466624 IMAGE Consortium	AA030987
3057 C	6 0210A1F4	IMAGE EST {IMAGE:467337}	467337 IMAGE Consortium	AA034601
3058 C	8 0210A1F4	IMAGE EST {IMAGE:466206}	466206 IMAGE Consortium	AA033390
3059 C	10 0210A1F4	IMAGE EST {IMAGE:466413}	466413 IMAGE Consortium	AA034767
3060 C	12 0210A1F4	IMAGE EST {IMAGE:466788}	466788 IMAGE Consortium	AA031142
3061 E	2 0210A1F4	ESTs, Weakly similar to P9513.2 gene product [S.cerevisiae] {IMAGE:464919}	464919 IMAGE Consortium	AA031240
3062 E	4 0210A1F4	IMAGE EST {IMAGE:466573}	466573 IMAGE Consortium	AA030915
3063 E	6 0210A1F4	Mus musculus hemin-sensitive initiation factor 2 alpha kinase mRNA, complete cds {IMAGE:466971}	466971 IMAGE Consortium	AA036546
3064 E	8 0210A1F4	ESTs {IMAGE:467055}	467055 IMAGE Consortium	AA034799
3065 E	10 0210A1F4	ESTs, Highly similar to Contains C3HC4 type zinc finger signature [H.sapiens] {IMAGE:467355}	467355 IMAGE Consortium	AA036618

3066 E	12 0210A1F4	IMAGE EST {IMAGE:466046}	466046 IMAGE Consortium	AA031116
3067 G	2 0210A1F4	IMAGE EST {IMAGE:466018}	466018 IMAGE Consortium	AA031105
3068 G	4 0210A1F4	IMAGE EST {IMAGE:466418}	466418 IMAGE Consortium	AA032326
3069 G	6 0210A1F4	ESTs {IMAGE:466141}	466141 IMAGE Consortium	AA036585
3070 G	8 0210A1F4	IMAGE EST {IMAGE:602403}	602403 IMAGE Consortium	AA146325
3071 G	10 0210A1F4	IMAGE EST {IMAGE:466780}	466780 IMAGE Consortium	AA031140
3072 G	12 0210A1F4	IMAGE EST {IMAGE:466546}	466546 IMAGE Consortium	AA030848
3073 A	2 0217A1F5	Solute carrier family 12, member 2 {IMAGE:466122}	466122 IMAGE Consortium	AA033344
3074 A	4 0217A1F5	Cytochrome P450, 2e1, ethanol inducible {IMAGE:468633}	468633 IMAGE Consortium	AA014577
3075 A	6 0217A1F5	IMAGE EST {IMAGE:466628}	466628 IMAGE Consortium	AA030989
3076 A	8 0217A1F5	ESTs, Weakly similar to Similar to sulfatase [C.elegans] {IMAGE:465592}	465592 IMAGE Consortium	AA032808
3077 A	10 0217A1F5	IMAGE EST {IMAGE:466013}	466013 IMAGE Consortium	AA031108
3078 A	12 0217A1F5	IMAGE EST {IMAGE:467364}	467364 IMAGE Consortium	AA034731
3079 C	2 0217A1F5	IMAGE EST {IMAGE:469019}	469019 IMAGE Consortium	AA036034
3080 C	4 0217A1F5	IMAGE EST {IMAGE:472562}	472562 IMAGE Consortium	AA036314
3081 C	6 0217A1F5	IMAGE EST {IMAGE:472432}	472432 IMAGE Consortium	AA036186
3082 C	8 0217A1F5	ESTs, Moderately similar to hypothetical protein [M.musculus] {IMAGE:472962}	472962 IMAGE Consortium	AA038043
3083 C	10 0217A1F5	ESTs {IMAGE:475512}	475512 IMAGE Consortium	AA042017
3084 C	12 0217A1F5	IMAGE EST {IMAGE:474365}	474365 IMAGE Consortium	AA038774
3085 E	2 0217A1F5	IMAGE EST {IMAGE:472444}	472444 IMAGE Consortium	AA036152
3086 E	4 0217A1F5	ESTs {IMAGE:472850}	472850 IMAGE Consortium	AA038004
3087 E	6 0217A1F5	ESTs {IMAGE:473415}	473415 IMAGE Consortium	AA038443
3088 E	8 0217A1F5	ESTs {IMAGE:475298}	475298 IMAGE Consortium	AA041964
3089 E	10 0217A1F5	ESTs, Moderately similar to HYPOTHETICAL 34.1 KD PROTEIN ZK1098.4 IN CHROMOSOME III [Caenorhabditis elegans] {IMAGE:473831}	473831 IMAGE Consortium	AA038433
3090 E	12 0217A1F5	IMAGE EST {IMAGE:468943}	468943 IMAGE Consortium	AA036279
3091 G	2 0217A1F5	ESTs, Moderately similar to ADENYLATE CYCLASE, TYPE VII [M.musculus] {IMAGE:472258}	472258 IMAGE Consortium	AA036065
3092 G	4 0217A1F5	T-LYMPHOCYTE ACTIVATED PROTEIN {IMAGE:472605}	472605 IMAGE Consortium	AA038052
3093 G	6 0217A1F5	ESTs {IMAGE:474037}	474037 IMAGE Consortium	AA038452
3094 G	8 0217A1F5	IMAGE EST {IMAGE:469021}	469021 IMAGE Consortium	AA036035
3095 G	10 0217A1F5	IMAGE EST {IMAGE:472282}	472282 IMAGE Consortium	AA036113
3096 G	12 0217A1F5	Treacher Collins Franceschetti syndrome 1, homolog {IMAGE:473404}	473404 IMAGE Consortium	AA038551
3097 A	2 021EA1F6	IMAGE EST {IMAGE:476321}	476321 IMAGE Consortium	AA050391
3098 A	4 021EA1F6	IMAGE EST {IMAGE:477905}	477905 IMAGE Consortium	AA049221
3099 A	6 021EA1F6	IMAGE EST {IMAGE:478154}	478154 IMAGE Consortium	AA049601
3100 A	8 021EA1F6	ESTs, Highly similar to PROBABLE PROTEIN DISULFIDE ISOMERASE P5 PRECURSOR [Mesocricetus auratus] {IMAGE:478021}	478021 IMAGE Consortium	AA049474

3101 A	10 021EA1F6	Protein kinase, cAMP dependent, catalytic, beta {IMAGE:478265}	478265 IMAGE Consortium	AA049744
3102 A	12 021EA1F6	IMAGE EST {IMAGE:475381}	475381 IMAGE Consortium	AA041819
3103 C	2 021EA1F6	IMAGE EST {IMAGE:477834}	477834 IMAGE Consortium	AA049391
3104 C	4 021EA1F6	IMAGE EST {IMAGE:476625}	476625 IMAGE Consortium	AA050521
3105 C	6 021EA1F6	IMAGE EST {IMAGE:476562}	476562 IMAGE Consortium	AA050741
3106 C	8 021EA1F6	IMAGE EST {IMAGE:478027}	478027 IMAGE Consortium	AA049361
3107 C	10 021EA1F6	ESTs, Highly similar to 33 KD HOUSEKEEPING PROTEIN [Homo sapiens] {IMAGE:476895}	476895 IMAGE Consortium	AA050837
3108 C	12 021EA1F6	ESTs, Highly similar to POSSIBLE GUSTATORY RECEPTOR TYPE B [Bos taurus] {IMAGE:476044}	476044 IMAGE Consortium	AA050273
3109 E	2 021EA1F6	ESTs {IMAGE:478323}	478323 IMAGE Consortium	AA049649
3110 E	4 021EA1F6	ESTs {IMAGE:478333}	478333 IMAGE Consortium	AA049697
3111 E	6 021EA1F6	Retinoid X receptor beta {IMAGE:475673}	475673 IMAGE Consortium	AA050298
3112 E	8 021EA1F6	IMAGE EST {IMAGE:477521}	477521 IMAGE Consortium	AA048486
3113 E	10 021EA1F6	ESTs {IMAGE:476274}	476274 IMAGE Consortium	AA050571
3114 E	12 021EA1F6	ESTs {IMAGE:477860}	477860 IMAGE Consortium	AA049399
3115 G	2 021EA1F6	ESTs {IMAGE:476387}	476387 IMAGE Consortium	AA050615
3116 G	4 021EA1F6	M.musculus es64 mRNA {IMAGE:476131}	476131 IMAGE Consortium	AA049878
3117 G	6 021EA1F6	Phosphoribosylglycinamide formyltransferase {IMAGE:476629}	476629 IMAGE Consortium	AA050540
3118 G	8 021EA1F6	ESTs, Moderately similar to ISOVALERYL-COA DEHYDROGENASE PRECURSOR [Rattus norvegicus] {IMAGE:478220}	478220 IMAGE Consortium	AA049730
3119 G	10 021EA1F6	ESTs {IMAGE:476887}	476887 IMAGE Consortium	AA050516
3120 G	12 021EA1F6	Lethal giant larvae homolog {IMAGE:475755}	475755 IMAGE Consortium	AA050402
3121 A	2 021LA1F7	Mouse DNA for tob family, complete cds {IMAGE:468859}	468859 IMAGE Consortium	AA036170
3122 A	4 021LA1F7	IMAGE EST {IMAGE:472332}	472332 IMAGE Consortium	AA036347
3123 A	6 021LA1F7	IMAGE EST {IMAGE:472942}	472942 IMAGE Consortium	AA038039
3124 A	8 021LA1F7	ESTs, Highly similar to transcription factor E2F-4 [M.musculus] {IMAGE:475443}	475443 IMAGE Consortium	AA041739
3125 A	10 021LA1F7	M.musculus mRNA for mTGIF protein {IMAGE:474339}	474339 IMAGE Consortium	AA038657
3126 A	12 021LA1F7	IMAGE EST {IMAGE:468849}	468849 IMAGE Consortium	AA035976
3127 C	2 021LA1F7	IMAGE EST {IMAGE:474275}	474275 IMAGE Consortium	AA038782
3128 C	4 021LA1F7	IMAGE EST {IMAGE:474305}	474305 IMAGE Consortium	AA038862
3129 C	6 021LA1F7	Opioid receptor, sigma 1 {IMAGE:469124}	469124 IMAGE Consortium	AA036319
3130 C	8 021LA1F7	ESTs {IMAGE:472407}	472407 IMAGE Consortium	AA036386
3131 C	10 021LA1F7	IMAGE EST {IMAGE:472722}	472722 IMAGE Consortium	AA038172
3132 C	12 021LA1F7	IMAGE EST {IMAGE:468817}	468817 IMAGE Consortium	AA035899
3133 E	2 021LA1F7	ESTs, Highly similar to HYPOTHETICAL ZINC FINGER PROTEIN ZK686.4 IN CHROMOSOME III [Caenorhabditis elegans] {IMAGE:475572}	475572 IMAGE Consortium	AA050347
3134 E	4 021LA1F7	IMAGE EST {IMAGE:475458}	475458 IMAGE Consortium	AA041916

3135 E	6 021LA1F7	ESTs {IMAGE:475631}	475631 IMAGE Consortium	AA049807
3136 E	8 021LA1F7	ESTs {IMAGE:477552}	477552 IMAGE Consortium	AA048449
3137 E	10 021LA1F7	ESTs, Highly similar to ISOCITRATE DEHYDROGENASE [Bos taurus] {IMAGE:475796}	475796 IMAGE Consortium	AA050135
3138 E	12 021LA1F7	EST {IMAGE:477541}	477541 IMAGE Consortium	AA048447
3139 G	2 021LA1F7	IMAGE EST {IMAGE:476022}	476022 IMAGE Consortium	AA049934
3140 G	4 021LA1F7	IMAGE EST {IMAGE:475725}	475725 IMAGE Consortium	AA050124
3141 G	6 021LA1F7	IMAGE EST {IMAGE:476342}	476342 IMAGE Consortium	AA050648
3142 G	8 021LA1F7	IMAGE EST {IMAGE:476325}	476325 IMAGE Consortium	AA050393
3143 G	10 021LA1F7	ESTs, Moderately similar to PHOSPHATIDYLINOSITOL [Homo sapiens] {IMAGE:476642}	476642 IMAGE Consortium	AA050487
3144 G	12 021LA1F7	Antigen identified by monoclonal antibodies 4F2 {IMAGE:478301}	478301 IMAGE Consortium	AA049696
3145 A	2 021SA1F8	ESTs {IMAGE:477055}	477055 IMAGE Consortium	AA048155
3146 A	4 021SA1F8	IMAGE EST {IMAGE:476993}	476993 IMAGE Consortium	AA048111
3147 A	6 021SA1F8	IMAGE EST {IMAGE:476879}	476879 IMAGE Consortium	AA050884
3148 A	8 021SA1F8	ESTs, Weakly similar to UBIQUITIN-CONJUGATING ENZYME E2-16 KD [Saccharomyces cerevisiae] {IMAGE:476358}	476358 IMAGE Consortium	AA050167
3149 A	10 021SA1F8	ESTs, Highly similar to KT112 PROTEIN [Saccharomyces cerevisiae] {IMAGE:476368}	476368 IMAGE Consortium	AA050211
3150 A	12 021SA1F8	ESTs, Moderately similar to RHOMBOTIN-1 [Homo sapiens] {IMAGE:476784}	476784 IMAGE Consortium	AA050915
3151 C	2 021SA1F8	ESTs {IMAGE:477796}	477796 IMAGE Consortium	AA048412
3152 C	4 021SA1F8	M.musculus mRNA for cytochrome C oxidase subunit VII homologue {IMAGE:476180}	476180 IMAGE Consortium	AA050684
3153 C	6 021SA1F8	IMAGE EST {IMAGE:476713}	476713 IMAGE Consortium	AA050895
3154 C	8 021SA1F8	IMAGE EST {IMAGE:478037}	478037 IMAGE Consortium	AA049486
3155 C	10 021SA1F8	ESTs, Weakly similar to Tid56 protein [D.melanogaster] {IMAGE:478167}	478167 IMAGE Consortium	AA049615
3156 C	12 021SA1F8	IMAGE EST {IMAGE:475404}	475404 IMAGE Consortium	AA041823
3157 E	2 021SA1F8	IMAGE EST {IMAGE:476758}	476758 IMAGE Consortium	AA050445
3158 E	4 021SA1F8	ESTs {IMAGE:477003}	477003 IMAGE Consortium	AA048121
3159 E	6 021SA1F8	ESTs {IMAGE:479858}	479858 IMAGE Consortium	AA051642
3160 E	8 021SA1F8	IMAGE EST {IMAGE:478692}	478692 IMAGE Consortium	AA051402
3161 E	10 021SA1F8	ESTs {IMAGE:478558}	478558 IMAGE Consortium	AA051331
3162 E	12 021SA1F8	ESTs {IMAGE:479913}	479913 IMAGE Consortium	AA051561
3163 G	2 021SA1F8	ESTs, Weakly similar to ORF YKL017c [S.cerevisiae] {IMAGE:478374}	478374 IMAGE Consortium	AA049766
3164 G	4 021SA1F8	IMAGE EST {IMAGE:479576}	479576 IMAGE Consortium	AA048985
3165 G	6 021SA1F8	M.musculus mRNA for calpain-like protease {IMAGE:478504}	478504 IMAGE Consortium	AA050030
3166 G	8 021SA1F8	IMAGE EST {IMAGE:480120}	480120 IMAGE Consortium	AA051720
3167 G	10 021SA1F8	IMAGE EST {IMAGE:478855}	478855 IMAGE Consortium	AA051256

3168 G	12 021SA1F8	IMAGE EST {IMAGE:479921}	479921 IMAGE Consortium	AA051759
3169 A	2 021ZA1F9	Period homolog (Drosophila) {IMAGE:478552}	478552 IMAGE Consortium	AA051328
3170 A	4 021ZA1F9	ESTs {IMAGE:480081}	480081 IMAGE Consortium	AA051780
3171 A	6 021ZA1F9	ESTs, Moderately similar to HYPOTHETICAL 167.8 KD PROTEIN CCE1-CAP1 INTERGENIC REGION [Saccharomyces cerevisiae] {IMAGE:480113}	480113 IMAGE Consortium	AA057999
3172 A	8 021ZA1F9	ESTs {IMAGE:480236}	480236 IMAGE Consortium	AA058079
3173 A	10 021ZA1F9	IMAGE EST {IMAGE:479382}	479382 IMAGE Consortium	AA048675
3174 A	12 021ZA1F9	Guanine nucleotide binding protein, beta-2, related sequence 1 {IMAGE:479249}	479249 IMAGE Consortium	AA048915
3175 C	2 021ZA1F9	ESTs, Moderately similar to ALPHA CRYSTALLIN B CHAIN [Gallus gallus] {IMAGE:479949}	479949 IMAGE Consortium	AA051767
3176 C	4 021ZA1F9	ESTs {IMAGE:479406}	479406 IMAGE Consortium	AA049133
3177 C	6 021ZA1F9	IMAGE EST {IMAGE:479462}	479462 IMAGE Consortium	AA049255
3178 C	8 021ZA1F9	IMAGE EST {IMAGE:480334}	480334 IMAGE Consortium	AA058112
3179 C	10 021ZA1F9	ESTs, Weakly similar to ORF YGL153w [S.cerevisiae] {IMAGE:479351}	479351 IMAGE Consortium	AA049018
3180 C	12 021ZA1F9	ESTs, Moderately similar to GTP:AMP PHOSPHOTRANSFERASE MITOCHONDRIAL [R.norvegicus] {IMAGE:479750}	479750 IMAGE Consortium	AA051422
3181 E	2 021ZA1F9	Mus musculus ETO/MTG8-related protein ETO-2 mRNA, complete cds {IMAGE:479774}	479774 IMAGE Consortium	AA051563
3182 E	4 021ZA1F9	ESTs {IMAGE:480026}	480026 IMAGE Consortium	AA051507
3183 E	6 021ZA1F9	ESTs {IMAGE:480197}	480197 IMAGE Consortium	AA058059
3184 E	8 021ZA1F9	IMAGE EST {IMAGE:480162}	480162 IMAGE Consortium	AA058019
3185 E	10 021ZA1F9	IMAGE EST {IMAGE:480325}	480325 IMAGE Consortium	AA058044
3186 E	12 021ZA1F9	IMAGE EST {IMAGE:479154}	479154 IMAGE Consortium	AA049325
3187 G	2 021ZA1F9	ESTs {IMAGE:479271}	479271 IMAGE Consortium	AA048540
3188 G	4 021ZA1F9	ESTs {IMAGE:479346}	479346 IMAGE Consortium	AA049012
3189 G	6 021ZA1F9	ESTs {IMAGE:478711}	478711 IMAGE Consortium	AA051384
3190 G	8 021ZA1F9	IMAGE EST {IMAGE:478571}	478571 IMAGE Consortium	AA051309
3191 G	10 021ZA1F9	IMAGE EST {IMAGE:479805}	479805 IMAGE Consortium	AA051355
3192 G	12 021ZA1F9	IMAGE EST {IMAGE:480035}	480035 IMAGE Consortium	AA051592
3193 A	2 021JA1FA	Mouse mRNA for dbpA murine homologue, complete cds {IMAGE:481949}	481949 IMAGE Consortium	AA059953
3194 A	4 021JA1FA	ESTs, Weakly similar to lactoferrin precursor [M.musculus] {IMAGE:483799}	483799 IMAGE Consortium	AA061451
3195 A	6 021JA1FA	IMAGE EST {IMAGE:483815}	483815 IMAGE Consortium	AA061508
3196 A	8 021JA1FA	Guanosine diphosphate (GDP) dissociation inhibitor 1 {IMAGE:482355}	482355 IMAGE Consortium	AA060905
3197 A	10 021JA1FA	House mouse; Musculus domesticus female mammary carcinoma mRNA for mCDC47, complete cds {IMAGE:480852}	480852 IMAGE Consortium	AA064230
3198 A	12 021JA1FA	TIS11B PROTEIN {IMAGE:481026}	481026 IMAGE Consortium	AA060205

3199 C	2 021JA1FA	CELL SURFACE GLYCOPROTEIN MAC-1 ALPHA SUBUNIT PRECURSOR {IMAGE:483591}	483591 IMAGE Consortium	AA061908
3200 C	4 021JA1FA	Mus musculus intersectin-EH binding protein Ibp2 mRNA, partial cds {IMAGE:483477}	483477 IMAGE Consortium	AA060038
3201 C	6 021JA1FA	IMAGE EST {IMAGE:483753}	483753 IMAGE Consortium	AA061408
3202 C	8 021JA1FA	ESTs {IMAGE:480721}	480721 IMAGE Consortium	AA064091
3203 C	10 021JA1FA	Alpha-2-macroglobulin {IMAGE:482459}	482459 IMAGE Consortium	AA061055
3204 C	12 021JA1FA	IMAGE EST {IMAGE:480811}	480811 IMAGE Consortium	AA064110
3205 E	2 021JA1FA	EST {IMAGE:481950}	481950 IMAGE Consortium	AA059950
3206 E	4 021JA1FA	IMAGE EST {IMAGE:481860}	481860 IMAGE Consortium	AA059783
3207 E	6 021JA1FA	Early growth response 1 {IMAGE:480793}	480793 IMAGE Consortium	AA060716
3208 E	8 021JA1FA	IMAGE EST {IMAGE:482196}	482196 IMAGE Consortium	AA059908
3209 E	10 021JA1FA	ESTs {IMAGE:482326}	482326 IMAGE Consortium	AA063761
3210 E	12 021JA1FA	ESTs {IMAGE:482885}	482885 IMAGE Consortium	AA061958
3211 G	2 021JA1FA	IMAGE EST {IMAGE:483645}	483645 IMAGE Consortium	AA061459
3212 G	4 021JA1FA	ESTs {IMAGE:483779}	483779 IMAGE Consortium	AA061440
3213 G	6 021JA1FA	Endoglin {IMAGE:484299}	484299 IMAGE Consortium	AA080278
3214 G	8 021JA1FA	IMAGE EST {IMAGE:482189}	482189 IMAGE Consortium	AA059837
3215 G	10 021JA1FA	ESTs {IMAGE:482198}	482198 IMAGE Consortium	AA059909
3216 G	12 021JA1FA	ESTs {IMAGE:482646}	482646 IMAGE Consortium	AA061977
3217 A	2 021QA1FB	IMAGE EST {IMAGE:478643}	478643 IMAGE Consortium	AA051389
3218 A	4 021QA1FB	IMAGE EST {IMAGE:479662}	479662 IMAGE Consortium	AA051452
3219 A	6 021QA1FB	IMAGE EST {IMAGE:479045}	479045 IMAGE Consortium	AA049029
3220 A	8 021QA1FB	ESTs {IMAGE:478810}	478810 IMAGE Consortium	AA051151
3221 A	10 021QA1FB	IMAGE EST {IMAGE:480139}	480139 IMAGE Consortium	AA058016
3222 A	12 021QA1FB	Solute carrier family 4 (anion exchanger), member 2 {IMAGE:479476}	479476 IMAGE Consortium	AA048952
3223 C	2 021QA1FB	Tyrosine kinase receptor 1 {IMAGE:479679}	479679 IMAGE Consortium	AA049318
3224 C	4 021QA1FB	Fibroblast growth factor 15 {IMAGE:479758}	479758 IMAGE Consortium	AA051675
3225 C	6 021QA1FB	IMAGE EST {IMAGE:480219}	480219 IMAGE Consortium	AA058084
3226 C	8 021QA1FB	IMAGE EST {IMAGE:480008}	480008 IMAGE Consortium	AA051581
3227 C	10 021QA1FB	IMAGE EST {IMAGE:479920}	479920 IMAGE Consortium	AA051756
3228 C	12 021QA1FB	M.musculus mRNA for novel stromal cell protein {IMAGE:479383}	479383 IMAGE Consortium	AA048752
3229 E	2 021QA1FB	ESTs {IMAGE:480204}	480204 IMAGE Consortium	AA058191
3230 E	4 021QA1FB	ESTs, Weakly similar to T14B4.2 gene product [C.elegans] {IMAGE:480112}	480112 IMAGE Consortium	AA057995
3231 E	6 021QA1FB	ESTs {IMAGE:479405}	479405 IMAGE Consortium	AA049136
3232 E	8 021QA1FB	IMAGE EST {IMAGE:480323}	480323 IMAGE Consortium	AA058043
3233 E	10 021QA1FB	ESTs {IMAGE:480281}	480281 IMAGE Consortium	AA058098
3234 E	12 021QA1FB	ESTs {IMAGE:479786}	479786 IMAGE Consortium	AA051735
3235 G	2 021QA1FB	ESTs {IMAGE:480552}	480552 IMAGE Consortium	AA064188
3236 G	4 021QA1FB	Mus musculus CDK-activating kinase assembly factor	480395 IMAGE Consortium	AA058165



		p36/MAT1, complete cds {IMAGE:480395}		
3237 G	6 021QA1FB	NUCLEOSIDE DIPHOSPHATE KINASE A {IMAGE:480329}	480329 IMAGE Consortium	AA058113
3238 G	8 021QA1FB	PEPTIDYL-PROLYL CIS-TRANS ISOMERASE B PRECURSOR {IMAGE:480816}	480816 IMAGE Consortium	AA060537
3239 G	10 021QA1FB	LIM and SH3 protein 1 {IMAGE:482313}	482313 IMAGE Consortium	AA060795
3240 G	12 021QA1FB	Mus musculus mRNA for NEFA protein, complete cds {IMAGE:482883}	482883 IMAGE Consortium	AA061957
3241 A	2 021XA1FC	Glycoprotein galactosyltransferase beta 1, 4 {IMAGE:484108}	484108 IMAGE Consortium	AA079961
3242 A	4 021XA1FC	Myelin basic protein {IMAGE:480753}	480753 IMAGE Consortium	AA059540
3243 A	6 021XA1FC	M.musculus mRNA for mSUG1 {IMAGE:480685}	480685 IMAGE Consortium	AA064154
3244 A	8 021XA1FC	IMAGE EST {IMAGE:482886}	482886 IMAGE Consortium	AA062144
3245 A	10 021XA1FC	EST {IMAGE:482801}	482801 IMAGE Consortium	AA062192
3246 A	12 021XA1FC	ESTs, Highly similar to MATERNAL PUMILIO PROTEIN [Drosophila melanogaster] {IMAGE:481054}	481054 IMAGE Consortium	AA060266
3247 C	2 021XA1FC	ESTs, Moderately similar to probable ataxia-telangiectasia group D protein [H.sapiens] {IMAGE:481011}	481011 IMAGE Consortium	AA059568
3248 C	4 021XA1FC	ESTs, Weakly similar to ZINC FINGER PROTEIN 91 [Homo sapiens] {IMAGE:480805}	480805 IMAGE Consortium	AA059554
3249 C	6 021XA1FC	IMAGE EST {IMAGE:480767}	480767 IMAGE Consortium	AA064098
3250 C	8 021XA1FC	ESTs, Highly similar to CD34 {IMAGE:481134}	481134 IMAGE Consortium	AA064307
3251 C	10 021XA1FC	IMAGE EST {IMAGE:481116}	481116 IMAGE Consortium	AA064296
3252 C	12 021XA1FC	ESTs, Highly similar to similar to human ZFY protein. [H.sapiens] {IMAGE:481738}	481738 IMAGE Consortium	AA060063
3253 E	2 021XA1FC	IMAGE EST {IMAGE:481297}	481297 IMAGE Consortium	AA060444
3254 E	4 021XA1FC	ESTs {IMAGE:481098}	481098 IMAGE Consortium	AA060321
3255 E	6 021XA1FC	ESTs, Weakly similar to T13H5.5 [C.elegans] {IMAGE:481469}	481469 IMAGE Consortium	AA060565
3256 E	8 021XA1FC	EST {IMAGE:481499}	481499 IMAGE Consortium	AA060615
3257 E	10 021XA1FC	IMAGE EST {IMAGE:481485}	481485 IMAGE Consortium	AA060308
3258 E	12 021XA1FC	IMAGE EST {IMAGE:482138}	482138 IMAGE Consortium	AA059643
3259 G	2 021XA1FC	IMAGE EST {IMAGE:482870}	482870 IMAGE Consortium	AA062137
3260 G	4 021XA1FC	ESTs, Highly similar to URIDYLATE KINASE [Saccharomyces cerevisiae] {IMAGE:482658}	482658 IMAGE Consortium	AA062050
3261 G	6 021XA1FC	IMAGE EST {IMAGE:483148}	483148 IMAGE Consortium	AA060909
3262 G	8 021XA1FC	EST {IMAGE:481542}	481542 IMAGE Consortium	AA060599
3263 G	10 021XA1FC	Cytochrome P450, 4a14 {IMAGE:481523}	481523 IMAGE Consortium	AA060595
3264 G	12 021XA1FC	IMAGE EST {IMAGE:481629}	481629 IMAGE Consortium	AA060256
3265 A	2 0214A1FD	IMAGE EST {IMAGE:480662}	480662 IMAGE Consortium	AA064128
3266 A	4 0214A1FD	Cytochrome P450, 2e1, ethanol inducible {IMAGE:481322}	481322 IMAGE Consortium	AA060334
3267 A	6 0214A1FD	EST {IMAGE:481308}	481308 IMAGE Consortium	AA060443
3268 A	8 0214A1FD	IMAGE EST {IMAGE:483196}	483196 IMAGE Consortium	AA060868
3269 A	10 0214A1FD	Mus musculus Ste-20 related kinase SPAK mRNA, complete cds {IMAGE:481691}	481691 IMAGE Consortium	AA060240

3270 A	12 0214A1FD	ESTs {IMAGE:481647}	481647 IMAGE Consortium	AA060228
3271 C	2 0214A1FD	Fas-associated factor 1 {IMAGE:481242}	481242 IMAGE Consortium	AA060511
3272 C	4 0214A1FD	ESTs {IMAGE:483014}	483014 IMAGE Consortium	AA060971
3273 C	6 0214A1FD	ESTs {IMAGE:483465}	483465 IMAGE Consortium	AA061732
3274 C	8 0214A1FD	ESTs, Weakly similar to UBIQUITIN CARBOXYL-TERMINAL HYDROLASE 13 [Saccharomyces cerevisiae] {IMAGE:483589}	483589 IMAGE Consortium	AA061907
3275 C	10 0214A1FD	ESTs, Weakly similar to FOUR-JOINTED PROTEIN [D.melanogaster] {IMAGE:482170}	482170 IMAGE Consortium	AA059698
3276 C	12 0214A1FD	ESTs, Highly similar to NEDD-4 PROTEIN [Homo sapiens] {IMAGE:482030}	482030 IMAGE Consortium	AA059891
3277 E	2 0214A1FD	EST {IMAGE:483016}	483016 IMAGE Consortium	AA060865
3278 E	4 0214A1FD	ESTs {IMAGE:481500}	481500 IMAGE Consortium	AA060591
3279 E	6 0214A1FD	IMAGE EST {IMAGE:481576}	481576 IMAGE Consortium	AA060099
3280 E	8 0214A1FD	IMAGE EST {IMAGE:481630}	481630 IMAGE Consortium	AA060255
3281 E	10 0214A1FD	Sialyltransferase 1 {IMAGE:481883}	481883 IMAGE Consortium	AA059687
3282 E	12 0214A1FD	ESTs, Moderately similar to SAV PROTEIN [Sulfolobus acidocaldarius] {IMAGE:484183}	484183 IMAGE Consortium	AA080005
3283 G	2 0214A1FD	IMAGE EST {IMAGE:570624}	570624 IMAGE Consortium	AA108622
3284 G	4 0214A1FD	IMAGE EST {IMAGE:570616}	570616 IMAGE Consortium	AA108591
3285 G	6 0214A1FD	IMAGE EST {IMAGE:573324}	573324 IMAGE Consortium	AA118718
3286 G	8 0214A1FD	ESTs {IMAGE:571300}	571300 IMAGE Consortium	AA108832
3287 G	10 0214A1FD	Prostaglandin D2 synthase (21 kDa, brain) {IMAGE:571621}	571621 IMAGE Consortium	AA108857
3288 G	12 0214A1FD	IMAGE EST {IMAGE:492526}	492526 IMAGE Consortium	AA097228
3289 A	2 021BA1FE	Mus musculus mRNA for nuclear protein SA3 {IMAGE:493305}	493305 IMAGE Consortium	AA097991
3290 A	4 021BA1FE	IMAGE EST {IMAGE:493024}	493024 IMAGE Consortium	AA098433
3291 A	6 021BA1FE	IMAGE EST {IMAGE:572919}	572919 IMAGE Consortium	AA111600
3292 A	8 021BA1FE	ESTs {IMAGE:570513}	570513 IMAGE Consortium	AA108514
3293 A	10 021BA1FE	IMAGE EST {IMAGE:573489}	573489 IMAGE Consortium	AA119260
3294 A	12 021BA1FE	Glutaryl Coenzyme A dehydrogenase {IMAGE:571328}	571328 IMAGE Consortium	AA108738
3295 C	2 021BA1FE	ESTs {IMAGE:493582}	493582 IMAGE Consortium	AA087131
3296 C	4 021BA1FE	ESTs, Highly similar to SEC PROTEIN [Homo sapiens] {IMAGE:570540}	570540 IMAGE Consortium	AA108520
3297 C	6 021BA1FE	ESTs, Weakly similar to putative Rab5-interacting protein {clone L1-94} [H.sapiens] {IMAGE:570898}	570898 IMAGE Consortium	AA110278
3298 C	8 021BA1FE	ESTs, Weakly similar to NIFS PROTEIN [Bacillus subtilis] {IMAGE:571199}	571199 IMAGE Consortium	AA107712
3299 C	10 021BA1FE	ESTs, Weakly similar to similar to DNA-J like domain [C.elegans] {IMAGE:571604}	571604 IMAGE Consortium	AA109705
3300 C	12 021BA1FE	IMAGE EST {IMAGE:493301}	493301 IMAGE Consortium	AA098079
3301 E	2 021BA1FE	IMAGE EST {IMAGE:573460}	573460 IMAGE Consortium	AA118783
3302 E	4 021BA1FE	Requiem {IMAGE:573346}	573346 IMAGE Consortium	AA118721

3303 E	6 021BA1FE	ESTs {IMAGE:492441}	492441 IMAGE Consortium	AA097191
3304 E	8 021BA1FE	IMAGE EST {IMAGE:493163}	493163 IMAGE Consortium	AA097783
3305 E	10 021BA1FE	ESTs, Highly similar to NADH-UBIQUINONE OXIDOREDUCTASE 49 KD SUBUNIT [Bos taurus] {IMAGE:572585}	572585 IMAGE Consortium	AA109715
3306 E	12 021BA1FE	IMAGE EST {IMAGE:493644}	493644 IMAGE Consortium	AA098237
3307 G	2 021BA1FE	IMAGE EST {IMAGE:572070}	572070 IMAGE Consortium	AA108466
3308 G	4 021BA1FE	ESTs, Highly similar to TRANSCRIPTIONAL REGULATORY PROTEIN RPD3 [Saccharomyces cerevisiae] {IMAGE:572157}	572157 IMAGE Consortium	AA105228
3309 G	6 021BA1FE	ATPase, Na <sup>+</sup> /K <sup>+</sup> transporting, beta 1 polypeptide {IMAGE:493488}	493488 IMAGE Consortium	AA087098
3310 G	8 021BA1FE	IMAGE EST {IMAGE:572889}	572889 IMAGE Consortium	AA110811
3311 G	10 021BA1FE	IMAGE EST {IMAGE:572824}	572824 IMAGE Consortium	AA110266
3312 G	12 021BA1FE	Mus musculus E2F-like transcriptional repressor protein mRNA, complete cds {IMAGE:570685}	570685 IMAGE Consortium	AA108727
3313 A	2 021IA1FF	IMAGE EST {IMAGE:484234}	484234 IMAGE Consortium	AA080220
3314 A	4 021IA1FF	ESTs {IMAGE:571310}	571310 IMAGE Consortium	AA108797
3315 A	6 021IA1FF	IMAGE EST {IMAGE:492446}	492446 IMAGE Consortium	AA097190
3316 A	8 021IA1FF	Periplakin {IMAGE:571984}	571984 IMAGE Consortium	AA105152
3317 A	10 021IA1FF	ESTs {IMAGE:572149}	572149 IMAGE Consortium	AA110042
3318 A	12 021IA1FF	Cytochrome P450, 4a14 {IMAGE:493244}	493244 IMAGE Consortium	AA098524
3319 C	2 021IA1FF	IMAGE EST {IMAGE:492457}	492457 IMAGE Consortium	AA097499
3320 C	4 021IA1FF	ESTs {IMAGE:492617}	492617 IMAGE Consortium	AA097261
3321 C	6 021IA1FF	IMAGE EST {IMAGE:572376}	572376 IMAGE Consortium	AA108396
3322 C	8 021IA1FF	IMAGE EST {IMAGE:492998}	492998 IMAGE Consortium	AA097520
3323 C	10 021IA1FF	ESTs, Highly similar to E1B 19K/Bcl-2-interacting protein Nip1 [H.sapiens] {IMAGE:572773}	572773 IMAGE Consortium	AA111758
3324 C	12 021IA1FF	Beta-2 microglobulin {IMAGE:572542}	572542 IMAGE Consortium	AA109951
3325 E	2 021IA1FF	Mus musculus cdc2/CDC28-like kinase 2 (Cdk2) mRNA, complete cds {IMAGE:492518}	492518 IMAGE Consortium	AA097860
3326 E	4 021IA1FF	ESTs {IMAGE:572041}	572041 IMAGE Consortium	AA108460
3327 E	6 021IA1FF	ESTs, Highly similar to UBIQUINOL-CYTOCHROME C REDUCTASE COMPLEX CORE PROTEIN 2 PRECURSOR [Bos taurus] {IMAGE:572127}	572127 IMAGE Consortium	AA108590
3328 E	8 021IA1FF	ESTs, Highly similar to HYPOTHETICAL 13.5 KD PROTEIN C45G9.7 IN CHROMOSOME III [Caenorhabditis elegans] {IMAGE:572510}	572510 IMAGE Consortium	AA111643
3329 E	10 021IA1FF	IMAGE EST {IMAGE:493687}	493687 IMAGE Consortium	AA087206
3330 E	12 021IA1FF	60S RIBOSOMAL PROTEIN L3 {IMAGE:570533}	570533 IMAGE Consortium	AA108363
3331 G	2 021IA1FF	ESTs, Weakly similar to putative enzyme [E.coli] {IMAGE:572151}	572151 IMAGE Consortium	AA110154
3332 G	4 021IA1FF	Cyclin D1 {IMAGE:572428}	572428 IMAGE Consortium	AA111722
3333 G	6 021IA1FF	Lipocalin 2 {IMAGE:493658}	493658 IMAGE Consortium	AA087193

3334 G	8 021IA1FF	IMAGE EST {IMAGE:493449}	493449 IMAGE Consortium	AA087037
3335 G	10 021IA1FF	ESTs, Weakly similar to ZK1058.5 [C.elegans] {IMAGE:573062}	573062 IMAGE Consortium	AA118419
3336 G	12 021IA1FF	IMAGE EST {IMAGE:572901}	572901 IMAGE Consortium	AA110353
3337 A	2 021PA1FG	ESTs, Highly similar to GLYCOPROTEIN 25L PRECURSOR [Canis familiaris] {IMAGE:572466}	572466 IMAGE Consortium	AA109932
3338 A	4 021PA1FG	IMAGE EST {IMAGE:572890}	572890 IMAGE Consortium	AA110808
3339 A	6 021PA1FG	ESTs {IMAGE:573432}	573432 IMAGE Consortium	AA118755
3340 A	8 021PA1FG	IMAGE EST {IMAGE:570997}	570997 IMAGE Consortium	AA108438
3341 A	10 021PA1FG	ESTs, Highly similar to TRICARBOXYLATE TRANSPORT PROTEIN PRECURSOR [Rattus norvegicus] {IMAGE:571633}	571633 IMAGE Consortium	AA108822
3342 A	12 021PA1FG	Mus musculus transcription factor BFCOL1 mRNA, complete cds {IMAGE:575405}	575405 IMAGE Consortium	AA125037
3343 C	2 021PA1FG	Plasminogen activator inhibitor, type II {IMAGE:574835}	574835 IMAGE Consortium	AA124868
3344 C	4 021PA1FG	ESTs {IMAGE:577019}	577019 IMAGE Consortium	AA124354
3345 C	6 021PA1FG	IMAGE EST {IMAGE:573843}	573843 IMAGE Consortium	AA119066
3346 C	8 021PA1FG	Cytochrome P450, 2e1, ethanol inducible {IMAGE:573803}	573803 IMAGE Consortium	AA118977
3347 C	10 021PA1FG	Tax-transcriptionally activated glycoprotein 1 {IMAGE:573690}	573690 IMAGE Consortium	AA110765
3348 C	12 021PA1FG	Mus domesticus ZNF127 (zfn127) gene, complete cds {IMAGE:574274}	574274 IMAGE Consortium	AA119257
3349 E	2 021PA1FG	IMAGE EST {IMAGE:576821}	576821 IMAGE Consortium	AA123850
3350 E	4 021PA1FG	ESTs {IMAGE:575087}	575087 IMAGE Consortium	AA120679
3351 E	6 021PA1FG	ESTs {IMAGE:575264}	575264 IMAGE Consortium	AA125391
3352 E	8 021PA1FG	ESTs {IMAGE:573851}	573851 IMAGE Consortium	AA119050
3353 E	10 021PA1FG	IMAGE EST {IMAGE:573906}	573906 IMAGE Consortium	AA111038
3354 E	12 021PA1FG	IMAGE EST {IMAGE:574227}	574227 IMAGE Consortium	AA119136
3355 G	2 021PA1FG	IMAGE EST {IMAGE:574914}	574914 IMAGE Consortium	AA120631
3356 G	4 021PA1FG	IMAGE EST {IMAGE:575336}	575336 IMAGE Consortium	AA125134
3357 G	6 021PA1FG	ESTs, Highly similar to H-2 class I histocompatibility antigen T15 [M.musculus] {IMAGE:575463}	575463 IMAGE Consortium	AA124667
3358 G	8 021PA1FG	IMAGE EST {IMAGE:575181}	575181 IMAGE Consortium	AA119154
3359 G	10 021PA1FG	ESTs, Highly similar to UBIQUITIN-ACTIVATING ENZYME E1 HOMOLOG [Homo sapiens] {IMAGE:575521}	575521 IMAGE Consortium	AA125011
3360 G	12 021PA1FG	ESTs, Weakly similar to ORF YLL039c [S.cerevisiae] {IMAGE:575698}	575698 IMAGE Consortium	AA123007
3361 A	2 021WA1FH	ESTs {IMAGE:575106}	575106 IMAGE Consortium	AA120681
3362 A	4 021WA1FH	IMAGE EST {IMAGE:576764}	576764 IMAGE Consortium	AA120011
3363 A	6 021WA1FH	IMAGE EST {IMAGE:573845}	573845 IMAGE Consortium	AA119067
3364 A	8 021WA1FH	Small inducible cytokine A6 {IMAGE:575397}	575397 IMAGE Consortium	AA119293
3365 A	10 021WA1FH	ESTs {IMAGE:575782}	575782 IMAGE Consortium	AA120432
3366 A	12 021WA1FH	ESTs, Weakly similar to coded for by C. elegans cDNA yk34a10.3 [C.elegans] {IMAGE:575965}	575965 IMAGE Consortium	AA119999

3367 C	2	021WA1FH	IMAGE EST {IMAGE:575330}	575330	IMAGE Consortium	AA123962
3368 C	4	021WA1FH	ESTs {IMAGE:573526}	573526	IMAGE Consortium	AA110654
3369 C	6	021WA1FH	IMAGE EST {IMAGE:573582}	573582	IMAGE Consortium	AA119372
3370 C	8	021WA1FH	IMAGE EST {IMAGE:573907}	573907	IMAGE Consortium	AA111040
3371 C	10	021WA1FH	ESTs, Moderately similar to serine/threonine protein kinase [M.musculus] {IMAGE:573966}	573966	IMAGE Consortium	AA118903
3372 C	12	021WA1FH	IMAGE EST {IMAGE:573973}	573973	IMAGE Consortium	AA118960
3373 E	2	021WA1FH	Histocompatibility 2, D region locus 1 {IMAGE:574771}	574771	IMAGE Consortium	AA120637
3374 E	4	021WA1FH	ESTs {IMAGE:575187}	575187	IMAGE Consortium	AA124918
3375 E	6	021WA1FH	IMAGE EST {IMAGE:575290}	575290	IMAGE Consortium	AA125399
3376 E	8	021WA1FH	IMAGE EST {IMAGE:575344}	575344	IMAGE Consortium	AA119275
3377 E	10	021WA1FH	ESTs, Weakly similar to similar to yeast hypothetical 51.3 kd protein in SMY2-RPS101 region [C.elegans] {IMAGE:573596}	573596	IMAGE Consortium	AA111707
3378 E	12	021WA1FH	ESTs {IMAGE:575605}	575605	IMAGE Consortium	AA124548
3379 G	2	021WA1FH	M.musculus mRNA for cell surface glycoprotein CD53 {IMAGE:576862}	576862	IMAGE Consortium	AA116505
3380 G	4	021WA1FH	IMAGE EST {IMAGE:575434}	575434	IMAGE Consortium	AA124928
3381 G	6	021WA1FH	ESTs, Highly similar to PROBABLE PHOSPHOSERINE AMINOTRANSFERASE [Oryctolagus cuniculus] {IMAGE:573784}	573784	IMAGE Consortium	AA110896
3382 G	8	021WA1FH	Rat generating islet-derived, mouse homolog 3 gamma {IMAGE:575773}	575773	IMAGE Consortium	AA123026
3383 G	10	021WA1FH	ESTs {IMAGE:575930}	575930	IMAGE Consortium	AA119941
3384 G	12	021WA1FH	ESTs, Highly similar to H-2 class I histocompatibility antigen T15 [M.musculus] {IMAGE:575960}	575960	IMAGE Consortium	AA119993
3385 A	2	0213A1FI	Mus musculus clone L5 uniform group of 2-cell-stage gene family mRNA, complete cds {IMAGE:582531}	582531	IMAGE Consortium	AA153114
3386 A	4	0213A1FI	IMAGE EST {IMAGE:582588}	582588	IMAGE Consortium	AA153968
3387 A	6	0213A1FI	IMAGE EST {IMAGE:582716}	582716	IMAGE Consortium	AA142459
3388 A	8	0213A1FI	ESTs, Highly similar to SUCCINATE DEHYDROGENASE CYTOCHROME B560 SUBUNIT PRECURSOR [Bos taurus] {IMAGE:580040}	580040	IMAGE Consortium	AA137762
3389 A	10	0213A1FI	ESTs {IMAGE:577155}	577155	IMAGE Consortium	AA124527
3390 A	12	0213A1FI	ESTs, Highly similar to NEUROLYSIN PRECURSOR [Rattus norvegicus] {IMAGE:577260}	577260	IMAGE Consortium	AA124221
3391 C	2	0213A1FI	IMAGE EST {IMAGE:577046}	577046	IMAGE Consortium	AA124423
3392 C	4	0213A1FI	IMAGE EST {IMAGE:577199}	577199	IMAGE Consortium	AA116515
3393 C	6	0213A1FI	IMAGE EST {IMAGE:577735}	577735	IMAGE Consortium	AA140352
3394 C	8	0213A1FI	IMAGE EST {IMAGE:581177}	581177	IMAGE Consortium	AA138559
3395 C	10	0213A1FI	ESTs {IMAGE:581652}	581652	IMAGE Consortium	AA152852
3396 C	12	0213A1FI	IMAGE EST {IMAGE:581680}	581680	IMAGE Consortium	AA146165
3397 E	2	0213A1FI	ATP synthase, H+ transporting, mitochondrial F0 complex, subunit c (subunit 9), isoform 1 {IMAGE:580898}	580898	IMAGE Consortium	AA139793
3398 E	4	0213A1FI	ESTs, Highly similar to PROFILIN II [Homo sapiens]	581193	IMAGE Consortium	AA139628

3399 E	6 0213A1FI	{IMAGE:581193} ESTs, Weakly similar to Des-1 protein [D.melanogaster] {IMAGE:577698}	577698 IMAGE Consortium	AA140530
3400 E	8 0213A1FI	ESTs, Highly similar to proteasome subunit p112 [H.sapiens] {IMAGE:581724}	581724 IMAGE Consortium	AA152739
3401 E	10 0213A1FI	ESTs {IMAGE:582695}	582695 IMAGE Consortium	AA144003
3402 E	12 0213A1FI	ESTs, Moderately similar to BIS(5'-NUCLEOSYL)- TETRAPHOSPHATASE [H.sapiens] {IMAGE:580384}	580384 IMAGE Consortium	AA123895
3403 G	2 0213A1FI	IMAGE EST {IMAGE:582719}	582719 IMAGE Consortium	AA142464
3404 G	4 0213A1FI	IMAGE EST {IMAGE:580019}	580019 IMAGE Consortium	AA137760
3405 G	6 0213A1FI	Mus musculus chromatin structural protein homolog Supt5hp (Supt5h) mRNA, complete cds {IMAGE:580108}	580108 IMAGE Consortium	AA137833
3406 G	8 0213A1FI	ESTs, Weakly similar to C08F8.2 [C.elegans] {IMAGE:577312}	577312 IMAGE Consortium	AA116572
3407 G	10 0213A1FI	Phenylalanine hydroxylase {IMAGE:581101}	581101 IMAGE Consortium	AA140114
3408 G	12 0213A1FI	IMAGE EST {IMAGE:581087}	581087 IMAGE Consortium	AA138659
3409 A	2 021AA1FJ	IMAGE EST {IMAGE:575244}	575244 IMAGE Consortium	AA125385
3410 A	4 021AA1FJ	ESTs {IMAGE:575441}	575441 IMAGE Consortium	AA119512
3411 A	6 021AA1FJ	IMAGE EST {IMAGE:573813}	573813 IMAGE Consortium	AA111051
3412 A	8 021AA1FJ	IMAGE EST {IMAGE:575606}	575606 IMAGE Consortium	AA124545
3413 A	10 021AA1FJ	IMAGE EST {IMAGE:575984}	575984 IMAGE Consortium	AA120279
3414 A	12 021AA1FJ	ESTs, Highly similar to PROBABLE UBIQUITIN CARBOXYL- TERMINAL HYDROLASE [Homo sapiens] {IMAGE:574425}	574425 IMAGE Consortium	AA119665
3415 C	2 021AA1FJ	ESTs {IMAGE:575436}	575436 IMAGE Consortium	AA124929
3416 C	4 021AA1FJ	IMAGE EST {IMAGE:573832}	573832 IMAGE Consortium	AA111053
3417 C	6 021AA1FJ	ESTs {IMAGE:573743}	573743 IMAGE Consortium	AA110910
3418 C	8 021AA1FJ	IMAGE EST {IMAGE:574309}	574309 IMAGE Consortium	AA117617
3419 C	10 021AA1FJ	Mus musculus TIA mRNA, complete cds {IMAGE:575920}	575920 IMAGE Consortium	AA117605
3420 C	12 021AA1FJ	IMAGE EST {IMAGE:576319}	576319 IMAGE Consortium	AA120086
3421 E	2 021AA1FJ	Mus musculus mRNA for putative tripeptidyl peptidase I {IMAGE:574070}	574070 IMAGE Consortium	AA119072
3422 E	4 021AA1FJ	ESTs, Highly similar to GCN20 PROTEIN [Saccharomyces cerevisiae] {IMAGE:574680}	574680 IMAGE Consortium	AA125260
3423 E	6 021AA1FJ	Mus musculus ubiquitin conjugating enzyme UBC9 mRNA, complete cds {IMAGE:576391}	576391 IMAGE Consortium	AA120351
3424 E	8 021AA1FJ	IMAGE EST {IMAGE:574921}	574921 IMAGE Consortium	AA124979
3425 E	10 021AA1FJ	IMAGE EST {IMAGE:576987}	576987 IMAGE Consortium	AA122542
3426 E	12 021AA1FJ	Fatty acid synthase {IMAGE:576881}	576881 IMAGE Consortium	AA116513
3427 G	2 021AA1FJ	IMAGE EST {IMAGE:574710}	574710 IMAGE Consortium	AA125271
3428 G	4 021AA1FJ	ESTs {IMAGE:576535}	576535 IMAGE Consortium	AA123823
3429 G	6 021AA1FJ	IMAGE EST {IMAGE:576518}	576518 IMAGE Consortium	AA123917
3430 G	8 021AA1FJ	IMAGE EST {IMAGE:576405}	576405 IMAGE Consortium	AA119678
3431 G	10 021AA1FJ	Mus musculus mRNA for neutral sphingomyelinase	576652 IMAGE Consortium	AA123762

		{IMAGE:576652}		
3432 G	12 021AA1FJ	IMAGE EST {IMAGE:576850}	576850 IMAGE Consortium	AA123455
3433 A	2 021HA1FK	ESTs, Highly similar to GTPASE ACTIVATING PROTEIN ROTUND [Drosophila melanogaster] {IMAGE:577676}	577676 IMAGE Consortium	AA140523
3434 A	4 021HA1FK	Mus musculus RW1 protein mRNA, complete cds {IMAGE:578120}	578120 IMAGE Consortium	AA124746
3435 A	6 021HA1FK	ESTs, Highly similar to The KIAA0130 gene is related to mouse genetic suppressor element [H.sapiens] {IMAGE:577821}	577821 IMAGE Consortium	AA116223
3436 A	8 021HA1FK	IMAGE EST {IMAGE:578380}	578380 IMAGE Consortium	AA124092
3437 A	10 021HA1FK	ESTs {IMAGE:578270}	578270 IMAGE Consortium	AA122636
3438 A	12 021HA1FK	Mus musculus mRNA for mGpi1p, complete cds {IMAGE:578206}	578206 IMAGE Consortium	AA120110
3439 C	2 021HA1FK	IMAGE EST {IMAGE:581965}	581965 IMAGE Consortium	AA154133
3440 C	4 021HA1FK	ESTs {IMAGE:581711}	581711 IMAGE Consortium	AA146177
3441 C	6 021HA1FK	ESTs {IMAGE:582475}	582475 IMAGE Consortium	AA143968
3442 C	8 021HA1FK	IMAGE EST {IMAGE:577135}	577135 IMAGE Consortium	AA124522
3443 C	10 021HA1FK	Mus musculus mRNA for mouse rabaptin-5, complete cds {IMAGE:580579}	580579 IMAGE Consortium	AA137859
3444 C	12 021HA1FK	ESTs {IMAGE:577737}	577737 IMAGE Consortium	AA139222
3445 E	2 021HA1FK	ESTs {IMAGE:582499}	582499 IMAGE Consortium	AA144080
3446 E	4 021HA1FK	IMAGE EST {IMAGE:583008}	583008 IMAGE Consortium	AA153697
3447 E	6 021HA1FK	ESTs, Weakly similar to HYPOTHETICAL 11.4 KD PROTEIN C13G6.04 IN CHROMOSOME I [Schizosaccharomyces pombe] {IMAGE:579733}	579733 IMAGE Consortium	AA116946
3448 E	8 021HA1FK	GA repeat binding protein, beta 1 {IMAGE:577184}	577184 IMAGE Consortium	AA124181
3449 E	10 021HA1FK	Small inducible cytokine A11 {IMAGE:577422}	577422 IMAGE Consortium	AA139170
3450 E	12 021HA1FK	ESTs {IMAGE:580935}	580935 IMAGE Consortium	AA139918
3451 G	2 021HA1FK	ESTs {IMAGE:581007}	581007 IMAGE Consortium	AA139837
3452 G	4 021HA1FK	IMAGE EST {IMAGE:581799}	581799 IMAGE Consortium	AA153905
3453 G	6 021HA1FK	IMAGE EST {IMAGE:578240}	578240 IMAGE Consortium	AA122593
3454 G	8 021HA1FK	Mus musculus peroxisome proliferator-activated receptor gamma binding protein mRNA, complete cds {IMAGE:582634}	582634 IMAGE Consortium	AA152649
3455 G	10 021HA1FK	ESTs, Weakly similar to PUTATIVE ACID PHOSPHATASE F26C11.1 [Caenorhabditis elegans] {IMAGE:582668}	582668 IMAGE Consortium	AA144194
3456 G	12 021HA1FK	ESTs {IMAGE:583033}	583033 IMAGE Consortium	AA153663
3457 A	2 021OA1FL	IMAGE EST {IMAGE:583495}	583495 IMAGE Consortium	AA142584
3458 A	4 021OA1FL	Spleen tyrosine kinase {IMAGE:583475}	583475 IMAGE Consortium	AA142676
3459 A	6 021OA1FL	IMAGE EST {IMAGE:583657}	583657 IMAGE Consortium	AA144094
3460 A	8 021OA1FL	IMAGE EST {IMAGE:583783}	583783 IMAGE Consortium	AA152817
3461 A	10 021OA1FL	IMAGE EST {IMAGE:583571}	583571 IMAGE Consortium	AA144047
3462 A	12 021OA1FL	IMAGE EST {IMAGE:583771}	583771 IMAGE Consortium	AA146402

3463 C	2 021OA1FL	ESTs {IMAGE:596893}	596893 IMAGE Consortium	AA145237
3464 C	4 021OA1FL	Myxovirus (influenza virus) resistance 2 {IMAGE:599128}	599128 IMAGE Consortium	AA145795
3465 C	6 021OA1FL	ESTs {IMAGE:597083}	597083 IMAGE Consortium	AA145564
3466 C	8 021OA1FL	IMAGE EST {IMAGE:597190}	597190 IMAGE Consortium	AA139085
3467 C	10 021OA1FL	IMAGE EST {IMAGE:597489}	597489 IMAGE Consortium	AA154417
3468 C	12 021OA1FL	Cytochrome P450, 2e1, ethanol inducible {IMAGE:597544}	597544 IMAGE Consortium	AA144383
3469 E	2 021OA1FL	IMAGE EST {IMAGE:598898}	598898 IMAGE Consortium	AA168504
3470 E	4 021OA1FL	IMAGE EST {IMAGE:598837}	598837 IMAGE Consortium	AA168538
3471 E	6 021OA1FL	ESTs {IMAGE:598870}	598870 IMAGE Consortium	AA162785
3472 E	8 021OA1FL	Cytochrome P450, 2e1, ethanol inducible {IMAGE:599296}	599296 IMAGE Consortium	AA145856
3473 E	10 021OA1FL	IMAGE EST {IMAGE:583238}	583238 IMAGE Consortium	AA154812
3474 E	12 021OA1FL	IMAGE EST {IMAGE:597513}	597513 IMAGE Consortium	AA154277
3475 G	2 021OA1FL	Promyelocytic leukemia {IMAGE:596564}	596564 IMAGE Consortium	AA138191
3476 G	4 021OA1FL	IMAGE EST {IMAGE:598894}	598894 IMAGE Consortium	AA162792
3477 G	6 021OA1FL	ESTs, Weakly similar to UBIQUITIN-CONJUGATING ENZYME E2-17 KD 1 [H.sapiens] {IMAGE:599190}	599190 IMAGE Consortium	AA154392
3478 G	8 021OA1FL	CD53 antigen {IMAGE:598986}	598986 IMAGE Consortium	AA162875
3479 G	10 021OA1FL	IMAGE EST {IMAGE:616807}	616807 IMAGE Consortium	AA174496
3480 G	12 021OA1FL	ESTs {IMAGE:583526}	583526 IMAGE Consortium	AA142687
3481 A	2 021VA1FM	Abl-interactor 1 {IMAGE:598116}	598116 IMAGE Consortium	AA266646
3482 A	4 021VA1FM	ESTs {IMAGE:598278}	598278 IMAGE Consortium	AA162654
3483 A	6 021VA1FM	IMAGE EST {IMAGE:598625}	598625 IMAGE Consortium	AA163165
3484 A	8 021VA1FM	Long (electrocardiographic) QT syndrome 2 {IMAGE:596823}	596823 IMAGE Consortium	AA145089
3485 A	10 021VA1FM	EST {IMAGE:599303}	599303 IMAGE Consortium	AA145901
3486 A	12 021VA1FM	IMAGE EST {IMAGE:597115}	597115 IMAGE Consortium	AA139065
3487 C	2 021VA1FM	IMAGE EST {IMAGE:598824}	598824 IMAGE Consortium	AA168416
3488 C	4 021VA1FM	Histocompatibility 2, Q region locus 7 {IMAGE:596447}	596447 IMAGE Consortium	AA122791
3489 C	6 021VA1FM	Caveolin, caveolae protein, 22 kDa {IMAGE:596968}	596968 IMAGE Consortium	AA138693
3490 C	8 021VA1FM	ESTs {IMAGE:597336}	597336 IMAGE Consortium	AA146020
3491 C	10 021VA1FM	IMAGE EST {IMAGE:616834}	616834 IMAGE Consortium	AA174484
3492 C	12 021VA1FM	ESTs {IMAGE:597020}	597020 IMAGE Consortium	AA138757
3493 E	2 021VA1FM	IMAGE EST {IMAGE:598537}	598537 IMAGE Consortium	AA162559
3494 E	4 021VA1FM	ESTs {IMAGE:596543}	596543 IMAGE Consortium	AA138313
3495 E	6 021VA1FM	Mouse thymidylate kinase homologue mRNA, complete cds {IMAGE:599164}	599164 IMAGE Consortium	AA154388
3496 E	8 021VA1FM	SH3 domain protein 2C {IMAGE:597291}	597291 IMAGE Consortium	AA146010
3497 E	10 021VA1FM	ESTs {IMAGE:597249}	597249 IMAGE Consortium	AA145337
3498 E	12 021VA1FM	IMAGE EST {IMAGE:597415}	597415 IMAGE Consortium	AA152882
3499 G	2 021VA1FM	IMAGE EST {IMAGE:598873}	598873 IMAGE Consortium	AA168500
3500 G	4 021VA1FM	ESTs {IMAGE:598698}	598698 IMAGE Consortium	AA163457
3501 G	6 021VA1FM	ESTs, Weakly similar to HYPOTHETICAL 24.1 KD PROTEIN ON CHROMOSOME 1 [M.musculus] {IMAGE:596825}	596825 IMAGE Consortium	AA138563
3502 G	8 021VA1FM	IMAGE EST {IMAGE:597321}	597321 IMAGE Consortium	AA146110



3503 G	10 021VA1FM	IMAGE EST {IMAGE:597357}	597357 IMAGE Consortium	AA138526
3504 G	12 021VA1FM	IMAGE EST {IMAGE:597679}	597679 IMAGE Consortium	AA153223
3505 A	2 0212A1FN	ESTs {IMAGE:596340}	596340 IMAGE Consortium	AA138854
3506 A	4 0212A1FN	Mouse mkf-1 mRNA, complete cds {IMAGE:596277}	596277 IMAGE Consortium	AA122720
3507 A	6 0212A1FN	ESTs, Moderately similar to HYPOTHETICAL 92.1 KD PROTEIN C24H6.03 IN CHROMOSOME I [Schizosaccharomyces pombe] {IMAGE:598918}	598918 IMAGE Consortium	AA162800
3508 A	8 0212A1FN	IMAGE EST {IMAGE:596708}	596708 IMAGE Consortium	AA138394
3509 A	10 0212A1FN	IMAGE EST {IMAGE:597342}	597342 IMAGE Consortium	AA146022
3510 A	12 0212A1FN	ESTs, Weakly similar to GLUCOSYLTRANSFERASE ALG8 [Saccharomyces cerevisiae] {IMAGE:583448}	583448 IMAGE Consortium	AA142817
3511 C	2 0212A1FN	Mus musculus mRNA for peroxisomal integral membrane protein PMP34 {IMAGE:596035}	596035 IMAGE Consortium	AA122711
3512 C	4 0212A1FN	ESTs {IMAGE:596570}	596570 IMAGE Consortium	AA146460
3513 C	6 0212A1FN	Aryl-hydrocarbon receptor {IMAGE:598827}	598827 IMAGE Consortium	AA168468
3514 C	8 0212A1FN	ESTs, Moderately similar to BIOTIN CARBOXYLASE [Anabaena pcc7120] {IMAGE:598743}	598743 IMAGE Consortium	AA168437
3515 C	10 0212A1FN	ESTs {IMAGE:598929}	598929 IMAGE Consortium	AA168640
3516 C	12 0212A1FN	ESTs {IMAGE:597031}	597031 IMAGE Consortium	AA145466
3517 E	2 0212A1FN	ESTs, Highly similar to KIAA0103 [H.sapiens] {IMAGE:596144}	596144 IMAGE Consortium	AA137986
3518 E	4 0212A1FN	ESTs {IMAGE:596179}	596179 IMAGE Consortium	AA122756
3519 E	6 0212A1FN	ESTs, Weakly similar to cDNA EST EMBL:C13590 comes from this gene [C.elegans] {IMAGE:596404}	596404 IMAGE Consortium	AA138546
3520 E	8 0212A1FN	IMAGE EST {IMAGE:596556}	596556 IMAGE Consortium	AA146363
3521 E	10 0212A1FN	IMAGE EST {IMAGE:596494}	596494 IMAGE Consortium	AA122801
3522 E	12 0212A1FN	ESTs, Highly similar to NEDD-4 PROTEIN [Homo sapiens] {IMAGE:596719}	596719 IMAGE Consortium	AA146534
3523 G	2 0212A1FN	ESTs, Highly similar to ISOCITRATE DEHYDROGENASE [Bos taurus] {IMAGE:596186}	596186 IMAGE Consortium	AA137848
3524 G	4 0212A1FN	ESTs, Moderately similar to KIAA0365 [H.sapiens] {IMAGE:598206}	598206 IMAGE Consortium	AA161817
3525 G	6 0212A1FN	Casein kinase II, alpha 2, polypeptide {IMAGE:598388}	598388 IMAGE Consortium	AA155019
3526 G	8 0212A1FN	Killer cell lectin-like receptor, subfamily D, member 1 {IMAGE:598616}	598616 IMAGE Consortium	AA161895
3527 G	10 0212A1FN	ESTs {IMAGE:598864}	598864 IMAGE Consortium	AA162782
3528 G	12 0212A1FN	IMAGE EST {IMAGE:598992}	598992 IMAGE Consortium	AA168602
3529 A	2 0219A1FO	IMAGE EST {IMAGE:598682}	598682 IMAGE Consortium	AA163911
3530 A	4 0219A1FO	IMAGE EST {IMAGE:596880}	596880 IMAGE Consortium	AA145137
3531 A	6 0219A1FO	IMAGE EST {IMAGE:596820}	596820 IMAGE Consortium	AA145023
3532 A	8 0219A1FO	ESTs {IMAGE:596897}	596897 IMAGE Consortium	AA138584
3533 A	10 0219A1FO	Mouse fibronectin (FN) mRNA {IMAGE:597005}	597005 IMAGE Consortium	AA145458
3534 A	12 0219A1FO	IMAGE EST {IMAGE:597058}	597058 IMAGE Consortium	AA145512
3535 C	2 0219A1FO	IMAGE EST {IMAGE:618306}	618306 IMAGE Consortium	AA170599

3536 C	4 0219A1FO	ESTs, Highly similar to progesterone receptor-related protein p23 [H.sapiens] {IMAGE:618630}	618630 IMAGE Consortium	AA174980
3537 C	6 0219A1FO	ESTs {IMAGE:620829}	620829 IMAGE Consortium	AA174729
3538 C	8 0219A1FO	IMAGE EST {IMAGE:619103}	619103 IMAGE Consortium	AA170642
3539 C	10 0219A1FO	ESTs {IMAGE:617593}	617593 IMAGE Consortium	AA174753
3540 C	12 0219A1FO	IL2-inducible T-cell kinase {IMAGE:619755}	619755 IMAGE Consortium	AA176042
3541 E	2 0219A1FO	IMAGE EST {IMAGE:618940}	618940 IMAGE Consortium	AA175377
3542 E	4 0219A1FO	ESTs, Weakly similar to unknown protein [R.norvegicus] {IMAGE:617309}	617309 IMAGE Consortium	AA174814
3543 E	6 0219A1FO	IMAGE EST {IMAGE:617270}	617270 IMAGE Consortium	AA174307
3544 E	8 0219A1FO	ATP-binding cassette 8 {IMAGE:619411}	619411 IMAGE Consortium	AA172527
3545 E	10 0219A1FO	ESTs {IMAGE:619708}	619708 IMAGE Consortium	AA175979
3546 E	12 0219A1FO	ESTs {IMAGE:619664}	619664 IMAGE Consortium	AA175990
3547 G	2 0219A1FO	ESTs, Moderately similar to PLASMA-CELL MEMBRANE GLYCOPROTEIN PC-1 [Homo sapiens] {IMAGE:621166}	621166 IMAGE Consortium	AA177363
3548 G	4 0219A1FO	H-2 CLASS II HISTOCOMPATIBILITY ANTIGEN, I-A BETA CHAIN PRECURSOR {IMAGE:617286}	617286 IMAGE Consortium	AA174620
3549 G	6 0219A1FO	ESTs {IMAGE:617294}	617294 IMAGE Consortium	AA174315
3550 G	8 0219A1FO	IMAGE EST {IMAGE:617529}	617529 IMAGE Consortium	AA174721
3551 G	10 0219A1FO	IMAGE EST {IMAGE:617884}	617884 IMAGE Consortium	AA175265
3552 G	12 0219A1FO	IMAGE EST {IMAGE:619845}	619845 IMAGE Consortium	AA172943
3553 A	2 021GA1FP	ESTs {IMAGE:616975}	616975 IMAGE Consortium	AA166336
3554 A	4 021GA1FP	IMAGE EST {IMAGE:617428}	617428 IMAGE Consortium	AA174441
3555 A	6 021GA1FP	IMAGE EST {IMAGE:620024}	620024 IMAGE Consortium	AA178098
3556 A	8 021GA1FP	ESTs, Highly similar to LYSYL-TRNA SYNTHETASE [Cricetulus longicaudatus] {IMAGE:619742}	619742 IMAGE Consortium	AA175815
3557 A	10 021GA1FP	IMAGE EST {IMAGE:618363}	618363 IMAGE Consortium	AA175078
3558 A	12 021GA1FP	IMAGE EST {IMAGE:618256}	618256 IMAGE Consortium	AA170586
3559 C	2 021GA1FP	IMAGE EST {IMAGE:619484}	619484 IMAGE Consortium	AA172598
3560 C	4 021GA1FP	ESTs, Moderately similar to hypothetical protein [M.musculus] {IMAGE:619685}	619685 IMAGE Consortium	AA175975
3561 C	6 021GA1FP	Mus musculus fibrillin 2 (fbn2) gene, complete cds {IMAGE:617885}	617885 IMAGE Consortium	AA175306
3562 C	8 021GA1FP	ESTs, Moderately similar to RIBOSE-PHOSPHATE PYROPHOSPHOKINASE II [Rattus norvegicus] {IMAGE:617752}	617752 IMAGE Consortium	AA170738
3563 C	10 021GA1FP	ESTs {IMAGE:618483}	618483 IMAGE Consortium	AA175284
3564 C	12 021GA1FP	Mus musculus ICH-3 mRNA, complete cds {IMAGE:618272}	618272 IMAGE Consortium	AA175651
3565 E	2 021GA1FP	IMAGE EST {IMAGE:617193}	617193 IMAGE Consortium	AA174774
3566 E	4 021GA1FP	RAB1, member RAS oncogene family {IMAGE:619501}	619501 IMAGE Consortium	AA175510
3567 E	6 021GA1FP	IMAGE EST {IMAGE:619686}	619686 IMAGE Consortium	AA175973
3568 E	8 021GA1FP	IMAGE EST {IMAGE:619958}	619958 IMAGE Consortium	AA178037
3569 E	10 021GA1FP	ESTs {IMAGE:619895}	619895 IMAGE Consortium	AA173013

3570 E	12 021GA1FP	IMAGE EST {IMAGE:620261}	620261 IMAGE Consortium	AA177841
3571 G	2 021GA1FP	ESTs {IMAGE:619347}	619347 IMAGE Consortium	AA175729
3572 G	4 021GA1FP	ESTs, Moderately similar to PROBABLE METHYLMALONATE-SEMIALDEHYDE DEHYDROGENASE PRECURSOR [C.elegans] {IMAGE:617478}	617478 IMAGE Consortium	AA174585
3573 G	6 021GA1FP	ESTs, Highly similar to similar to human DNA-binding protein 5. [H.sapiens] {IMAGE:617365}	617365 IMAGE Consortium	AA174517
3574 G	8 021GA1FP	IMAGE EST {IMAGE:620030}	620030 IMAGE Consortium	AA178100
3575 G	10 021GA1FP	ESTs {IMAGE:620209}	620209 IMAGE Consortium	AA177920
3576 G	12 021GA1FP	IMAGE EST {IMAGE:618201}	618201 IMAGE Consortium	AA175313
3577 A	2 021NA1FQ	ESTs, Weakly similar to ZINC FINGER PROTEIN 91 [H.sapiens] {IMAGE:622878}	622878 IMAGE Consortium	AA184296
3578 A	4 021NA1FQ	ESTs, Weakly similar to Similarity to Yeast hypothetical protein 00926 [C.elegans] {IMAGE:623077}	623077 IMAGE Consortium	AA186084
3579 A	6 021NA1FQ	IMAGE EST {IMAGE:636324}	636324 IMAGE Consortium	AA189232
3580 A	8 021NA1FQ	ESTs, Weakly similar to Similarity to litosperm LEC14B protein [C.elegans] {IMAGE:634469}	634469 IMAGE Consortium	AA184958
3581 A	10 021NA1FQ	ESTs {IMAGE:634356}	634356 IMAGE Consortium	AA185432
3582 A	12 021NA1FQ	YY1 transcription factor {IMAGE:621788}	621788 IMAGE Consortium	AA177341
3583 C	2 021NA1FQ	ESTs {IMAGE:634589}	634589 IMAGE Consortium	AA185943
3584 C	4 021NA1FQ	IMAGE EST {IMAGE:634525}	634525 IMAGE Consortium	AA185617
3585 C	6 021NA1FQ	IMAGE EST {IMAGE:634460}	634460 IMAGE Consortium	AA185688
3586 C	8 021NA1FQ	IMAGE EST {IMAGE:621964}	621964 IMAGE Consortium	AA183748
3587 C	10 021NA1FQ	ESTs, Weakly similar to Rga [D.melanogaster] {IMAGE:634769}	634769 IMAGE Consortium	AA154673
3588 C	12 021NA1FQ	Mus musculus sphingosine kinase (SPHK1b) mRNA, complete cds {IMAGE:634846}	634846 IMAGE Consortium	AA154763
3589 E	2 021NA1FQ	IMAGE EST {IMAGE:621774}	621774 IMAGE Consortium	AA184084
3590 E	4 021NA1FQ	IMAGE EST {IMAGE:634484}	634484 IMAGE Consortium	AA185695
3591 E	6 021NA1FQ	IMAGE EST {IMAGE:634753}	634753 IMAGE Consortium	AA163328
3592 E	8 021NA1FQ	ESTs {IMAGE:634882}	634882 IMAGE Consortium	AA163624
3593 E	10 021NA1FQ	ESTs {IMAGE:635257}	635257 IMAGE Consortium	AA189498
3594 E	12 021NA1FQ	IMAGE EST {IMAGE:622364}	622364 IMAGE Consortium	AA177745
3595 G	2 021NA1FQ	ESTs {IMAGE:622039}	622039 IMAGE Consortium	AA183836
3596 G	4 021NA1FQ	ESTs {IMAGE:635114}	635114 IMAGE Consortium	AA162386
3597 G	6 021NA1FQ	IMAGE EST {IMAGE:622359}	622359 IMAGE Consortium	AA184200
3598 G	8 021NA1FQ	ESTs {IMAGE:636124}	636124 IMAGE Consortium	AA189375
3599 G	10 021NA1FQ	IMAGE EST {IMAGE:621460}	621460 IMAGE Consortium	AA178652
3600 G	12 021NA1FQ	IMAGE EST {IMAGE:634455}	634455 IMAGE Consortium	AA185484
3601 A	2 021UA1FR	Mouse mRNA for ASCT2, complete cds {IMAGE:619563}	619563 IMAGE Consortium	AA175769
3602 A	4 021UA1FR	ESTs {IMAGE:619430}	619430 IMAGE Consortium	AA175625
3603 A	6 021UA1FR	IMAGE EST {IMAGE:617437}	617437 IMAGE Consortium	AA174397
3604 A	8 021UA1FR	ESTs {IMAGE:617687}	617687 IMAGE Consortium	AA170661

3605 A	10 021UA1FR	ESTs {IMAGE:619696}	619696 IMAGE Consortium	AA172813
3606 A	12 021UA1FR	ZINC FINGER PROTEIN ZFP-26 {IMAGE:620364}	620364 IMAGE Consortium	AA178164
3607 C	2 021UA1FR	ESTs {IMAGE:619922}	619922 IMAGE Consortium	AA177731
3608 C	4 021UA1FR	ESTs {IMAGE:619926}	619926 IMAGE Consortium	AA177733
3609 C	6 021UA1FR	Mus musculus mRNA for cathepsin S {IMAGE:620027}	620027 IMAGE Consortium	AA178121
3610 C	8 021UA1FR	ESTs {IMAGE:619745}	619745 IMAGE Consortium	AA172771
3611 C	10 021UA1FR	IMAGE EST {IMAGE:620101}	620101 IMAGE Consortium	AA176037
3612 C	12 021UA1FR	IMAGE EST {IMAGE:620326}	620326 IMAGE Consortium	AA172559
3613 E	2 021UA1FR	ESTs {IMAGE:619970}	619970 IMAGE Consortium	AA178026
3614 E	4 021UA1FR	IMAGE EST {IMAGE:619950}	619950 IMAGE Consortium	AA178021
3615 E	6 021UA1FR	ESTs, Highly similar to GLUTATHIONE REDUCTASE [Escherichia coli] {IMAGE:619836}	619836 IMAGE Consortium	AA177872
3616 E	8 021UA1FR	ESTs {IMAGE:619746}	619746 IMAGE Consortium	AA172768
3617 E	10 021UA1FR	ESTs {IMAGE:620221}	620221 IMAGE Consortium	AA177976
3618 E	12 021UA1FR	Mouse calcineurin catalytic subunit mRNA, complete cds {IMAGE:620546}	620546 IMAGE Consortium	AA178283
3619 G	2 021UA1FR	ESTs, Highly similar to E1B 19K/Bcl-2-interacting protein Nip1 [H.sapiens] {IMAGE:620341}	620341 IMAGE Consortium	AA178161
3620 G	4 021UA1FR	IMAGE EST {IMAGE:618687}	618687 IMAGE Consortium	AA175097
3621 G	6 021UA1FR	ESTs {IMAGE:621012}	621012 IMAGE Consortium	AA178628
3622 G	8 021UA1FR	IMAGE EST {IMAGE:621459}	621459 IMAGE Consortium	AA178673
3623 G	10 021UA1FR	IMAGE EST {IMAGE:621393}	621393 IMAGE Consortium	AA178708
3624 G	12 021UA1FR	IMAGE EST {IMAGE:634588}	634588 IMAGE Consortium	AA185939
3625 A	2 0211A1FS	IMAGE EST {IMAGE:635943}	635943 IMAGE Consortium	AA183285
3626 A	4 0211A1FS	Alcohol dehydrogenase 5 {IMAGE:636207}	636207 IMAGE Consortium	AA183192
3627 A	6 0211A1FS	IMAGE EST {IMAGE:634233}	634233 IMAGE Consortium	AA182113
3628 A	8 0211A1FS	IMAGE EST {IMAGE:634466}	634466 IMAGE Consortium	AA184909
3629 A	10 0211A1FS	IMAGE EST {IMAGE:621954}	621954 IMAGE Consortium	AA177432
3630 A	12 0211A1FS	IMAGE EST {IMAGE:634658}	634658 IMAGE Consortium	AA154741
3631 C	2 0211A1FS	ESTs, Highly similar to N-TERMINAL ACETYLTRANSFERASE COMPLEX ARD1 SUBUNIT HOMOLOG [Homo sapiens] {IMAGE:634371}	634371 IMAGE Consortium	AA185526
3632 C	4 0211A1FS	IMAGE EST {IMAGE:634503}	634503 IMAGE Consortium	AA185562
3633 C	6 0211A1FS	Solute carrier family 12, member 4 {IMAGE:634461}	634461 IMAGE Consortium	AA185691
3634 C	8 0211A1FS	IMAGE EST {IMAGE:622085}	622085 IMAGE Consortium	AA184015
3635 C	10 0211A1FS	IMAGE EST {IMAGE:622257}	622257 IMAGE Consortium	AA183609
3636 C	12 0211A1FS	ESTs {IMAGE:635116}	635116 IMAGE Consortium	AA162387
3637 E	2 0211A1FS	IMAGE EST {IMAGE:634400}	634400 IMAGE Consortium	AA185444
3638 E	4 0211A1FS	ESTs, Weakly similar to cDNA EST EMBL:D75506 comes from this gene [C.elegans] {IMAGE:634360}	634360 IMAGE Consortium	AA185218
3639 E	6 0211A1FS	IMAGE EST {IMAGE:634581}	634581 IMAGE Consortium	AA189648
3640 E	8 0211A1FS	Mus musculus plasma membrane protein syntaxin-4 mRNA, complete cds {IMAGE:634982}	634982 IMAGE Consortium	AA164051

3641 E	10 0211A1FS	ESTs, Weakly similar to reverse transcriptase [M.musculus] {IMAGE:634820}	634820 IMAGE Consortium	AA154685
3642 E	12 0211A1FS	IMAGE EST {IMAGE:635117}	635117 IMAGE Consortium	AA162417
3643 G	2 0211A1FS	ESTs {IMAGE:634515}	634515 IMAGE Consortium	AA184991
3644 G	4 0211A1FS	IMAGE EST {IMAGE:621635}	621635 IMAGE Consortium	AA183654
3645 G	6 0211A1FS	ESTs, Weakly similar to zinc finger protein [M.musculus] {IMAGE:621763}	621763 IMAGE Consortium	AA177740
3646 G	8 0211A1FS	IMAGE EST {IMAGE:622128}	622128 IMAGE Consortium	AA184039
3647 G	10 0211A1FS	Regulatory protein, T lymphocyte 1 {IMAGE:622181}	622181 IMAGE Consortium	AA183432
3648 G	12 0211A1FS	IMAGE EST {IMAGE:634677}	634677 IMAGE Consortium	AA155388
3649 A	2 0218A1FT	IMAGE EST {IMAGE:621829}	621829 IMAGE Consortium	AA184026
3650 A	4 0218A1FT	ESTs {IMAGE:621787}	621787 IMAGE Consortium	AA177344
3651 A	6 0218A1FT	IMAGE EST {IMAGE:622130}	622130 IMAGE Consortium	AA184040
3652 A	8 0218A1FT	ESTs {IMAGE:622253}	622253 IMAGE Consortium	AA183369
3653 A	10 0218A1FT	IMAGE EST {IMAGE:634749}	634749 IMAGE Consortium	AA154648
3654 A	12 0218A1FT	EST {IMAGE:635213}	635213 IMAGE Consortium	AA189528
3655 C	2 0218A1FT	ESTs {IMAGE:634354}	634354 IMAGE Consortium	AA185431
3656 C	4 0218A1FT	ESTs {IMAGE:634625}	634625 IMAGE Consortium	AA189661
3657 C	6 0218A1FT	IMAGE EST {IMAGE:634656}	634656 IMAGE Consortium	AA154740
3658 C	8 0218A1FT	IMAGE EST {IMAGE:634685}	634685 IMAGE Consortium	AA154789
3659 C	10 0218A1FT	IMAGE EST {IMAGE:634792}	634792 IMAGE Consortium	AA154676
3660 C	12 0218A1FT	IMAGE EST {IMAGE:622174}	622174 IMAGE Consortium	AA177599
3661 E	2 0218A1FT	IMAGE EST {IMAGE:637183}	637183 IMAGE Consortium	AA189703
3662 E	4 0218A1FT	IMAGE EST {IMAGE:639709}	639709 IMAGE Consortium	AA197399
3663 E	6 0218A1FT	ESTs, Highly similar to GLYCINE AMIDINOTRANSFERASE PRECURSOR [R.norvegicus] {IMAGE:637453}	637453 IMAGE Consortium	AA185055
3664 E	8 0218A1FT	ESTs {IMAGE:637913}	637913 IMAGE Consortium	AA190006
3665 E	10 0218A1FT	Mus musculus mRNA for N-glycan alpha 2,8-sialyltransferase {IMAGE:638406}	638406 IMAGE Consortium	AA217008
3666 E	12 0218A1FT	ESTs {IMAGE:638389}	638389 IMAGE Consortium	AA200043
3667 G	2 0218A1FT	IMAGE EST {IMAGE:638454}	638454 IMAGE Consortium	AA216987
3668 G	4 0218A1FT	ESTs {IMAGE:638198}	638198 IMAGE Consortium	AA200321
3669 G	6 0218A1FT	IMAGE EST {IMAGE:641143}	641143 IMAGE Consortium	AA204455
3670 G	8 0218A1FT	Biglycan {IMAGE:638662}	638662 IMAGE Consortium	AA200264
3671 G	10 0218A1FT	IMAGE EST {IMAGE:639056}	639056 IMAGE Consortium	AA199297
3672 G	12 0218A1FT	IMAGE EST {IMAGE:636753}	636753 IMAGE Consortium	AA183317
3673 A	2 021FA1FU	IMAGE EST {IMAGE:637440}	637440 IMAGE Consortium	AA185050
3674 A	4 021FA1FU	IMAGE EST {IMAGE:640320}	640320 IMAGE Consortium	AA199429
3675 A	6 021FA1FU	ESTs, Highly similar to HYPOTHETICAL PROTEIN KIAA0036 [H.sapiens] {IMAGE:640287}	640287 IMAGE Consortium	AA198784
3676 A	8 021FA1FU	IMAGE EST {IMAGE:640732}	640732 IMAGE Consortium	AA203877
3677 A	10 021FA1FU	Frequently rearranged in advanced T-cell lymphomas {IMAGE:638394}	638394 IMAGE Consortium	AA200201

3678 A	12 021FA1FU	IMAGE EST {IMAGE:638596}	638596 IMAGE Consortium	AA200888
3679 C	2 021FA1FU	IMAGE EST {IMAGE:639640}	639640 IMAGE Consortium	AA197461
3680 C	4 021FA1FU	ESTs {IMAGE:637930}	637930 IMAGE Consortium	AA189993
3681 C	6 021FA1FU	IMAGE EST {IMAGE:638284}	638284 IMAGE Consortium	AA200747
3682 C	8 021FA1FU	ESTs, Weakly similar to DNAJ PROTEIN HOMOLOG 2 [R.norvegicus] {IMAGE:641016}	641016 IMAGE Consortium	AA204094
3683 C	10 021FA1FU	ESTs {IMAGE:641299}	641299 IMAGE Consortium	AA200596
3684 C	12 021FA1FU	TPA regulated locus {IMAGE:639530}	639530 IMAGE Consortium	AA198785
3685 E	2 021FA1FU	IMAGE EST {IMAGE:640501}	640501 IMAGE Consortium	AA204066
3686 E	4 021FA1FU	IMAGE EST {IMAGE:638599}	638599 IMAGE Consortium	AA200892
3687 E	6 021FA1FU	ESTs {IMAGE:638682}	638682 IMAGE Consortium	AA250415
3688 E	8 021FA1FU	INTERFERON REGULATORY FACTOR 2 {IMAGE:639197}	639197 IMAGE Consortium	AA200609
3689 E	10 021FA1FU	ESTs {IMAGE:636660}	636660 IMAGE Consortium	AA189474
3690 E	12 021FA1FU	ESTs, Highly similar to PUTATIVE SERYL-TRNA SYNTHETASE YHR011W [Saccharomyces cerevisiae] {IMAGE:637278}	637278 IMAGE Consortium	AA189797
3691 G	2 021FA1FU	Mus musculus BALB/c mannosyl-oligosaccharide alpha-1,2-mannosidase mRNA, complete cds {IMAGE:638791}	638791 IMAGE Consortium	AA199988
3692 G	4 021FA1FU	IMAGE EST {IMAGE:638613}	638613 IMAGE Consortium	AA200252
3693 G	6 021FA1FU	Acid beta glucosidase {IMAGE:639442}	639442 IMAGE Consortium	AA198668
3694 G	8 021FA1FU	IMAGE EST {IMAGE:639572}	639572 IMAGE Consortium	AA200835
3695 G	10 021FA1FU	IMAGE EST {IMAGE:637442}	637442 IMAGE Consortium	AA185051
3696 G	12 021FA1FU	IMAGE EST {IMAGE:637333}	637333 IMAGE Consortium	AA184831
3697 A	2 021MA1FV	ESTs {IMAGE:638495}	638495 IMAGE Consortium	AA198993
3698 A	4 021MA1FV	Mus musculus CTP synthetase homolog (CTPsH) mRNA, complete cds {IMAGE:641397}	641397 IMAGE Consortium	AA200393
3699 A	6 021MA1FV	Mus musculus mRNA for M-Ras, complete cds {IMAGE:638879}	638879 IMAGE Consortium	AA200952
3700 A	8 021MA1FV	IMAGE EST {IMAGE:639415}	639415 IMAGE Consortium	AA199123
3701 A	10 021MA1FV	IMAGE EST {IMAGE:639519}	639519 IMAGE Consortium	AA198708
3702 A	12 021MA1FV	ESTs {IMAGE:639991}	639991 IMAGE Consortium	AA197478
3703 C	2 021MA1FV	IMAGE EST {IMAGE:641484}	641484 IMAGE Consortium	AA200464
3704 C	4 021MA1FV	IMAGE EST {IMAGE:641904}	641904 IMAGE Consortium	AA210212
3705 C	6 021MA1FV	ESTs {IMAGE:638975}	638975 IMAGE Consortium	AA200135
3706 C	8 021MA1FV	ESTs {IMAGE:639440}	639440 IMAGE Consortium	AA198667
3707 C	10 021MA1FV	ESTs {IMAGE:639376}	639376 IMAGE Consortium	AA200797
3708 C	12 021MA1FV	IMAGE EST {IMAGE:639704}	639704 IMAGE Consortium	AA197393
3709 E	2 021MA1FV	IMAGE EST {IMAGE:636407}	636407 IMAGE Consortium	AA182094
3710 E	4 021MA1FV	IMAGE EST {IMAGE:636682}	636682 IMAGE Consortium	AA189273
3711 E	6 021MA1FV	ESTs {IMAGE:639592}	639592 IMAGE Consortium	AA200894
3712 E	8 021MA1FV	IMAGE EST {IMAGE:639865}	639865 IMAGE Consortium	AA197643
3713 E	10 021MA1FV	ESTs {IMAGE:637078}	637078 IMAGE Consortium	AA185500
3714 E	12 021MA1FV	Mus musculus Fas-binding protein (Daxx) mRNA, complete	637568 IMAGE Consortium	AA184772

		cds {IMAGE:637568}		
3715 G	2	021MA1FV ESTs {IMAGE:639481}	639481	IMAGE Consortium AA198560
3716 G	4	021MA1FV ESTs {IMAGE:636569}	636569	IMAGE Consortium AA204571
3717 G	6	021MA1FV IMAGE EST {IMAGE:637042}	637042	IMAGE Consortium AA185657
3718 G	8	021MA1FV ESTs {IMAGE:639807}	639807	IMAGE Consortium AA197571
3719 G	10	021MA1FV IMAGE EST {IMAGE:640191}	640191	IMAGE Consortium AA197618
3720 G	12	021MA1FV ESTs {IMAGE:638265}	638265	IMAGE Consortium AA200196
3721 A	2	021TA1FW ESTs {IMAGE:644982}	644982	IMAGE Consortium AA210344
3722 A	4	021TA1FW ESTs {IMAGE:641806}	641806	IMAGE Consortium AA210170
3723 A	6	021TA1FW Mouse (clone RAD11) T-cell receptor rearranged delta-chain mRNA V-D-J-C region, 3' end {IMAGE:642223}	642223	IMAGE Consortium AA212375
3724 A	8	021TA1FW ESTs {IMAGE:652923}	652923	IMAGE Consortium AA216865
3725 A	10	021TA1FW IMAGE EST {IMAGE:652956}	652956	IMAGE Consortium AA217125
3726 A	12	021TA1FW ESTs {IMAGE:653402}	653402	IMAGE Consortium AA240299
3727 C	2	021TA1FW ESTs, Moderately similar to RAF SERINE/THREONINE-PROTEIN KINASE TRANSFORMING PROTEIN [Murine sarcoma virus 3611] {IMAGE:641845}	641845	IMAGE Consortium AA209788
3728 C	4	021TA1FW Signal transducer and activator of transcription 6 {IMAGE:644971}	644971	IMAGE Consortium AA210400
3729 C	6	021TA1FW IMAGE EST {IMAGE:652593}	652593	IMAGE Consortium AA216829
3730 C	8	021TA1FW ESTs {IMAGE:652601}	652601	IMAGE Consortium AA217059
3731 C	10	021TA1FW IMAGE EST {IMAGE:642115}	642115	IMAGE Consortium AA210152
3732 C	12	021TA1FW Dlk1-like homolog (Drosophila) {IMAGE:642510}	642510	IMAGE Consortium AA185478
3733 E	2	021TA1FW ESTs {IMAGE:641630}	641630	IMAGE Consortium AA209892
3734 E	4	021TA1FW ESTs, Highly similar to CAMP-DEPENDENT 3',5'-CYCLIC PHOSPHODIESTERASE 4B [Homo sapiens] {IMAGE:641682}	641682	IMAGE Consortium AA209984
3735 E	6	021TA1FW ESTs {IMAGE:641902}	641902	IMAGE Consortium AA209939
3736 E	8	021TA1FW ESTs {IMAGE:642205}	642205	IMAGE Consortium AA212243
3737 E	10	021TA1FW IMAGE EST {IMAGE:642165}	642165	IMAGE Consortium AA212307
3738 E	12	021TA1FW ESTs {IMAGE:642379}	642379	IMAGE Consortium AA185202
3739 G	2	021TA1FW Cytochrome P450, 2e1, ethanol inducible {IMAGE:642224}	642224	IMAGE Consortium AA212531
3740 G	4	021TA1FW ESTs {IMAGE:642301}	642301	IMAGE Consortium AA212559
3741 G	6	021TA1FW ESTs {IMAGE:652983}	652983	IMAGE Consortium AA217415
3742 G	8	021TA1FW IMAGE EST {IMAGE:653167}	653167	IMAGE Consortium AA217116
3743 G	10	021TA1FW IMAGE EST {IMAGE:642941}	642941	IMAGE Consortium AA185404
3744 G	12	021TA1FW ESTs {IMAGE:643131}	643131	IMAGE Consortium AA189876
3745 A	2	0210A1FX IMAGE EST {IMAGE:653031}	653031	IMAGE Consortium AA217394
3746 A	4	0210A1FX IMAGE EST {IMAGE:653219}	653219	IMAGE Consortium AA217290
3747 A	6	0210A1FX IMAGE EST {IMAGE:642847}	642847	IMAGE Consortium AA185648
3748 A	8	0210A1FX Mus musculus RING zinc finger protein (Rzf) mRNA, complete cds {IMAGE:643108}	643108	IMAGE Consortium AA189868
3749 A	10	0210A1FX IMAGE EST {IMAGE:643815}	643815	IMAGE Consortium AA203914

3750 A	12 0210A1FX	Kinesin family member 3b {IMAGE:641650}	641650 IMAGE Consortium	AA209952
3751 C	2 0210A1FX	Mus musculus apoptotic protease activating factor 1 mRNA, complete cds {IMAGE:657503}	657503 IMAGE Consortium	AA198703
3752 C	4 0210A1FX	ESTs {IMAGE:657851}	657851 IMAGE Consortium	AA199219
3753 C	6 0210A1FX	ESTs, Weakly similar to CYSTEINE-STRING PROTEIN [D.melanogaster] {IMAGE:643696}	643696 IMAGE Consortium	AA204613
3754 C	8 0210A1FX	ESTs {IMAGE:643914}	643914 IMAGE Consortium	AA203942
3755 C	10 0210A1FX	ESTs {IMAGE:644952}	644952 IMAGE Consortium	AA210317
3756 C	12 0210A1FX	ESTs {IMAGE:644964}	644964 IMAGE Consortium	AA210380
3757 E	2 0210A1FX	IMAGE EST {IMAGE:585041}	585041 IMAGE Consortium	AA125425
3758 E	4 0210A1FX	ESTs {IMAGE:643792}	643792 IMAGE Consortium	AA204021
3759 E	6 0210A1FX	ESTs {IMAGE:644109}	644109 IMAGE Consortium	AA208763
3760 E	8 0210A1FX	ESTs {IMAGE:644857}	644857 IMAGE Consortium	AA210597
3761 E	10 0210A1FX	ESTs {IMAGE:641637}	641637 IMAGE Consortium	AA221291
3762 E	12 0210A1FX	ESTs, Highly similar to GLUTAMINASE, LIVER ISOFORM PRECURSOR [Rattus norvegicus] {IMAGE:652644}	652644 IMAGE Consortium	AA216842
3763 G	2 0210A1FX	ESTs {IMAGE:641795}	641795 IMAGE Consortium	AA210022
3764 G	4 0210A1FX	Mus musculus cleavage and polyadenylation specificity factor (MCPSF) mRNA, complete cds {IMAGE:641608}	641608 IMAGE Consortium	AA221269
3765 G	6 0210A1FX	IMAGE EST {IMAGE:641661}	641661 IMAGE Consortium	AA209981
3766 G	8 0210A1FX	ESTs {IMAGE:652800}	652800 IMAGE Consortium	AA217508
3767 G	10 0210A1FX	IMAGE EST {IMAGE:642625}	642625 IMAGE Consortium	AA185346
3768 G	12 0210A1FX	IMAGE EST {IMAGE:653633}	653633 IMAGE Consortium	AA217820
3769 A	2 0217A1FY	ESTs, Highly similar to HYPOTHETICAL PROTEIN KIAA0174 [H.sapiens] {IMAGE:832713}	832713 IMAGE Consortium	AA427055
3770 A	4 0217A1FY	TRANSCRIPTION FACTOR MTF-1 {IMAGE:849197}	849197 IMAGE Consortium	AA469826
3771 A	6 0217A1FY	ESTs, Weakly similar to hypertension-related protein [R.norvegicus] {IMAGE:846294}	846294 IMAGE Consortium	AA416078
3772 A	8 0217A1FY	Signal transducing adaptor molecule (SH3 domain and ITAM motif) 1 {IMAGE:849879}	849879 IMAGE Consortium	AA450729
3773 A	10 0217A1FY	ESTs {IMAGE:850280}	850280 IMAGE Consortium	AA451495
3774 A	12 0217A1FY	ESTs {IMAGE:850078}	850078 IMAGE Consortium	AA450631
3775 C	2 0217A1FY	ESTs {IMAGE:851223}	851223 IMAGE Consortium	AA462944
3776 C	4 0217A1FY	IMAGE EST {IMAGE:860259}	860259 IMAGE Consortium	AA500910
3777 C	6 0217A1FY	Mus musculus syntaxin-like protein 3I35 mRNA, complete cds {IMAGE:850251}	850251 IMAGE Consortium	AA451480
3778 C	8 0217A1FY	IMAGE EST {IMAGE:851064}	851064 IMAGE Consortium	AA462611
3779 C	10 0217A1FY	IMAGE EST {IMAGE:851329}	851329 IMAGE Consortium	AA461661
3780 C	12 0217A1FY	IMAGE EST {IMAGE:860380}	860380 IMAGE Consortium	AA499577
3781 E	2 0217A1FY	ESTs, Highly similar to ARGINYL-TRNA SYNTHETASE [Cricetulus longicaudatus] {IMAGE:850426}	850426 IMAGE Consortium	AA451273
3782 E	4 0217A1FY	IMAGE EST {IMAGE:851205}	851205 IMAGE Consortium	AA462871
3783 E	6 0217A1FY	Mus musculus phospholipid scramblase mRNA, complete cds	850206 IMAGE Consortium	AA451429



3784 E	8 0217A1FY	{IMAGE:850206} Mus musculus CD-1 orphan receptor TAK1 (TAK1) mRNA, complete cds {IMAGE:860347}	860347 IMAGE Consortium	AA501045
3785 E	10 0217A1FY	Split hand/foot deleted gene 1 {IMAGE:850971}	850971 IMAGE Consortium	AA462396
3786 E	12 0217A1FY	ESTs {IMAGE:851014}	851014 IMAGE Consortium	AA462416
3787 G	2 0217A1FY	ESTs {IMAGE:870973}	870973 IMAGE Consortium	AA461746
3788 G	4 0217A1FY	IMAGE EST {IMAGE:871502}	871502 IMAGE Consortium	AA462082
3789 G	6 0217A1FY	ESTs, Highly similar to HYPOTHETICAL 77.3 KD PROTEIN T05G5.8 IN CHROMOSOME III [Caenorhabditis elegans] {IMAGE:874128}	874128 IMAGE Consortium	AA471754
3790 G	8 0217A1FY	IMAGE EST {IMAGE:874345}	874345 IMAGE Consortium	AA474937
3791 G	10 0217A1FY	IMAGE EST {IMAGE:871701}	871701 IMAGE Consortium	AA462383
3792 G	12 0217A1FY	ESTs {IMAGE:872986}	872986 IMAGE Consortium	AA466235
3793 A	2 021EA1FZ	IMAGE EST {IMAGE:652835}	652835 IMAGE Consortium	AA217632
3794 A	4 021EA1FZ	IMAGE EST {IMAGE:653112}	653112 IMAGE Consortium	AA217528
3795 A	6 021EA1FZ	ESTs {IMAGE:653078}	653078 IMAGE Consortium	AA216884
3796 A	8 021EA1FZ	ESTs {IMAGE:653761}	653761 IMAGE Consortium	AA218059
3797 A	10 021EA1FZ	ESTs, Weakly similar to similar to Zinc finger, C2H2 type [C.elegans] {IMAGE:643152}	643152 IMAGE Consortium	AA189879
3798 A	12 021EA1FZ	IMAGE EST {IMAGE:643384}	643384 IMAGE Consortium	AA186107
3799 C	2 021EA1FZ	ESTs {IMAGE:652753}	652753 IMAGE Consortium	AA217200
3800 C	4 021EA1FZ	ESTs {IMAGE:652859}	652859 IMAGE Consortium	AA217639
3801 C	6 021EA1FZ	ESTs {IMAGE:653137}	653137 IMAGE Consortium	AA217603
3802 C	8 021EA1FZ	IMAGE EST {IMAGE:653491}	653491 IMAGE Consortium	AA217572
3803 C	10 021EA1FZ	ESTs, Weakly similar to C56C10.7 [C.elegans] {IMAGE:643122}	643122 IMAGE Consortium	AA190060
3804 C	12 021EA1FZ	ESTs {IMAGE:643725}	643725 IMAGE Consortium	AA203775
3805 E	2 021EA1FZ	IMAGE EST {IMAGE:642497}	642497 IMAGE Consortium	AA189196
3806 E	4 021EA1FZ	IMAGE EST {IMAGE:653813}	653813 IMAGE Consortium	AA218073
3807 E	6 021EA1FZ	ESTs {IMAGE:643273}	643273 IMAGE Consortium	AA189590
3808 E	8 021EA1FZ	Mus musculus mRNA for CDC6 protein {IMAGE:643194}	643194 IMAGE Consortium	AA189836
3809 E	10 021EA1FZ	ESTs {IMAGE:643474}	643474 IMAGE Consortium	AA204106
3810 E	12 021EA1FZ	ESTs {IMAGE:658713}	658713 IMAGE Consortium	AA221880
3811 G	2 021EA1FZ	Mus musculus signal recognition particle receptor beta subunit mRNA, complete cds {IMAGE:847006}	847006 IMAGE Consortium	AA419748
3812 G	4 021EA1FZ	Peripheral myelin protein, 22 kDa {IMAGE:846064}	846064 IMAGE Consortium	AA416246
3813 G	6 021EA1FZ	ESTs {IMAGE:832183}	832183 IMAGE Consortium	AA423601
3814 G	8 021EA1FZ	Retinoblastoma 1 {IMAGE:832714}	832714 IMAGE Consortium	AA427019
3815 G	10 021EA1FZ	IMAGE EST {IMAGE:832451}	832451 IMAGE Consortium	AA427236
3816 G	12 021EA1FZ	IMAGE EST {IMAGE:832428}	832428 IMAGE Consortium	AA427166
3817 A	2 021EA1G0	Max interacting protein 1 {IMAGE:874202}	874202 IMAGE Consortium	AA472395
3818 A	4 021EA1G0	ESTs {IMAGE:874194}	874194 IMAGE Consortium	AA472300
3819 A	6 021EA1G0	IMAGE EST {IMAGE:874572}	874572 IMAGE Consortium	AA472340

3820 A	8 021EA1G0	ESTs {IMAGE:874629}	874629 IMAGE Consortium	AA472465
3821 A	10 021EA1G0	IMAGE EST {IMAGE:873758}	873758 IMAGE Consortium	AA471929
3822 A	12 021EA1G0	IMAGE EST {IMAGE:874131}	874131 IMAGE Consortium	AA471764
3823 C	2 021EA1G0	ESTs {IMAGE:874349}	874349 IMAGE Consortium	AA474939
3824 C	4 021EA1G0	IMAGE EST {IMAGE:871585}	871585 IMAGE Consortium	AA462223
3825 C	6 021EA1G0	IMAGE EST {IMAGE:873812}	873812 IMAGE Consortium	AA471802
3826 C	8 021EA1G0	IMAGE EST {IMAGE:874541}	874541 IMAGE Consortium	AA472325
3827 C	10 021EA1G0	ESTs {IMAGE:871705}	871705 IMAGE Consortium	AA462486
3828 C	12 021EA1G0	ESTs, Highly similar to PUTATIVE ATP-DEPENDENT RNA HELICASE K03H1.2 IN CHROMOSOME III [Caenorhabditis elegans] {IMAGE:864368}	864368 IMAGE Consortium	AA458128
3829 E	2 021EA1G0	IMAGE EST {IMAGE:870852}	870852 IMAGE Consortium	AA462035
3830 E	4 021EA1G0	ESTs {IMAGE:871226}	871226 IMAGE Consortium	AA462874
3831 E	6 021EA1G0	ESTs, Weakly similar to Y38A8.1 gene product [C.elegans] {IMAGE:871562}	871562 IMAGE Consortium	AA462190
3832 E	8 021EA1G0	ESTs {IMAGE:873489}	873489 IMAGE Consortium	AA473904
3833 E	10 021EA1G0	Xanthine dehydrogenase {IMAGE:873928}	873928 IMAGE Consortium	AA472074
3834 E	12 021EA1G0	IMAGE EST {IMAGE:874806}	874806 IMAGE Consortium	AA472743
3835 G	2 021EA1G0	Annexin VI, p68 {IMAGE:876698}	876698 IMAGE Consortium	AA499296
3836 G	4 021EA1G0	ESTs {IMAGE:888577}	888577 IMAGE Consortium	AA498622
3837 G	6 021EA1G0	IMAGE EST {IMAGE:902466}	902466 IMAGE Consortium	AA517387
3838 G	8 021EA1G0	Nucleolin {IMAGE:903284}	903284 IMAGE Consortium	AA518999
3839 G	10 021EA1G0	ESTs {IMAGE:890837}	890837 IMAGE Consortium	AA510686
3840 G	12 021EA1G0	IMAGE EST {IMAGE:891477}	891477 IMAGE Consortium	AA521989
3841 A	2 021LA1G1	M.musculus mRNA for Sui1 {IMAGE:888602}	888602 IMAGE Consortium	AA498630
3842 A	4 021LA1G1	Pyruvate dehydrogenase E1alpha subunit {IMAGE:888842}	888842 IMAGE Consortium	AA466268
3843 A	6 021LA1G1	Transcription factor 4 {IMAGE:903046}	903046 IMAGE Consortium	AA518455
3844 A	8 021LA1G1	ESTs {IMAGE:903335}	903335 IMAGE Consortium	AA518231
3845 A	10 021LA1G1	IMAGE EST {IMAGE:890760}	890760 IMAGE Consortium	AA510479
3846 A	12 021LA1G1	Cathepsin C {IMAGE:876326}	876326 IMAGE Consortium	AA475774
3847 C	2 021LA1G1	ESTs {IMAGE:888605}	888605 IMAGE Consortium	AA498640
3848 C	4 021LA1G1	Lamin B1 {IMAGE:889295}	889295 IMAGE Consortium	AA498281
3849 C	6 021LA1G1	ESTs {IMAGE:890679}	890679 IMAGE Consortium	AA509979
3850 C	8 021LA1G1	ESTs {IMAGE:888687}	888687 IMAGE Consortium	AA498757
3851 C	10 021LA1G1	Eukaryotic translation initiation factor 3 {IMAGE:889816}	889816 IMAGE Consortium	AA498480
3852 C	12 021LA1G1	ESTs {IMAGE:876313}	876313 IMAGE Consortium	AA475778
3853 E	2 021LA1G1	ESTs {IMAGE:876838}	876838 IMAGE Consortium	AA497550
3854 E	4 021LA1G1	ESTs {IMAGE:889868}	889868 IMAGE Consortium	AA498565
3855 E	6 021LA1G1	Mus musculus eosinophil secondary granule protein (mEAR-2) mRNA, complete cds {IMAGE:891131}	891131 IMAGE Consortium	AA510161
3856 E	8 021LA1G1	ESTs {IMAGE:876479}	876479 IMAGE Consortium	AA475938
3857 E	10 021LA1G1	Mus musculus mRNA for ganglioside-induced differentiation associated protein 2 {IMAGE:876663}	876663 IMAGE Consortium	AA499267

3858 E	12 021LA1G1	Small proline-rich protein 2A {IMAGE:888512}	888512 IMAGE Consortium	AA497620
3859 G	2 021LA1G1	IMAGE EST {IMAGE:891094}	891094 IMAGE Consortium	AA510391
3860 G	4 021LA1G1	Mus musculus ERG-associated protein ESET mRNA, complete cds {IMAGE:876624}	876624 IMAGE Consortium	AA476068
3861 G	6 021LA1G1	40S RIBOSOMAL PROTEIN SA {IMAGE:888465}	888465 IMAGE Consortium	AA497590
3862 G	8 021LA1G1	ESTs, Moderately similar to H-2 class I histocompatibility antigen T15 [M.musculus] {IMAGE:889449}	889449 IMAGE Consortium	AA498469
3863 G	10 021LA1G1	ESTs {IMAGE:890568}	890568 IMAGE Consortium	AA509729
3864 G	12 021LA1G1	Amyloid beta (A4) precursor-like protein 2 {IMAGE:876058}	876058 IMAGE Consortium	AA475422
3865 A	2 021SA1G2	ESTs, Weakly similar to similar to Saccharomyces cerevisiae nuclear protein SNF7 [C.elegans] {IMAGE:922455}	922455 IMAGE Consortium	AA511304
3866 A	4 021SA1G2	ESTs {IMAGE:949758}	949758 IMAGE Consortium	AA537468
3867 A	6 021SA1G2	IMAGE EST {IMAGE:949623}	949623 IMAGE Consortium	AA537280
3868 A	8 021SA1G2	IMAGE EST {IMAGE:949845}	949845 IMAGE Consortium	AA537547
3869 A	10 021SA1G2	Prostaglandin E receptor EP4 subtype {IMAGE:904890}	904890 IMAGE Consortium	AA518372
3870 A	12 021SA1G2	IMAGE EST {IMAGE:920235}	920235 IMAGE Consortium	AA538228
3871 C	2 021SA1G2	ESTs {IMAGE:949042}	949042 IMAGE Consortium	AA543897
3872 C	4 021SA1G2	IMAGE EST {IMAGE:949241}	949241 IMAGE Consortium	AA544904
3873 C	6 021SA1G2	Cytochrome P450, 2e1, ethanol inducible {IMAGE:949657}	949657 IMAGE Consortium	AA537352
3874 C	8 021SA1G2	ESTs {IMAGE:949666}	949666 IMAGE Consortium	AA537347
3875 C	10 021SA1G2	ESTs {IMAGE:905429}	905429 IMAGE Consortium	AA529137
3876 C	12 021SA1G2	ESTs {IMAGE:949060}	949060 IMAGE Consortium	AA543916
3877 E	2 021SA1G2	ESTs {IMAGE:935463}	935463 IMAGE Consortium	AA543155
3878 E	4 021SA1G2	ESTs {IMAGE:947933}	947933 IMAGE Consortium	AA542278
3879 E	6 021SA1G2	IMAGE EST {IMAGE:949539}	949539 IMAGE Consortium	AA537140
3880 E	8 021SA1G2	ESTs {IMAGE:949715}	949715 IMAGE Consortium	AA537438
3881 E	10 021SA1G2	TIS11B PROTEIN {IMAGE:671377}	671377 IMAGE Consortium	AA222201
3882 E	12 021SA1G2	IMAGE EST {IMAGE:671530}	671530 IMAGE Consortium	AA240047
3883 G	2 021SA1G2	ESTs, Highly similar to NADH-UBIQUINONE OXIDOREDUCTASE SGDH SUBUNIT PRECURSOR [Bos taurus] {IMAGE:671212}	671212 IMAGE Consortium	AA222463
3884 G	4 021SA1G2	ESTs {IMAGE:671736}	671736 IMAGE Consortium	AA241808
3885 G	6 021SA1G2	ESTs {IMAGE:671867}	671867 IMAGE Consortium	AA242316
3886 G	8 021SA1G2	ESTs {IMAGE:671590}	671590 IMAGE Consortium	AA242064
3887 G	10 021SA1G2	IMAGE EST {IMAGE:672250}	672250 IMAGE Consortium	AA231370
3888 G	12 021SA1G2	ESTs, Highly similar to PROTEIN PHOSPHATASE 2C HOMOLOG 3 [Schizosaccharomyces pombe] {IMAGE:672294}	672294 IMAGE Consortium	AA231522
3889 A	2 021ZA1G3	Niemann Pick type C1 {IMAGE:902448}	902448 IMAGE Consortium	AA517379
3890 A	4 021ZA1G3	ESTs {IMAGE:902733}	902733 IMAGE Consortium	AA518757
3891 A	6 021ZA1G3	ESTs {IMAGE:903370}	903370 IMAGE Consortium	AA518354
3892 A	8 021ZA1G3	ESTs, Highly similar to HYPOTHETICAL 19.9 KD PROTEIN IN PUT3-CCE1 INTERGENIC REGION [Saccharomyces	903379 IMAGE Consortium	AA518369

		cerevisiae] {IMAGE:903379}		
3893 A	10 021ZA1G3	Cytochrome P450, 2e1, ethanol inducible {IMAGE:904738}	904738	IMAGE Consortium AA521764
3894 A	12 021ZA1G3	Mus musculus mRNA for N-acetylglucosamine-6-O-sulfotransferase, complete cds {IMAGE:920327}	920327	IMAGE Consortium AA538322
3895 C	2 021ZA1G3	IMAGE EST {IMAGE:920344}	920344	IMAGE Consortium AA538370
3896 C	4 021ZA1G3	Carbonic anhydrase 1 {IMAGE:922088}	922088	IMAGE Consortium AA512372
3897 C	6 021ZA1G3	ESTs, Weakly similar to P9513.2 gene product [S.cerevisiae] {IMAGE:922371}	922371	IMAGE Consortium AA512195
3898 C	8 021ZA1G3	IMAGE EST {IMAGE:949732}	949732	IMAGE Consortium AA537445
3899 C	10 021ZA1G3	IMAGE EST {IMAGE:949787}	949787	IMAGE Consortium AA537499
3900 C	12 021ZA1G3	ESTs, Highly similar to NADH-UBIQUINONE OXIDOREDUCTASE 9 KD SUBUNIT PRECURSOR [Bos taurus] {IMAGE:903911}	903911	IMAGE Consortium AA521758
3901 E	2 021ZA1G3	IMAGE EST {IMAGE:904680}	904680	IMAGE Consortium AA521581
3902 E	4 021ZA1G3	IMAGE EST {IMAGE:920146}	920146	IMAGE Consortium AA500129
3903 E	6 021ZA1G3	ESTs, Weakly similar to PEREGRIN [H.sapiens] {IMAGE:920640}	920640	IMAGE Consortium AA538272
3904 E	8 021ZA1G3	IMAGE EST {IMAGE:922018}	922018	IMAGE Consortium AA512269
3905 E	10 021ZA1G3	ESTs, Weakly similar to RING zinc finger protein [M.musculus] {IMAGE:922965}	922965	IMAGE Consortium AA511365
3906 E	12 021ZA1G3	ESTs, Highly similar to HYPOTHETICAL AMINOTRANSFERASE YJL060W [Saccharomyces cerevisiae] {IMAGE:935315}	935315	IMAGE Consortium AA542160
3907 G	2 021ZA1G3	IMAGE EST {IMAGE:904900}	904900	IMAGE Consortium AA519027
3908 G	4 021ZA1G3	ESTs {IMAGE:904769}	904769	IMAGE Consortium AA521797
3909 G	6 021ZA1G3	ESTs {IMAGE:948567}	948567	IMAGE Consortium AA544713
3910 G	8 021ZA1G3	Mus musculus peroxisome proliferator-activated receptor gamma binding protein mRNA, complete cds {IMAGE:948961}	948961	IMAGE Consortium AA544549
3911 G	10 021ZA1G3	ESTs, Moderately similar to TRANSCRIPTION FACTOR BF-1 [Rattus norvegicus] {IMAGE:948896}	948896	IMAGE Consortium AA544949
3912 G	12 021ZA1G3	ESTs {IMAGE:949824}	949824	IMAGE Consortium AA537527
3913 A	2 0216A1G4	ESTs {IMAGE:671169}	671169	IMAGE Consortium AA222347
3914 A	4 0216A1G4	ESTs, Highly similar to herpesvirus associated ubiquitin-specific protease [H.sapiens] {IMAGE:671784}	671784	IMAGE Consortium AA242262
3915 A	6 0216A1G4	ESTs, Highly similar to TRANSCRIPTION INITIATION FACTOR IIF, ALPHA SUBUNIT [Homo sapiens] {IMAGE:671628}	671628	IMAGE Consortium AA240978
3916 A	8 0216A1G4	Mus musculus protein phosphatase type 2A catalytic subunit alpha isoform mRNA, complete cds {IMAGE:671987}	671987	IMAGE Consortium AA245165
3917 A	10 0216A1G4	IMAGE EST {IMAGE:672344}	672344	IMAGE Consortium AA231557
3918 A	12 0216A1G4	IMAGE EST {IMAGE:672659}	672659	IMAGE Consortium AA231462
3919 C	2 0216A1G4	Cadherin 2 {IMAGE:671834}	671834	IMAGE Consortium AA242226
3920 C	4 0216A1G4	IMAGE EST {IMAGE:659759}	659759	IMAGE Consortium AA220077

3921 C	6 0216A1G4	ESTs, Highly similar to GLUTATHIONE S-TRANSFERASE 1 [Dianthus caryophyllus] {IMAGE:672972}	672972 IMAGE Consortium	AA231621
3922 C	8 0216A1G4	ESTs, Highly similar to CARBOXYPEPTIDASE B PRECURSOR [Rattus norvegicus] {IMAGE:676636}	676636 IMAGE Consortium	AA208905
3923 C	10 0216A1G4	Signal transducer and activator of transcription 5B {IMAGE:676651}	676651 IMAGE Consortium	AA212445
3924 C	12 0216A1G4	ESTs {IMAGE:676900}	676900 IMAGE Consortium	AA212964
3925 E	2 0216A1G4	ESTs, Weakly similar to HYPOTHETICAL 55.2 KD TRP-ASP REPEATS CONTAINING PROTEIN T10F2.4 IN CHROMOSOME III [C.elegans] {IMAGE:672881}	672881 IMAGE Consortium	AA231428
3926 E	4 0216A1G4	Coagulation factor IX {IMAGE:676179}	676179 IMAGE Consortium	AA209011
3927 E	6 0216A1G4	ESTs {IMAGE:676266}	676266 IMAGE Consortium	AA208946
3928 E	8 0216A1G4	IMAGE EST {IMAGE:676780}	676780 IMAGE Consortium	AA212598
3929 E	10 0216A1G4	Aconitase 1 {IMAGE:677092}	677092 IMAGE Consortium	AA212704
3930 E	12 0216A1G4	ESTs, Weakly similar to HYPOTHETICAL 23.1 KD PROTEIN IN SHP1-SEC17 INTERGENIC REGION [S.cerevisiae] {IMAGE:670753}	670753 IMAGE Consortium	AA221871
3931 G	2 0216A1G4	IMAGE EST {IMAGE:660884}	660884 IMAGE Consortium	AA216931
3932 G	4 0216A1G4	ESTs {IMAGE:669905}	669905 IMAGE Consortium	AA230647
3933 G	6 0216A1G4	ESTs {IMAGE:676532}	676532 IMAGE Consortium	AA209058
3934 G	8 0216A1G4	ESTs, Weakly similar to similar to nucleotide translocator [C.elegans] {IMAGE:677033}	677033 IMAGE Consortium	AA213247
3935 G	10 0216A1G4	ESTs {IMAGE:670788}	670788 IMAGE Consortium	AA230842
3936 G	12 0216A1G4	Mus musculus MEK kinase 3, mRNA, partial cds {IMAGE:671038}	671038 IMAGE Consortium	AA239991
3937 A	2 021DA1G5	IMAGE EST {IMAGE:676748}	676748 IMAGE Consortium	AA212577
3938 A	4 021DA1G5	ESTs, Highly similar to CHLORINE CHANNEL PROTEIN P64 [Bos taurus] {IMAGE:670375}	670375 IMAGE Consortium	AA221246
3939 A	6 021DA1G5	EGF-like module containing, mucin-like, hormone receptor-like sequence 1 {IMAGE:677598}	677598 IMAGE Consortium	AA212896
3940 A	8 021DA1G5	ESTs, Highly similar to HYPOTHETICAL 44.2 KD PROTEIN IN SCO2-MRF1 INTERGENIC REGION [Saccharomyces cerevisiae] {IMAGE:658480}	658480 IMAGE Consortium	AA221692
3941 A	10 021DA1G5	IMAGE EST {IMAGE:659248}	659248 IMAGE Consortium	AA217848
3942 A	12 021DA1G5	IMAGE EST {IMAGE:672261}	672261 IMAGE Consortium	AA230994
3943 C	2 021DA1G5	RAB11B, member RAS oncogene family {IMAGE:659369}	659369 IMAGE Consortium	AA218406
3944 C	4 021DA1G5	ESTs {IMAGE:660012}	660012 IMAGE Consortium	AA220714
3945 C	6 021DA1G5	IMAGE EST {IMAGE:660676}	660676 IMAGE Consortium	AA222039
3946 C	8 021DA1G5	Adaptor-related protein complex AP-1, beta 1 subunit {IMAGE:660866}	660866 IMAGE Consortium	AA221073
3947 C	10 021DA1G5	Mus musculus muskelin mRNA, complete cds {IMAGE:676111}	676111 IMAGE Consortium	AA208930
3948 C	12 021DA1G5	IMAGE EST {IMAGE:676523}	676523 IMAGE Consortium	AA209594
3949 E	2 021DA1G5	Follistatin-like {IMAGE:671910}	671910 IMAGE Consortium	AA242611

3950 E	4 021DA1G5	TAR (HIV) RNA binding protein 2 {IMAGE:661071}	661071 IMAGE Consortium	AA222734
3951 E	6 021DA1G5	Nascent polypeptide-associated complex alpha polypeptide {IMAGE:660958}	660958 IMAGE Consortium	AA230684
3952 E	8 021DA1G5	ESTs {IMAGE:676502}	676502 IMAGE Consortium	AA209568
3953 E	10 021DA1G5	ESTs, Highly similar to PROLINE OXIDASE PRECURSOR [Drosophila melanogaster] {IMAGE:676559}	676559 IMAGE Consortium	AA209015
3954 E	12 021DA1G5	Deiodinase, iodothyronine, type I {IMAGE:677180}	677180 IMAGE Consortium	AA212899
3955 G	2 021DA1G5	IMAGE EST {IMAGE:676921}	676921 IMAGE Consortium	AA212990
3956 G	4 021DA1G5	ESTs {IMAGE:670697}	670697 IMAGE Consortium	AA265657
3957 G	6 021DA1G5	ESTs, Highly similar to HYPOTHETICAL 31.1 KD PROTEIN B0228.7 IN CHROMOSOME II [Caenorhabditis elegans] {IMAGE:670830}	670830 IMAGE Consortium	AA221942
3958 G	8 021DA1G5	IMAGE EST {IMAGE:671257}	671257 IMAGE Consortium	AA222477
3959 G	10 021DA1G5	IMAGE EST {IMAGE:671865}	671865 IMAGE Consortium	AA242315
3960 G	12 021DA1G5	ESTs, Weakly similar to truncated form platelet-activating factor acetylhydrolase 45kD subunit [M.musculus] {IMAGE:672056}	672056 IMAGE Consortium	AA245181
3961 A	2 021KA1G6	IMAGE EST {IMAGE:692239}	692239 IMAGE Consortium	AA238239
3962 A	4 021KA1G6	IMAGE EST {IMAGE:679585}	679585 IMAGE Consortium	AA242673
3963 A	6 021KA1G6	IMAGE EST {IMAGE:680042}	680042 IMAGE Consortium	AA244954
3964 A	8 021KA1G6	IMAGE EST {IMAGE:680195}	680195 IMAGE Consortium	AA237922
3965 A	10 021KA1G6	ESTs, Highly similar to RAS-RELATED PROTEIN RAB-8 [Homo sapiens; Canis familiaris] {IMAGE:680060}	680060 IMAGE Consortium	AA244791
3966 A	12 021KA1G6	IMAGE EST {IMAGE:693774}	693774 IMAGE Consortium	AA238911
3967 C	2 021KA1G6	IMAGE EST {IMAGE:680148}	680148 IMAGE Consortium	AA237871
3968 C	4 021KA1G6	Mus musculus mRNA for alpha-albumine protein {IMAGE:693581}	693581 IMAGE Consortium	AA266651
3969 C	6 021KA1G6	ESTs {IMAGE:680658}	680658 IMAGE Consortium	AA237415
3970 C	8 021KA1G6	ESTs {IMAGE:681367}	681367 IMAGE Consortium	AA241826
3971 C	10 021KA1G6	ESTs {IMAGE:678633}	678633 IMAGE Consortium	AA238116
3972 C	12 021KA1G6	Kallikrein-3, plasma {IMAGE:678991}	678991 IMAGE Consortium	AA245959
3973 E	2 021KA1G6	Glial cell line derived neurotrophic factor family receptor alpha 1 {IMAGE:680894}	680894 IMAGE Consortium	AA250120
3974 E	4 021KA1G6	ESTs {IMAGE:681152}	681152 IMAGE Consortium	AA242567
3975 E	6 021KA1G6	Angiogenin {IMAGE:681634}	681634 IMAGE Consortium	AA237829
3976 E	8 021KA1G6	IMAGE EST {IMAGE:679075}	679075 IMAGE Consortium	AA244641
3977 E	10 021KA1G6	ESTs {IMAGE:679209}	679209 IMAGE Consortium	AA245985
3978 E	12 021KA1G6	Mus musculus dihydroorotate dehydrogenase mRNA, partial cds {IMAGE:692981}	692981 IMAGE Consortium	AA238399
3979 G	2 021KA1G6	ESTs, Moderately similar to RETINOL DEHYDROGENASE TYPE II [R.norvegicus] {IMAGE:681236}	681236 IMAGE Consortium	AA244820
3980 G	4 021KA1G6	ESTs, Weakly similar to cDNA EST EMBL:D75703 comes from this gene [C.elegans] {IMAGE:678992}	678992 IMAGE Consortium	AA244510
3981 G	6 021KA1G6	IMAGE EST {IMAGE:692439}	692439 IMAGE Consortium	AA238090

3982 G	8 021KA1G6	Fatty acid binding protein intestinal {IMAGE:679661}	679661 IMAGE Consortium	AA245078
3983 G	10 021KA1G6	ESTs, Weakly similar to BROWN PROTEIN [D.melanogaster] {IMAGE:680198}	680198 IMAGE Consortium	AA237916
3984 G	12 021KA1G6	ESTs, Moderately similar to LIVER 60 KD CARBOXYLESTERASE 2 [Oryctolagus cuniculus] {IMAGE:680448}	680448 IMAGE Consortium	AA237173
3985 A	2 021RA1G7	Esterase 1 {IMAGE:678863}	678863 IMAGE Consortium	AA249900
3986 A	4 021RA1G7	Proteasome (prosome, macropain) 28 subunit, 3 {IMAGE:679379}	679379 IMAGE Consortium	AA245903
3987 A	6 021RA1G7	ESTs, Highly similar to TRANSCRIPTION FACTOR BTF3 [Homo sapiens] {IMAGE:692936}	692936 IMAGE Consortium	AA238461
3988 A	8 021RA1G7	IMAGE EST {IMAGE:693024}	693024 IMAGE Consortium	AA253654
3989 A	10 021RA1G7	Mus musculus Masp-2 mRNA for mannose-binding lectin associated serine protease-2, complete cds {IMAGE:680104}	680104 IMAGE Consortium	AA244853
3990 A	12 021RA1G7	ESTs, Moderately similar to UDP-GLUCURONOSYLTRANSFERASE 2B5 PRECURSOR, MICROSOMAL [M.musculus] {IMAGE:680497}	680497 IMAGE Consortium	AA237191
3991 C	2 021RA1G7	ESTs, Moderately similar to STEROL 26-HYDROXYLASE MITOCHONDRIAL PRECURSOR [R.norvegicus] {IMAGE:680217}	680217 IMAGE Consortium	AA237744
3992 C	4 021RA1G7	IMAGE EST {IMAGE:679985}	679985 IMAGE Consortium	AA242287
3993 C	6 021RA1G7	ESTs, Highly similar to LZIP-1 and LZIP-2 [M.musculus] {IMAGE:680426}	680426 IMAGE Consortium	AA237169
3994 C	8 021RA1G7	IMAGE EST {IMAGE:694084}	694084 IMAGE Consortium	AA239050
3995 C	10 021RA1G7	Mus musculus thioredoxin mRNA, nuclear gene encoding mitochondrial protein, complete cds {IMAGE:681159}	681159 IMAGE Consortium	AA242573
3996 C	12 021RA1G7	ESTs {IMAGE:679163}	679163 IMAGE Consortium	AA245698
3997 E	2 021RA1G7	Annexin VII {IMAGE:681168}	681168 IMAGE Consortium	AA242060
3998 E	4 021RA1G7	ESTs, Highly similar to CYSTATHIONINE BETA-SYNTHASE [R.norvegicus] {IMAGE:678750}	678750 IMAGE Consortium	AA239480
3999 E	6 021RA1G7	IMAGE EST {IMAGE:678961}	678961 IMAGE Consortium	AA245654
4000 E	8 021RA1G7	Mus musculus D9 splice variant 2 mRNA, complete cds {IMAGE:679322}	679322 IMAGE Consortium	AA244657
4001 E	10 021RA1G7	IMAGE EST {IMAGE:692738}	692738 IMAGE Consortium	AA239881
4002 E	12 021RA1G7	Carbonic anhydrase-like sequence {IMAGE:692913}	692913 IMAGE Consortium	AA238417
4003 G	2 021RA1G7	ESTs, Highly similar to HYPOTHETICAL AMINOTRANSFERASE YJL060W [Saccharomyces cerevisiae] {IMAGE:679214}	679214 IMAGE Consortium	AA245984
4004 G	4 021RA1G7	IMAGE EST {IMAGE:679920}	679920 IMAGE Consortium	AA237592
4005 G	6 021RA1G7	IMAGE EST {IMAGE:692656}	692656 IMAGE Consortium	AA239841
4006 G	8 021RA1G7	DNA primase, p58 subunit {IMAGE:680147}	680147 IMAGE Consortium	AA237894
4007 G	10 021RA1G7	ESTs {IMAGE:693648}	693648 IMAGE Consortium	AA266660
4008 G	12 021RA1G7	IMAGE EST {IMAGE:693478}	693478 IMAGE Consortium	AA243947
4009 A	2 021YA1G8	ESTs, Highly similar to ARGININOSUCCINATE LYASE	692781 IMAGE Consortium	AA239933

		[Homo sapiens] {IMAGE:692781}		
4010 A	4 021YA1G8	Methylmalonyl coenzyme A mutase {IMAGE:693139}	693139 IMAGE Consortium	AA250181
4011 A	6 021YA1G8	ESTs {IMAGE:680435}	680435 IMAGE Consortium	AA237224
4012 A	8 021YA1G8	IMAGE EST {IMAGE:694065}	694065 IMAGE Consortium	AA243954
4013 A	10 021YA1G8	IMAGE EST {IMAGE:693837}	693837 IMAGE Consortium	AA238435
4014 A	12 021YA1G8	IMAGE EST {IMAGE:694929}	694929 IMAGE Consortium	AA218231
4015 C	2 021YA1G8	ESTs, Weakly similar to F35G2.2 [C.elegans] {IMAGE:680103}	680103 IMAGE Consortium	AA244842
4016 C	4 021YA1G8	IMAGE EST {IMAGE:680806}	680806 IMAGE Consortium	AA242626
4017 C	6 021YA1G8	ESTs, Highly similar to carnitine/acylcarnitine carrier protein [R.norvegicus] {IMAGE:699181}	699181 IMAGE Consortium	AA245413
4018 C	8 021YA1G8	Mus musculus suppressor of cytokine signalling-2 (SOCS-2) mRNA, complete cds {IMAGE:699428}	699428 IMAGE Consortium	AA245505
4019 C	10 021YA1G8	ESTs {IMAGE:696542}	696542 IMAGE Consortium	AA239742
4020 C	12 021YA1G8	ESTs {IMAGE:697191}	697191 IMAGE Consortium	AA221819
4021 E	2 021YA1G8	Mus musculus Crk-associated substrate (Cas-b) mRNA, complete cds {IMAGE:696710}	696710 IMAGE Consortium	AA240272
4022 E	4 021YA1G8	Protein phosphatase 1B, magnesium dependent, beta isoform {IMAGE:717394}	717394 IMAGE Consortium	AA404189
4023 E	6 021YA1G8	Inositol 1,4,5-triphosphate receptor 1 {IMAGE:697571}	697571 IMAGE Consortium	AA230925
4024 E	8 021YA1G8	Low density lipoprotein receptor related protein, associated protein 1 {IMAGE:697460}	697460 IMAGE Consortium	AA253890
4025 E	10 021YA1G8	Erythropoietin receptor {IMAGE:718293}	718293 IMAGE Consortium	AA261101
4026 E	12 021YA1G8	ESTs, Highly similar to ATP SYNTHASE LIPID-BINDING PROTEIN P2 PRECURSOR [Rattus norvegicus] {IMAGE:698109}	698109 IMAGE Consortium	AA239148
4027 G	2 021YA1G8	ESTs {IMAGE:717835}	717835 IMAGE Consortium	AA250361
4028 G	4 021YA1G8	Tumor necrosis factor super family 3-like {IMAGE:697864}	697864 IMAGE Consortium	AA238752
4029 G	6 021YA1G8	ESTs {IMAGE:698119}	698119 IMAGE Consortium	AA239367
4030 G	8 021YA1G8	Elastin {IMAGE:698181}	698181 IMAGE Consortium	AA239171
4031 G	10 021YA1G8	Biliary glycoprotein 1 {IMAGE:699461}	699461 IMAGE Consortium	AA245546
4032 G	12 021YA1G8	ESTs, Highly similar to HYPOTHETICAL 17.0 KD PROTEIN F56D1.3 IN CHROMOSOME II [Caenorhabditis elegans] {IMAGE:699422}	699422 IMAGE Consortium	AA245451
4033 A	2 0215A1G9	ESTs {IMAGE:697786}	697786 IMAGE Consortium	AA239093
4034 A	4 0215A1G9	IMAGE EST {IMAGE:697912}	697912 IMAGE Consortium	AA239100
4035 A	6 0215A1G9	IMAGE EST {IMAGE:718351}	718351 IMAGE Consortium	AA267699
4036 A	8 0215A1G9	ESTs {IMAGE:718474}	718474 IMAGE Consortium	AA266382
4037 A	10 0215A1G9	CALPONIN H2, SMOOTH MUSCLE {IMAGE:695687}	695687 IMAGE Consortium	AA220193
4038 A	12 0215A1G9	ESTs, Weakly similar to CYSTEINE-STRING PROTEIN [D.melanogaster] {IMAGE:716933}	716933 IMAGE Consortium	AA266043
4039 C	2 0215A1G9	IMAGE EST {IMAGE:716427}	716427 IMAGE Consortium	AA265354
4040 C	4 0215A1G9	ESTs, Highly similar to TXBP151 [H.sapiens] {IMAGE:717005}	717005 IMAGE Consortium	AA266531



4041 C	6 0215A1G9	Ankyrin 3, epithelial {IMAGE:717135}	717135 IMAGE Consortium	AA266234
4042 C	8 0215A1G9	IMAGE EST {IMAGE:697155}	697155 IMAGE Consortium	AA221680
4043 C	10 0215A1G9	IMAGE EST {IMAGE:717722}	717722 IMAGE Consortium	AA250453
4044 C	12 0215A1G9	ESTs, Moderately similar to putative Rab5-interacting protein {clone L1-94} [H.sapiens] {IMAGE:697407}	697407 IMAGE Consortium	AA244814
4045 E	2 0215A1G9	Mus musculus ribosomal protein L8 (RPL8) mRNA, complete cds {IMAGE:716896}	716896 IMAGE Consortium	AA266197
4046 E	4 0215A1G9	ESTs {IMAGE:697180}	697180 IMAGE Consortium	AA221743
4047 E	6 0215A1G9	ESTs, Highly similar to p20-CGGBP [H.sapiens] {IMAGE:697512}	697512 IMAGE Consortium	AA250640
4048 E	8 0215A1G9	Opioid receptor, sigma 1 {IMAGE:697551}	697551 IMAGE Consortium	AA230920
4049 E	10 0215A1G9	ESTs, Highly similar to HYPOTHETICAL 24.1 KD PROTEIN ON CHROMOSOME 1 [M.musculus] {IMAGE:718326}	718326 IMAGE Consortium	AA261092
4050 E	12 0215A1G9	IMAGE EST {IMAGE:718393}	718393 IMAGE Consortium	AA267730
4051 G	2 0215A1G9	ESTs {IMAGE:697225}	697225 IMAGE Consortium	AA242478
4052 G	4 0215A1G9	Mus musculus non-muscle myosin light chain 3 (MLC3nm) mRNA, partial cds {IMAGE:697569}	697569 IMAGE Consortium	AA230924
4053 G	6 0215A1G9	IMAGE EST {IMAGE:717991}	717991 IMAGE Consortium	AA261540
4054 G	8 0215A1G9	Mus musculus MEK binding partner 1 (Mp1) mRNA, complete cds {IMAGE:718194}	718194 IMAGE Consortium	AA266352
4055 G	10 0215A1G9	Nidogen {IMAGE:698175}	698175 IMAGE Consortium	AA239404
4056 G	12 0215A1G9	ESTs {IMAGE:718715}	718715 IMAGE Consortium	AA265789
4057 A	2 021PA1GA	ESTs {IMAGE:720736}	720736 IMAGE Consortium	AA267416
4058 A	4 021PA1GA	Gene trap ROSA 26 antisense {IMAGE:720812}	720812 IMAGE Consortium	AA267363
4059 A	6 021PA1GA	ESTs {IMAGE:721040}	721040 IMAGE Consortium	AA267373
4060 A	8 021PA1GA	Peroxisome proliferator activator receptor gamma {IMAGE:721051}	721051 IMAGE Consortium	AA267605
4061 A	10 021PA1GA	ESTs {IMAGE:734086}	734086 IMAGE Consortium	AA272115
4062 A	12 021PA1GA	ESTs, Weakly similar to HYPOTHETICAL 16.1 KD PROTEIN IN SEC17-QCR1 INTERGENIC REGION [Saccharomyces cerevisiae] {IMAGE:721503}	721503 IMAGE Consortium	AA266817
4063 C	2 021PA1GA	IMAGE EST {IMAGE:723580}	723580 IMAGE Consortium	AA260909
4064 C	4 021PA1GA	ESTs {IMAGE:733532}	733532 IMAGE Consortium	AA261229
4065 C	6 021PA1GA	ESTs {IMAGE:733832}	733832 IMAGE Consortium	AA268154
4066 C	8 021PA1GA	House mouse; Musculus domesticus postnatal 0-day old brain mRNA for KAP3A, complete cds {IMAGE:734376}	734376 IMAGE Consortium	AA268683
4067 C	10 021PA1GA	ESTs, Highly similar to ALPHA-1,6-MANNOSYL-GLYCOPROTEIN BETA-1,2-N-ACETYLGLUCOSAMINYLTRANSFERASE [Rattus norvegicus] {IMAGE:721625}	721625 IMAGE Consortium	AA260868
4068 C	12 021PA1GA	IMAGE EST {IMAGE:721996}	721996 IMAGE Consortium	AA268451
4069 E	2 021PA1GA	ESTs {IMAGE:733594}	733594 IMAGE Consortium	AA259986
4070 E	4 021PA1GA	ESTs, Moderately similar to developmentally regulated protein [R.norvegicus] {IMAGE:733557}	733557 IMAGE Consortium	AA259357

4071 E	6 021PA1GA	Calpain 5 {IMAGE:733856}	733856 IMAGE Consortium	AA268162
4072 E	8 021PA1GA	IMAGE EST {IMAGE:734305}	734305 IMAGE Consortium	AA268953
4073 E	10 021PA1GA	House mouse; Musculus domesticus male brain mRNA for ARF2, complete cds {IMAGE:721482}	721482 IMAGE Consortium	AA266938
4074 E	12 021PA1GA	ESTs {IMAGE:722068}	722068 IMAGE Consortium	AA268529
4075 G	2 021PA1GA	IMAGE EST {IMAGE:720952}	720952 IMAGE Consortium	AA266856
4076 G	4 021PA1GA	RAB11B, member RAS oncogene family {IMAGE:733701}	733701 IMAGE Consortium	AA260075
4077 G	6 021PA1GA	IMAGE EST {IMAGE:733746}	733746 IMAGE Consortium	AA268333
4078 G	8 021PA1GA	IMAGE EST {IMAGE:719075}	719075 IMAGE Consortium	AA266024
4079 G	10 021PA1GA	ESTs, Weakly similar to retinal-specific heterotrimeric GTP-binding protein beta subunit, G beta5L [M.musculus] {IMAGE:718989}	718989 IMAGE Consortium	AA265966
4080 G	12 021PA1GA	Dlk1-like homolog (Drosophila) {IMAGE:722057}	722057 IMAGE Consortium	AA268477
4081 A	2 021WA1GB	ESTs {IMAGE:718218}	718218 IMAGE Consortium	AA266360
4082 A	4 021WA1GB	ESTs, Highly similar to GLYCOGEN SYNTHASE KINASE-3 ALPHA [Rattus norvegicus] {IMAGE:718568}	718568 IMAGE Consortium	AA267750
4083 A	6 021WA1GB	Dlk1-like homolog (Drosophila) {IMAGE:699417}	699417 IMAGE Consortium	AA245452
4084 A	8 021WA1GB	Surfeit gene 6 {IMAGE:699378}	699378 IMAGE Consortium	AA245490
4085 A	10 021WA1GB	IMAGE EST {IMAGE:716768}	716768 IMAGE Consortium	AA266156
4086 A	12 021WA1GB	IMAGE EST {IMAGE:696469}	696469 IMAGE Consortium	AA239667
4087 C	2 021WA1GB	IMAGE EST {IMAGE:696517}	696517 IMAGE Consortium	AA239721
4088 C	4 021WA1GB	IMAGE EST {IMAGE:717845}	717845 IMAGE Consortium	AA250548
4089 C	6 021WA1GB	ESTs {IMAGE:718033}	718033 IMAGE Consortium	AA261553
4090 C	8 021WA1GB	Zinc finger protein 147 {IMAGE:718148}	718148 IMAGE Consortium	AA266342
4091 C	10 021WA1GB	Notch gene homolog 1, (Drosophila) {IMAGE:718688}	718688 IMAGE Consortium	AA265694
4092 C	12 021WA1GB	Mouse calcineurin catalytic subunit mRNA, complete cds {IMAGE:699236}	699236 IMAGE Consortium	AA245461
4093 E	2 021WA1GB	IMAGE EST {IMAGE:718670}	718670 IMAGE Consortium	AA265562
4094 E	4 021WA1GB	IMAGE EST {IMAGE:718806}	718806 IMAGE Consortium	AA265845
4095 E	6 021WA1GB	M.musculus mRNA for F25B3.3 kinase like protein from C.elegans {IMAGE:718866}	718866 IMAGE Consortium	AA265803
4096 E	8 021WA1GB	ESTs {IMAGE:719142}	719142 IMAGE Consortium	AA267310
4097 E	10 021WA1GB	IMAGE EST {IMAGE:719390}	719390 IMAGE Consortium	AA255102
4098 E	12 021WA1GB	ESTs {IMAGE:719691}	719691 IMAGE Consortium	AA254586
4099 G	2 021WA1GB	IMAGE EST {IMAGE:720716}	720716 IMAGE Consortium	AA267505
4100 G	4 021WA1GB	ESTs {IMAGE:721397}	721397 IMAGE Consortium	AA267286
4101 G	6 021WA1GB	X-linked B-cell surface antigen 3b {IMAGE:721409}	721409 IMAGE Consortium	AA267141
4102 G	8 021WA1GB	IMAGE EST {IMAGE:734062}	734062 IMAGE Consortium	AA272107
4103 G	10 021WA1GB	Platelet derived growth factor receptor, alpha polypeptide {IMAGE:721790}	721790 IMAGE Consortium	AA260832
4104 G	12 021WA1GB	ESTs {IMAGE:721970}	721970 IMAGE Consortium	AA268402
4105 A	2 0213A1GC	IMAGE EST {IMAGE:720886}	720886 IMAGE Consortium	AA267064
4106 A	4 0213A1GC	ESTs {IMAGE:721307}	721307 IMAGE Consortium	AA266972

4107 A	6 0213A1GC	ESTs, Moderately similar to Rabin3 [R.norvegicus] {IMAGE:733914}	733914 IMAGE Consortium	AA268382
4108 A	8 0213A1GC	ESTs, Weakly similar to glycoprotein specific UDP-glucuronyltransferase [R.norvegicus] {IMAGE:734282}	734282 IMAGE Consortium	AA268927
4109 A	10 0213A1GC	ESTs, Weakly similar to weak similarity to rat TEGT protein [C.elegans] {IMAGE:721531}	721531 IMAGE Consortium	AA267658
4110 A	12 0213A1GC	CD59 antigen {IMAGE:721831}	721831 IMAGE Consortium	AA267720
4111 C	2 0213A1GC	ESTs {IMAGE:733752}	733752 IMAGE Consortium	AA272913
4112 C	4 0213A1GC	IMAGE EST {IMAGE:733738}	733738 IMAGE Consortium	AA272878
4113 C	6 0213A1GC	ESTs, Highly similar to NADH-CYTOCHROME B5 REDUCTASE [Homo sapiens] {IMAGE:733988}	733988 IMAGE Consortium	AA268398
4114 C	8 0213A1GC	Mus musculus mRNA for ubiquitin conjugating enzyme {IMAGE:721747}	721747 IMAGE Consortium	AA260747
4115 C	10 0213A1GC	Seven in absentia 1A {IMAGE:721928}	721928 IMAGE Consortium	AA267965
4116 C	12 0213A1GC	IMAGE EST {IMAGE:722332}	722332 IMAGE Consortium	AA254238
4117 E	2 0213A1GC	ESTs {IMAGE:721334}	721334 IMAGE Consortium	AA267109
4118 E	4 0213A1GC	IMAGE EST {IMAGE:721537}	721537 IMAGE Consortium	AA266966
4119 E	6 0213A1GC	ESTs {IMAGE:721532}	721532 IMAGE Consortium	AA267655
4120 E	8 0213A1GC	IMAGE EST {IMAGE:722000}	722000 IMAGE Consortium	AA268453
4121 E	10 0213A1GC	ESTs, Weakly similar to Similarity to Yeast D-lactate dehydrogenase [C.elegans] {IMAGE:719593}	719593 IMAGE Consortium	AA255086
4122 E	12 0213A1GC	Mus musculus mRNA for very-long-chain acyl-CoA synthetase related protein (VLACSR) {IMAGE:719777}	719777 IMAGE Consortium	AA254935
4123 G	2 0213A1GC	ESTs {IMAGE:721700}	721700 IMAGE Consortium	AA260887
4124 G	4 0213A1GC	IMAGE EST {IMAGE:721929}	721929 IMAGE Consortium	AA267975
4125 G	6 0213A1GC	Nuclear transcription factor-Y alpha {IMAGE:719809}	719809 IMAGE Consortium	AA261057
4126 G	8 0213A1GC	ESTs {IMAGE:719849}	719849 IMAGE Consortium	AA260621
4127 G	10 0213A1GC	IMAGE EST {IMAGE:722566}	722566 IMAGE Consortium	AA254159
4128 G	12 0213A1GC	IMAGE EST {IMAGE:722582}	722582 IMAGE Consortium	AA260573
4129 A	2 021AA1GD	IMAGE EST {IMAGE:719522}	719522 IMAGE Consortium	AA254777
4130 A	4 021AA1GD	ESTs, Highly similar to GTP-BINDING PROTEIN SARA [Mus musculus] {IMAGE:719779}	719779 IMAGE Consortium	AA254936
4131 A	6 021AA1GD	Mus musculus Ankhzn mRNA, complete cds {IMAGE:720035}	720035 IMAGE Consortium	AA261035
4132 A	8 021AA1GD	ESTs {IMAGE:723127}	723127 IMAGE Consortium	AA267828
4133 A	10 021AA1GD	Mus musculus putative G protein-coupled receptor TDAG8 (TDAG8) mRNA, complete cds {IMAGE:723208}	723208 IMAGE Consortium	AA254500
4134 A	12 021AA1GD	ESTs {IMAGE:733685}	733685 IMAGE Consortium	AA259510
4135 C	2 021AA1GD	ESTs {IMAGE:722412}	722412 IMAGE Consortium	AA254265
4136 C	4 021AA1GD	IMAGE EST {IMAGE:722351}	722351 IMAGE Consortium	AA254115
4137 C	6 021AA1GD	IMAGE EST {IMAGE:723271}	723271 IMAGE Consortium	AA267787
4138 C	8 021AA1GD	IMAGE EST {IMAGE:720942}	720942 IMAGE Consortium	AA267577
4139 C	10 021AA1GD	ESTs {IMAGE:733337}	733337 IMAGE Consortium	AA259576

4140 C	12 021AA1GD	Faciogenital dysplasia homolog {IMAGE:733780}	733780 IMAGE Consortium	AA272942
4141 E	2 021AA1GD	CD37 antigen {IMAGE:746381}	746381 IMAGE Consortium	AA259959
4142 E	4 021AA1GD	ESTs {IMAGE:735903}	735903 IMAGE Consortium	AA269614
4143 E	6 021AA1GD	IMAGE EST {IMAGE:736250}	736250 IMAGE Consortium	AA270415
4144 E	8 021AA1GD	ESTs, Moderately similar to PROTEIN PHOSPHATASE PP2A, 65 KD REGULATORY SUBUNIT, BETA ISOFORM [Homo sapiens] {IMAGE:736162}	736162 IMAGE Consortium	AA270019
4145 E	10 021AA1GD	IMAGE EST {IMAGE:747376}	747376 IMAGE Consortium	AA272806
4146 E	12 021AA1GD	ESTs {IMAGE:747332}	747332 IMAGE Consortium	AA274881
4147 G	2 021AA1GD	Mus musculus mRNA encoding lysine-ketoglutarate reductase/saccharopine dehydrogenase {IMAGE:738285}	738285 IMAGE Consortium	AA271121
4148 G	4 021AA1GD	ESTs {IMAGE:735526}	735526 IMAGE Consortium	AA269842
4149 G	6 021AA1GD	ESTs {IMAGE:747105}	747105 IMAGE Consortium	AA274978
4150 G	8 021AA1GD	M.musculus mRNA for high mobility group 2 protein {IMAGE:747136}	747136 IMAGE Consortium	AA275111
4151 G	10 021AA1GD	ESTs {IMAGE:747377}	747377 IMAGE Consortium	AA272810
4152 G	12 021AA1GD	ESTs {IMAGE:737031}	737031 IMAGE Consortium	AA272049
4153 A	2 021HA1GE	Mannose binding lectin, liver (A) {IMAGE:737373}	737373 IMAGE Consortium	AA277810
4154 A	4 021HA1GE	ESTs, Weakly similar to cytochrome P-450 [M.musculus] {IMAGE:737648}	737648 IMAGE Consortium	AA277565
4155 A	6 021HA1GE	IMAGE EST {IMAGE:737802}	737802 IMAGE Consortium	AA277325
4156 A	8 021HA1GE	IMAGE EST {IMAGE:735576}	735576 IMAGE Consortium	AA270037
4157 A	10 021HA1GE	ESTs {IMAGE:735564}	735564 IMAGE Consortium	AA270329
4158 A	12 021HA1GE	IMAGE EST {IMAGE:735500}	735500 IMAGE Consortium	AA269710
4159 C	2 021HA1GE	ESTs, Weakly similar to transcriptional repressor, AEBP1 protein [M.musculus] {IMAGE:736854}	736854 IMAGE Consortium	AA271275
4160 C	4 021HA1GE	ESTs {IMAGE:737532}	737532 IMAGE Consortium	AA277718
4161 C	6 021HA1GE	IMAGE EST {IMAGE:734319}	734319 IMAGE Consortium	AA268692
4162 C	8 021HA1GE	IMAGE EST {IMAGE:734651}	734651 IMAGE Consortium	AA272717
4163 C	10 021HA1GE	Carnitine palmitoyltransferase 1, liver {IMAGE:737898}	737898 IMAGE Consortium	AA277495
4164 C	12 021HA1GE	ESTs {IMAGE:735413}	735413 IMAGE Consortium	AA272826
4165 E	2 021HA1GE	ESTs {IMAGE:747211}	747211 IMAGE Consortium	AA260494
4166 E	4 021HA1GE	Solute carrier family 22, member 1 {IMAGE:747431}	747431 IMAGE Consortium	AA272847
4167 E	6 021HA1GE	IMAGE EST {IMAGE:747656}	747656 IMAGE Consortium	AA268817
4168 E	8 021HA1GE	IMAGE EST {IMAGE:734367}	734367 IMAGE Consortium	AA268708
4169 E	10 021HA1GE	COPROPORPHYRINOGEN III OXIDASE PRECURSOR {IMAGE:734795}	734795 IMAGE Consortium	AA259342
4170 E	12 021HA1GE	H-2 CLASS II HISTOCOMPATIBILITY ANTIGEN, I-A BETA CHAIN PRECURSOR {IMAGE:737803}	737803 IMAGE Consortium	AA277329
4171 G	2 021HA1GE	ESTs, Weakly similar to HYPOTHETICAL PROTEIN F44G4.1 IN CHROMOSOME II [C.elegans] {IMAGE:736660}	736660 IMAGE Consortium	AA270506
4172 G	4 021HA1GE	ESTs, Weakly similar to D1007.5 [C.elegans] {IMAGE:736530}	736530 IMAGE Consortium	AA270737

4173 G	6 021HA1GE	ESTs {IMAGE:736999}	736999 IMAGE Consortium	AA271883
4174 G	8 021HA1GE	Mus musculus mRNA for GM3 synthase, complete cds {IMAGE:748275}	748275 IMAGE Consortium	AA274576
4175 G	10 021HA1GE	Mus musculus WSB-2 mRNA, complete cds {IMAGE:735057}	735057 IMAGE Consortium	AA259925
4176 G	12 021HA1GE	Platelet derived growth factor receptor, alpha polypeptide {IMAGE:735605}	735605 IMAGE Consortium	AA270169
4177 A	2 021OA1GF	ESTs {IMAGE:735401}	735401 IMAGE Consortium	AA269761
4178 A	4 021OA1GF	ESTs {IMAGE:736379}	736379 IMAGE Consortium	AA270649
4179 A	6 021OA1GF	Mus musculus aquaporin-8 mRNA, complete cds {IMAGE:747401}	747401 IMAGE Consortium	AA272836
4180 A	8 021OA1GF	ESTs {IMAGE:747524}	747524 IMAGE Consortium	AA268444
4181 A	10 021OA1GF	ESTs, Weakly similar to SKAP55 [H.sapiens] {IMAGE:747589}	747589 IMAGE Consortium	AA268862
4182 A	12 021OA1GF	ESTs, Highly similar to CYSTATHIONINE BETA-SYNTHASE [R.norvegicus] {IMAGE:737490}	737490 IMAGE Consortium	AA277347
4183 C	2 021OA1GF	Mus musculus maf10 (mAF10) mRNA, complete cds {IMAGE:736069}	736069 IMAGE Consortium	AA270281
4184 C	4 021OA1GF	Retinol dehydrogenase type 5 {IMAGE:747449}	747449 IMAGE Consortium	AA268423
4185 C	6 021OA1GF	IMAGE EST {IMAGE:747189}	747189 IMAGE Consortium	AA260299
4186 C	8 021OA1GF	Mouse mRNA for 8-oxo-dGTPase, complete cds {IMAGE:737058}	737058 IMAGE Consortium	AA271997
4187 C	10 021OA1GF	ESTs, Moderately similar to ALCOHOL SULFOTRANSFERASE 2 [M.musculus] {IMAGE:737384}	737384 IMAGE Consortium	AA277580
4188 C	12 021OA1GF	Transforming growth factor, beta induced, 68 kDa {IMAGE:734101}	734101 IMAGE Consortium	AA268592
4189 E	2 021OA1GF	IMAGE EST {IMAGE:736299}	736299 IMAGE Consortium	AA270608
4190 E	4 021OA1GF	Mus musculus eIF-1A (eIF-1A) mRNA, complete cds {IMAGE:747322}	747322 IMAGE Consortium	AA274946
4191 E	6 021OA1GF	ESTs, Weakly similar to HYPOTHETICAL 91.2 KD PROTEIN IN RPS7A-SCH9 INTERGENIC REGION [Saccharomyces cerevisiae] {IMAGE:747381}	747381 IMAGE Consortium	AA272812
4192 E	8 021OA1GF	ESTs, Highly similar to 40S RIBOSOMAL PROTEIN S25 [Homo sapiens; Rattus norvegicus] {IMAGE:736987}	736987 IMAGE Consortium	AA272124
4193 E	10 021OA1GF	Mus musculus (clone C7/B9) S-adenosyl homocysteine hydrolase (ahcy) mRNA, complete cds {IMAGE:737180}	737180 IMAGE Consortium	AA272567
4194 E	12 021OA1GF	IMAGE EST {IMAGE:737669}	737669 IMAGE Consortium	AA277646
4195 G	2 021OA1GF	ESTs, Moderately similar to betaine:homocysteine methyltransferase [H.sapiens] {IMAGE:737041}	737041 IMAGE Consortium	AA272010
4196 G	4 021OA1GF	IMAGE EST {IMAGE:737084}	737084 IMAGE Consortium	AA272501
4197 G	6 021OA1GF	ESTs {IMAGE:747960}	747960 IMAGE Consortium	AA260958
4198 G	8 021OA1GF	Mus musculus secretory carrier membrane protein (SCAMP3) mRNA, complete cds {IMAGE:737492}	737492 IMAGE Consortium	AA277348
4199 G	10 021OA1GF	ESTs {IMAGE:735147}	735147 IMAGE Consortium	AA259439
4200 G	12 021OA1GF	IMAGE EST {IMAGE:734967}	734967 IMAGE Consortium	AA259967

4201 A	2	021VA1GG	ESTs {IMAGE:737425}	737425	IMAGE Consortium	AA277183
4202 A	4	021VA1GG	Fatty acid amide hydrolase {IMAGE:748132}	748132	IMAGE Consortium	AA260227
4203 A	6	021VA1GG	ESTs, Weakly similar to weakly similar to serine/threonine protein kinase [C.elegans] {IMAGE:734850}	734850	IMAGE Consortium	AA259352
4204 A	8	021VA1GG	Mouse mRNA for elongation factor 1-alpha (EF 1-alpha) {IMAGE:735225}	735225	IMAGE Consortium	AA259551
4205 A	10	021VA1GG	ESTs {IMAGE:735090}	735090	IMAGE Consortium	AA259717
4206 A	12	021VA1GG	IMAGE EST {IMAGE:738165}	738165	IMAGE Consortium	AA271054
4207 C	2	021VA1GG	ESTs {IMAGE:748639}	748639	IMAGE Consortium	AA269741
4208 C	4	021VA1GG	ESTs {IMAGE:751298}	751298	IMAGE Consortium	AA290369
4209 C	6	021VA1GG	ESTs {IMAGE:751399}	751399	IMAGE Consortium	AA395963
4210 C	8	021VA1GG	ESTs {IMAGE:749382}	749382	IMAGE Consortium	AA289526
4211 C	10	021VA1GG	ESTs {IMAGE:752233}	752233	IMAGE Consortium	AA289093
4212 C	12	021VA1GG	IMAGE EST {IMAGE:752464}	752464	IMAGE Consortium	AA289513
4213 E	2	021VA1GG	ESTs {IMAGE:748881}	748881	IMAGE Consortium	AA289889
4214 E	4	021VA1GG	ESTs {IMAGE:748913}	748913	IMAGE Consortium	AA288092
4215 E	6	021VA1GG	IMAGE EST {IMAGE:749353}	749353	IMAGE Consortium	AA289472
4216 E	8	021VA1GG	ESTs, Moderately similar to HYPOTHETICAL 45.3 KD PROTEIN ZK370.5 IN CHROMOSOME III [Caenorhabditis elegans] {IMAGE:752497}	752497	IMAGE Consortium	AA289387
4217 E	10	021VA1GG	ESTs {IMAGE:752201}	752201	IMAGE Consortium	AA289075
4218 E	12	021VA1GG	ESTs, Weakly similar to ENV POLYPROTEIN [Mouse mammary tumor virus (strain br6)] {IMAGE:749513}	749513	IMAGE Consortium	AA288274
4219 G	2	021VA1GG	IMAGE EST {IMAGE:751687}	751687	IMAGE Consortium	AA396025
4220 G	4	021VA1GG	Inositol polyphosphate-5-phosphatase, 75 kDa {IMAGE:749425}	749425	IMAGE Consortium	AA289577
4221 G	6	021VA1GG	ESTs {IMAGE:749144}	749144	IMAGE Consortium	AA290557
4222 G	8	021VA1GG	IMAGE EST {IMAGE:749175}	749175	IMAGE Consortium	AA289599
4223 G	10	021VA1GG	IMAGE EST {IMAGE:749569}	749569	IMAGE Consortium	AA388690
4224 G	12	021VA1GG	IMAGE EST {IMAGE:749778}	749778	IMAGE Consortium	AA288612
4225 A	2	0212A1GH	IMAGE EST {IMAGE:749066}	749066	IMAGE Consortium	AA290517
4226 A	4	0212A1GH	ESTs {IMAGE:749073}	749073	IMAGE Consortium	AA290526
4227 A	6	0212A1GH	ESTs, Weakly similar to ORF YKR086w [S.cerevisiae] {IMAGE:752140}	752140	IMAGE Consortium	AA289001
4228 A	8	0212A1GH	ESTs {IMAGE:749507}	749507	IMAGE Consortium	AA290511
4229 A	10	0212A1GH	Abl-interactor 1 {IMAGE:749467}	749467	IMAGE Consortium	AA288264
4230 A	12	0212A1GH	ESTs {IMAGE:749934}	749934	IMAGE Consortium	AA388607
4231 C	2	0212A1GH	CD14 antigen {IMAGE:751756}	751756	IMAGE Consortium	AA396117
4232 C	4	0212A1GH	Prostaglandin E receptor EP1 subtype {IMAGE:751958}	751958	IMAGE Consortium	AA396341
4233 C	6	0212A1GH	ESTs {IMAGE:752308}	752308	IMAGE Consortium	AA289983
4234 C	8	0212A1GH	IMAGE EST {IMAGE:752156}	752156	IMAGE Consortium	AA289059
4235 C	10	0212A1GH	ESTs, Moderately similar to envelope polyprotein [M.musculus] {IMAGE:761667}	761667	IMAGE Consortium	AA271814

4236 C	12 0212A1GH	ESTs, Highly similar to NIFS-LIKE 54.5 KD PROTEIN [Saccharomyces cerevisiae] {IMAGE:750126}	750126 IMAGE Consortium	AA396372
4237 E	2 0212A1GH	LIM only 2 {IMAGE:752429}	752429 IMAGE Consortium	AA290294
4238 E	4 0212A1GH	ESTs {IMAGE:752276}	752276 IMAGE Consortium	AA290015
4239 E	6 0212A1GH	IG ALPHA CHAIN C REGION {IMAGE:749660}	749660 IMAGE Consortium	AA388939
4240 E	8 0212A1GH	Biliary glycoprotein 1 {IMAGE:762299}	762299 IMAGE Consortium	AA277736
4241 E	10 0212A1GH	IMAGE EST {IMAGE:750450}	750450 IMAGE Consortium	AA472882
4242 E	12 0212A1GH	Ubiquitously transcribed tetratricopeptide repeat gene, X Chromosome {IMAGE:750794}	750794 IMAGE Consortium	AA290145
4243 G	2 0212A1GH	Mus musculus Na <sup>+</sup> /H <sup>+</sup> exchanger (NHE-1) mRNA, complete cds {IMAGE:749780}	749780 IMAGE Consortium	AA288613
4244 G	4 0212A1GH	ESTs {IMAGE:749865}	749865 IMAGE Consortium	AA389152
4245 G	6 0212A1GH	IMAGE EST {IMAGE:749853}	749853 IMAGE Consortium	AA389257
4246 G	8 0212A1GH	ESTs, Highly similar to KIAA0228 protein [H.sapiens] {IMAGE:750627}	750627 IMAGE Consortium	AA290102
4247 G	10 0212A1GH	ESTs {IMAGE:751354}	751354 IMAGE Consortium	AA289439
4248 G	12 0212A1GH	IMAGE EST {IMAGE:751540}	751540 IMAGE Consortium	AA396037
4249 A	2 0219A1GI	ESTs, Highly similar to NUCLEAR TRANSPORT FACTOR 2 [Homo sapiens; Rattus norvegicus] {IMAGE:762516}	762516 IMAGE Consortium	AA271353
4250 A	4 0219A1GI	ESTs, Weakly similar to CARBONIC ANHYDRASE III [H.sapiens] {IMAGE:775836}	775836 IMAGE Consortium	AA276019
4251 A	6 0219A1GI	Mus musculus ubiquitin conjugating enzyme (ubc4) mRNA, complete cds {IMAGE:775795}	775795 IMAGE Consortium	AA276008
4252 A	8 0219A1GI	ESTs, Weakly similar to similar to sodium/phosphate transporter protein [C.elegans] {IMAGE:776148}	776148 IMAGE Consortium	AA276173
4253 A	10 0219A1GI	ESTs {IMAGE:764732}	764732 IMAGE Consortium	AA273704
4254 A	12 0219A1GI	ESTs, Moderately similar to HYPOTHETICAL 54.7 KD PROTEIN F37A4.1 IN CHROMOSOME III [Caenorhabditis elegans] {IMAGE:765045}	765045 IMAGE Consortium	AA275058
4255 C	2 0219A1GI	ESTs {IMAGE:764163}	764163 IMAGE Consortium	AA285845
4256 C	4 0219A1GI	IMAGE EST {IMAGE:764224}	764224 IMAGE Consortium	AA285703
4257 C	6 0219A1GI	Branched chain aminotransferase 2, mitochondrial {IMAGE:776036}	776036 IMAGE Consortium	AA286063
4258 C	8 0219A1GI	Retinol dehydrogenase type 5 {IMAGE:776350}	776350 IMAGE Consortium	AA275664
4259 C	10 0219A1GI	IMAGE EST {IMAGE:764998}	764998 IMAGE Consortium	AA273426
4260 C	12 0219A1GI	Interleukin 15 receptor, alpha chain {IMAGE:777103}	777103 IMAGE Consortium	AA276366
4261 E	2 0219A1GI	IMAGE EST {IMAGE:764272}	764272 IMAGE Consortium	AA285418
4262 E	4 0219A1GI	ESTs {IMAGE:764473}	764473 IMAGE Consortium	AA274099
4263 E	6 0219A1GI	IMAGE EST {IMAGE:764651}	764651 IMAGE Consortium	AA273175
4264 E	8 0219A1GI	M.musculus mRNA for histone H3.3A {IMAGE:776614}	776614 IMAGE Consortium	AA275906
4265 E	10 0219A1GI	Hydroxysteroid dehydrogenase-4, delta<5>-3-beta {IMAGE:777059}	777059 IMAGE Consortium	AA276254
4266 E	12 0219A1GI	IMAGE EST {IMAGE:777099}	777099 IMAGE Consortium	AA276364
4267 G	2 0219A1GI	ESTs {IMAGE:764212}	764212 IMAGE Consortium	AA285661

4268 G	4 0219A1GI	IMAGE EST {IMAGE:764344}	764344 IMAGE Consortium	AA277492
4269 G	6 0219A1GI	Cadherin 1 {IMAGE:776133}	776133 IMAGE Consortium	AA276110
4270 G	8 0219A1GI	ESTs {IMAGE:776447}	776447 IMAGE Consortium	AA278090
4271 G	10 0219A1GI	IMAGE EST {IMAGE:777036}	777036 IMAGE Consortium	AA276138
4272 G	12 0219A1GI	IMAGE EST {IMAGE:777171}	777171 IMAGE Consortium	AA276192
4273 A	2 021GA1GJ	ESTs {IMAGE:750538}	750538 IMAGE Consortium	AA472919
4274 A	4 021GA1GJ	ESTs, Highly similar to PUTATIVE ADENOSINE KINASE [Saccharomyces cerevisiae] {IMAGE:748517}	748517 IMAGE Consortium	AA272821
4275 A	6 021GA1GJ	Mus musculus retinoid X receptor interacting protein (RIP14-1No.6) mRNA, complete cds {IMAGE:748622}	748622 IMAGE Consortium	AA270183
4276 A	8 021GA1GJ	IMAGE EST {IMAGE:748390}	748390 IMAGE Consortium	AA268125
4277 A	10 021GA1GJ	ESTs, Highly similar to INTERFERON-INDUCIBLE PROTEIN [Rattus norvegicus] {IMAGE:748975}	748975 IMAGE Consortium	AA289992
4278 A	12 021GA1GJ	ESTs {IMAGE:751596}	751596 IMAGE Consortium	AA396050
4279 C	2 021GA1GJ	IMAGE EST {IMAGE:750228}	750228 IMAGE Consortium	AA390087
4280 C	4 021GA1GJ	Mus musculus ribosomal protein L21 mRNA, complete cds {IMAGE:748374}	748374 IMAGE Consortium	AA274667
4281 C	6 021GA1GJ	ESTs {IMAGE:748670}	748670 IMAGE Consortium	AA270197
4282 C	8 021GA1GJ	M.musculus mRNA for serine/threonine kinase {IMAGE:751216}	751216 IMAGE Consortium	AA288220
4283 C	10 021GA1GJ	TRANSCRIPTION FACTOR S-II-RELATED PROTEIN {IMAGE:748744}	748744 IMAGE Consortium	AA288128
4284 C	12 021GA1GJ	IMAGE EST {IMAGE:751530}	751530 IMAGE Consortium	AA389944
4285 E	2 021GA1GJ	IMAGE EST {IMAGE:750695}	750695 IMAGE Consortium	AA288248
4286 E	4 021GA1GJ	Mus musculus cytokine receptor-like molecule (EBI3) mRNA, complete cds {IMAGE:751312}	751312 IMAGE Consortium	AA289482
4287 E	6 021GA1GJ	IMAGE EST {IMAGE:751494}	751494 IMAGE Consortium	AA396001
4288 E	8 021GA1GJ	ESTs {IMAGE:749113}	749113 IMAGE Consortium	AA290548
4289 E	10 021GA1GJ	IMAGE EST {IMAGE:749099}	749099 IMAGE Consortium	AA290539
4290 E	12 021GA1GJ	IMAGE EST {IMAGE:749766}	749766 IMAGE Consortium	AA396408
4291 G	2 021GA1GJ	ESTs {IMAGE:748896}	748896 IMAGE Consortium	AA289893
4292 G	4 021GA1GJ	Mus musculus NET1 homolog (Net1) mRNA, complete cds {IMAGE:751710}	751710 IMAGE Consortium	AA396114
4293 G	6 021GA1GJ	IMAGE EST {IMAGE:751501}	751501 IMAGE Consortium	AA396009
4294 G	8 021GA1GJ	IMAGE EST {IMAGE:751389}	751389 IMAGE Consortium	AA396314
4295 G	10 021GA1GJ	Cyclin B1, related sequence 1 {IMAGE:751977}	751977 IMAGE Consortium	AA396324
4296 G	12 021GA1GJ	ESTs {IMAGE:752137}	752137 IMAGE Consortium	AA289003
4297 A	2 021NA1GK	IMAGE EST {IMAGE:763629}	763629 IMAGE Consortium	AA285584
4298 A	4 021NA1GK	ESTs {IMAGE:764227}	764227 IMAGE Consortium	AA285653
4299 A	6 021NA1GK	Cell division cycle 7-like 1 {IMAGE:764785}	764785 IMAGE Consortium	AA274194
4300 A	8 021NA1GK	ESTs, Weakly similar to weak similarity to rat TEGT protein [C.elegans] {IMAGE:764677}	764677 IMAGE Consortium	AA273209
4301 A	10 021NA1GK	Solute carrier family 2 (facilitated glucose transporter),	776543 IMAGE Consortium	AA275871



		member 2 {IMAGE:776543}		
4302 A	12 021NA1GK	IMAGE EST {IMAGE:765559}	765559 IMAGE Consortium	AA275141
4303 C	2 021NA1GK	IMAGE EST {IMAGE:764359}	764359 IMAGE Consortium	AA277466
4304 C	4 021NA1GK	IMAGE EST {IMAGE:764690}	764690 IMAGE Consortium	AA273211
4305 C	6 021NA1GK	IMAGE EST {IMAGE:764606}	764606 IMAGE Consortium	AA273858
4306 C	8 021NA1GK	ESTs {IMAGE:765182}	765182 IMAGE Consortium	AA273950
4307 C	10 021NA1GK	ESTs {IMAGE:777363}	777363 IMAGE Consortium	AA276938
4308 C	12 021NA1GK	ESTs {IMAGE:777321}	777321 IMAGE Consortium	AA276913
4309 E	2 021NA1GK	ATPase-like vacuolar proton channel {IMAGE:776055}	776055 IMAGE Consortium	AA276030
4310 E	4 021NA1GK	IMAGE EST {IMAGE:764727}	764727 IMAGE Consortium	AA273603
4311 E	6 021NA1GK	IMAGE EST {IMAGE:765015}	765015 IMAGE Consortium	AA273573
4312 E	8 021NA1GK	ESTs {IMAGE:765512}	765512 IMAGE Consortium	AA274737
4313 E	10 021NA1GK	ESTs {IMAGE:765332}	765332 IMAGE Consortium	AA275245
4314 E	12 021NA1GK	IMAGE EST {IMAGE:765774}	765774 IMAGE Consortium	AA275349
4315 G	2 021NA1GK	ESTs {IMAGE:765194}	765194 IMAGE Consortium	AA273249
4316 G	4 021NA1GK	ESTs {IMAGE:777139}	777139 IMAGE Consortium	AA276512
4317 G	6 021NA1GK	IMAGE EST {IMAGE:765703}	765703 IMAGE Consortium	AA277071
4318 G	8 021NA1GK	IMAGE EST {IMAGE:762549}	762549 IMAGE Consortium	AA271085
4319 G	10 021NA1GK	ESTs, Moderately similar to DNA-DIRECTED RNA POLYMERASE II 32 KD POLYPEPTIDE [ <i>Saccharomyces cerevisiae</i> ] {IMAGE:763048}	763048 IMAGE Consortium	AA271483
4320 G	12 021NA1GK	Uroporphyrinogen III synthase {IMAGE:763423}	763423 IMAGE Consortium	AA272198
4321 A	2 021UA1GL	ESTs {IMAGE:765261}	765261 IMAGE Consortium	AA273453
4322 A	4 021UA1GL	IMAGE EST {IMAGE:762237}	762237 IMAGE Consortium	AA277405
4323 A	6 021UA1GL	IMAGE EST {IMAGE:762655}	762655 IMAGE Consortium	AA271235
4324 A	8 021UA1GL	IMAGE EST {IMAGE:762599}	762599 IMAGE Consortium	AA271142
4325 A	10 021UA1GL	ESTs {IMAGE:763144}	763144 IMAGE Consortium	AA271147
4326 A	12 021UA1GL	ESTs {IMAGE:775596}	775596 IMAGE Consortium	AA275920
4327 C	2 021UA1GL	<i>M.musculus</i> mRNA for Pinin protein {IMAGE:765518}	765518 IMAGE Consortium	AA274739
4328 C	4 021UA1GL	ESTs, Weakly similar to ZINC FINGER PROTEIN 76 [ <i>Homo sapiens</i> ] {IMAGE:762142}	762142 IMAGE Consortium	AA277793
4329 C	6 021UA1GL	ESTs {IMAGE:762728}	762728 IMAGE Consortium	AA271551
4330 C	8 021UA1GL	Matrix metalloproteinase 2 {IMAGE:762791}	762791 IMAGE Consortium	AA271593
4331 C	10 021UA1GL	ESTs {IMAGE:763099}	763099 IMAGE Consortium	AA271329
4332 C	12 021UA1GL	ESTs, Weakly similar to F56D2.5 [ <i>C.elegans</i> ] {IMAGE:763376}	763376 IMAGE Consortium	AA272245
4333 E	2 021UA1GL	ESTs {IMAGE:777384}	777384 IMAGE Consortium	AA276540
4334 E	4 021UA1GL	IMAGE EST {IMAGE:777349}	777349 IMAGE Consortium	AA276928
4335 E	6 021UA1GL	ESTs {IMAGE:763202}	763202 IMAGE Consortium	AA286654
4336 E	8 021UA1GL	ESTs, Moderately similar to protein kinase C-binding protein Beta 15 [ <i>R.norvegicus</i> ] {IMAGE:763028}	763028 IMAGE Consortium	AA271411
4337 E	10 021UA1GL	ESTs, Highly similar to hnRNP G protein [ <i>H.sapiens</i> ] {IMAGE:763362}	763362 IMAGE Consortium	AA285673

4338 E	12 021UA1GL	IMAGE EST {IMAGE:763966}	763966 IMAGE Consortium	AA285518
4339 G	2 021UA1GL	ESTs {IMAGE:808736}	808736 IMAGE Consortium	AA467340
4340 G	4 021UA1GL	IMAGE EST {IMAGE:808891}	808891 IMAGE Consortium	AA465786
4341 G	6 021UA1GL	IMAGE EST {IMAGE:809011}	809011 IMAGE Consortium	AA466063
4342 G	8 021UA1GL	IMAGE EST {IMAGE:790197}	790197 IMAGE Consortium	AA388644
4343 G	10 021UA1GL	ESTs {IMAGE:818884}	818884 IMAGE Consortium	AA434883
4344 G	12 021UA1GL	Mus musculus vesicle-associated membrane protein cellubrevin mRNA, complete cds {IMAGE:818971}	818971 IMAGE Consortium	AA435160
4345 A	2 0211A1GM	IMAGE EST {IMAGE:817923}	817923 IMAGE Consortium	AA466517
4346 A	4 0211A1GM	ESTs {IMAGE:790569}	790569 IMAGE Consortium	AA290020
4347 A	6 0211A1GM	IMAGE EST {IMAGE:790674}	790674 IMAGE Consortium	AA388377
4348 A	8 0211A1GM	IMAGE EST {IMAGE:803275}	803275 IMAGE Consortium	AA396306
4349 A	10 0211A1GM	ESTs, Weakly similar to GRAVIN [Homo sapiens] {IMAGE:806940}	806940 IMAGE Consortium	AA403828
4350 A	12 0211A1GM	ESTs {IMAGE:807462}	807462 IMAGE Consortium	AA412944
4351 C	2 0211A1GM	IMAGE EST {IMAGE:820179}	820179 IMAGE Consortium	AA423218
4352 C	4 0211A1GM	ESTs, Moderately similar to p15 protein [H.sapiens] {IMAGE:803541}	803541 IMAGE Consortium	AA473349
4353 C	6 0211A1GM	ESTs, Weakly similar to CELL CYCLE REGULATION AND TRANSLATION INITIATION PROTEIN [Saccharomyces cerevisiae] {IMAGE:820518}	820518 IMAGE Consortium	AA423253
4354 C	8 0211A1GM	ESTs {IMAGE:831653}	831653 IMAGE Consortium	AA467013
4355 C	10 0211A1GM	IMAGE EST {IMAGE:777529}	777529 IMAGE Consortium	AA276682
4356 C	12 0211A1GM	ESTs {IMAGE:808780}	808780 IMAGE Consortium	AA467476
4357 E	2 0211A1GM	Glycoprotein 49 A {IMAGE:820498}	820498 IMAGE Consortium	AA423373
4358 E	4 0211A1GM	ESTs, Weakly similar to located at OATL1 [H.sapiens] {IMAGE:807817}	807817 IMAGE Consortium	AA432631
4359 E	6 0211A1GM	ESTs, Weakly similar to F25H9.7 [C.elegans] {IMAGE:807855}	807855 IMAGE Consortium	AA470218
4360 E	8 0211A1GM	IMAGE EST {IMAGE:777555}	777555 IMAGE Consortium	AA276691
4361 E	10 0211A1GM	ESTs {IMAGE:777823}	777823 IMAGE Consortium	AA286400
4362 E	12 0211A1GM	IMAGE EST {IMAGE:777589}	777589 IMAGE Consortium	AA276847
4363 G	2 0211A1GM	IMAGE EST {IMAGE:831691}	831691 IMAGE Consortium	AA466102
4364 G	4 0211A1GM	IMAGE EST {IMAGE:777560}	777560 IMAGE Consortium	AA276755
4365 G	6 0211A1GM	Mouse chromosome 6 BAC-284H12 (Research Genetics mouse BAC library) complete sequence {IMAGE:808651}	808651 IMAGE Consortium	AA467142
4366 G	8 0211A1GM	ESTs, Weakly similar to 60S RIBOSOMAL PROTEIN L30A [Saccharomyces cerevisiae] {IMAGE:809055}	809055 IMAGE Consortium	AA466052
4367 G	10 0211A1GM	ESTs, Highly similar to NADH-UBIQUINONE OXIDOREDUCTASE B15 SUBUNIT [Bos taurus] {IMAGE:818906}	818906 IMAGE Consortium	AA434897
4368 G	12 0211A1GM	Platelet derived growth factor receptor, alpha polypeptide {IMAGE:818704}	818704 IMAGE Consortium	AA466573
4369 A	2 0218A1GN	Mus musculus serine/threonine kinase (sak-a) mRNA,	777598 IMAGE Consortium	AA276998

		complete cds {IMAGE:777598}		
4370 A	4 0218A1GN	ESTs {IMAGE:818188}	818188 IMAGE Consortium	AA435275
4371 A	6 0218A1GN	ESTs, Weakly similar to KIAA0364 [H.sapiens] {IMAGE:818908}	818908 IMAGE Consortium	AA434898
4372 A	8 0218A1GN	ESTs {IMAGE:819021}	819021 IMAGE Consortium	AA435176
4373 A	10 0218A1GN	Mus musculus cytoplasmic protein Ndr1 (Ndr1) mRNA, complete cds {IMAGE:803416}	803416 IMAGE Consortium	AA473269
4374 A	12 0218A1GN	IMAGE EST {IMAGE:820195}	820195 IMAGE Consortium	AA423224
4375 C	2 0218A1GN	IMAGE EST {IMAGE:820320}	820320 IMAGE Consortium	AA423099
4376 C	4 0218A1GN	IMAGE EST {IMAGE:803488}	803488 IMAGE Consortium	AA473289
4377 C	6 0218A1GN	ESTs {IMAGE:806740}	806740 IMAGE Consortium	AA403565
4378 C	8 0218A1GN	Mast cell growth factor {IMAGE:806850}	806850 IMAGE Consortium	AA403846
4379 C	10 0218A1GN	IMAGE EST {IMAGE:807687}	807687 IMAGE Consortium	AA470257
4380 C	12 0218A1GN	ESTs {IMAGE:777237}	777237 IMAGE Consortium	AA276895
4381 E	2 0218A1GN	Mus musculus manic fringe precursor mRNA, complete cds {IMAGE:831674}	831674 IMAGE Consortium	AA467019
4382 E	4 0218A1GN	IMAGE EST {IMAGE:831735}	831735 IMAGE Consortium	AA444938
4383 E	6 0218A1GN	Solute carrier family 8 (sodium/calcium exchanger), member 1 {IMAGE:808226}	808226 IMAGE Consortium	AA433080
4384 E	8 0218A1GN	ESTs, Highly similar to hypothetical protein, 100K [R.norvegicus] {IMAGE:808777}	808777 IMAGE Consortium	AA467485
4385 E	10 0218A1GN	ESTs, Moderately similar to elastin like protein [D.melanogaster] {IMAGE:808594}	808594 IMAGE Consortium	AA467099
4386 E	12 0218A1GN	ESTs, Weakly similar to F42A6.6 [C.elegans] {IMAGE:809206}	809206 IMAGE Consortium	AA434709
4387 G	2 0218A1GN	ESTs {IMAGE:808658}	808658 IMAGE Consortium	AA467199
4388 G	4 0218A1GN	ESTs {IMAGE:808595}	808595 IMAGE Consortium	AA467108
4389 G	6 0218A1GN	ESTs {IMAGE:809255}	809255 IMAGE Consortium	AA445144
4390 G	8 0218A1GN	ESTs {IMAGE:818818}	818818 IMAGE Consortium	AA467406
4391 G	10 0218A1GN	ESTs {IMAGE:820154}	820154 IMAGE Consortium	AA423114
4392 G	12 0218A1GN	ESTs, Highly similar to RIBONUCLEASE PL3 PRECURSOR [Sus scrofa] {IMAGE:803251}	803251 IMAGE Consortium	AA396298
4393 A	2 021FA1GO	ESTs {IMAGE:777813}	777813 IMAGE Consortium	AA286507
4394 A	4 021FA1GO	ESTs, Highly similar to PINCH PROTEIN [Homo sapiens] {IMAGE:790449}	790449 IMAGE Consortium	AA289280
4395 A	6 021FA1GO	Mus musculus cell death activator CIDE-B (Cide-b) mRNA, complete cds {IMAGE:790532}	790532 IMAGE Consortium	AA290390
4396 A	8 021FA1GO	IMAGE EST {IMAGE:818779}	818779 IMAGE Consortium	AA467376
4397 A	10 021FA1GO	Mus musculus mRNA for eIF3 p66, complete cds {IMAGE:820355}	820355 IMAGE Consortium	AA423277
4398 A	12 021FA1GO	ESTs, Moderately similar to KIAA0429 [H.sapiens] {IMAGE:807447}	807447 IMAGE Consortium	AA412824
4399 C	2 021FA1GO			
4400 C	4 021FA1GO			

4401 C	6 021FA1GO			
4402 C	8 021FA1GO			
4403 C	10 021FA1GO			
4404 C	12 021FA1GO			
4405 E	2 021FA1GO			
4406 E	4 021FA1GO			
4407 E	6 021FA1GO			
4408 E	8 021FA1GO			
4409 E	10 021FA1GO			
4410 E	12 021FA1GO			
4411 G	2 021FA1GO			
4412 G	4 021FA1GO			
4413 G	6 021FA1GO			
4414 G	8 021FA1GO			
4415 G	10 021FA1GO			
4416 G	12 021FA1GO			
4417 A	2 021MA1GP	ESTs {IMAGE:523814}	523814 IMAGE Consortium	AA067146
4418 A	4 021MA1GP	IMAGE EST {IMAGE:524133}	524133 IMAGE Consortium	AA066133
4419 A	6 021MA1GP	ESTs, Highly similar to HYPOTHETICAL 44.2 KD PROTEIN IN SCO2-MRF1 INTERGENIC REGION [ <i>Saccharomyces cerevisiae</i> ] {IMAGE:524792}	524792 IMAGE Consortium	AA067828
4420 A	8 021MA1GP	IMAGE EST {IMAGE:408040}	408040 IMAGE Consortium	W85448
4421 A	10 021MA1GP	ESTs, Moderately similar to MHC Class I region proline rich protein [ <i>H.sapiens</i> ] {IMAGE:535955}	535955 IMAGE Consortium	AA073345
4422 A	12 021MA1GP	Apoptosis inhibitor 3 {IMAGE:550702}	550702 IMAGE Consortium	AA097958
4423 C	2 021MA1GP	IMAGE EST {IMAGE:522558}	522558 IMAGE Consortium	AA067659
4424 C	4 021MA1GP	IMAGE EST {IMAGE:523057}	523057 IMAGE Consortium	AA066452
4425 C	6 021MA1GP	Mus musculus coding region determinant binding protein mRNA, complete cds {IMAGE:535036}	535036 IMAGE Consortium	AA073514
4426 C	8 021MA1GP	IMAGE EST {IMAGE:535184}	535184 IMAGE Consortium	AA086732
4427 C	10 021MA1GP	M.musculus mRNA for s17 protein {IMAGE:535199}	535199 IMAGE Consortium	AA086796
4428 C	12 021MA1GP	ESTs {IMAGE:535718}	535718 IMAGE Consortium	AA106962
4429 E	2 021MA1GP	IMAGE EST {IMAGE:535886}	535886 IMAGE Consortium	AA105485
4430 E	4 021MA1GP	ESTs {IMAGE:525134}	525134 IMAGE Consortium	AA096594
4431 E	6 021MA1GP	ESTs {IMAGE:536412}	536412 IMAGE Consortium	AA073864
4432 E	8 021MA1GP	ESTs {IMAGE:550811}	550811 IMAGE Consortium	AA098149
4433 E	10 021MA1GP	ESTs {IMAGE:551184}	551184 IMAGE Consortium	AA080443
4434 E	12 021MA1GP	IMAGE EST {IMAGE:533489}	533489 IMAGE Consortium	AA105093
4435 G	2 021MA1GP	Cytochrome P450, 4a10 {IMAGE:550993}	550993 IMAGE Consortium	AA097980
4436 G	4 021MA1GP	IMAGE EST {IMAGE:533085}	533085 IMAGE Consortium	AA068665
4437 G	6 021MA1GP	IMAGE EST {IMAGE:385448}	385448 IMAGE Consortium	W62937
4438 G	8 021MA1GP	IMAGE EST {IMAGE:523842}	523842 IMAGE Consortium	AA067204
4439 G	10 021MA1GP	IMAGE EST {IMAGE:524510}	524510 IMAGE Consortium	AA068537

4440 G	12	021MA1GP	IMAGE EST {IMAGE:535949}	535949	IMAGE Consortium	AA073211
4441 A	2	021TA1GQ	IMAGE EST {IMAGE:536310}	536310	IMAGE Consortium	AA073606
4442 A	4	021TA1GQ	IMAGE EST {IMAGE:550973}	550973	IMAGE Consortium	AA098099
4443 A	6	021TA1GQ	IMAGE EST {IMAGE:522594}	522594	IMAGE Consortium	AA068007
4444 A	8	021TA1GQ	IMAGE EST {IMAGE:523322}	523322	IMAGE Consortium	AA080196
4445 A	10	021TA1GQ	UDP-glucose dehydrogenase {IMAGE:523796}	523796	IMAGE Consortium	AA067191
4446 A	12	021TA1GQ	ESTs {IMAGE:524641}	524641	IMAGE Consortium	AA068605
4447 C	2	021TA1GQ	ESTs {IMAGE:532253}	532253	IMAGE Consortium	AA066741
4448 C	4	021TA1GQ	IMAGE EST {IMAGE:532169}	532169	IMAGE Consortium	AA067045
4449 C	6	021TA1GQ	IMAGE EST {IMAGE:533100}	533100	IMAGE Consortium	AA068980
4450 C	8	021TA1GQ	IMAGE EST {IMAGE:524122}	524122	IMAGE Consortium	AA066174
4451 C	10	021TA1GQ	Heparan sulfate proteoglycan 1, cell surface-associated (fibroglycan) {IMAGE:535688}	535688	IMAGE Consortium	AA106952
4452 C	12	021TA1GQ	ESTs {IMAGE:535794}	535794	IMAGE Consortium	AA105996
4453 E	2	021TA1GQ	ESTs {IMAGE:536526}	536526	IMAGE Consortium	AA073872
4454 E	4	021TA1GQ	Lymphocyte protein tyrosine kinase {IMAGE:550845}	550845	IMAGE Consortium	AA098002
4455 E	6	021TA1GQ	ESTs, Highly similar to HYPOTHETICAL 29.1 KD PROTEIN IN URA7-POL12 INTERGENIC REGION [Saccharomyces cerevisiae] {IMAGE:522841}	522841	IMAGE Consortium	AA087650
4456 E	8	021TA1GQ	Mus musculus proline-rich protein (Bprp) mRNA, complete cds {IMAGE:374501}	374501	IMAGE Consortium	W64075
4457 E	10	021TA1GQ	Platelet-activating factor acetylhydrolase, isoform 1b, beta1 subunit {IMAGE:523706}	523706	IMAGE Consortium	AA067153
4458 E	12	021TA1GQ	ESTs {IMAGE:523742}	523742	IMAGE Consortium	AA067324
4459 G	2	021TA1GQ	ESTs, Highly similar to KINESIN-II 85 KD SUBUNIT [Strongylocentrotus purpuratus] {IMAGE:438886}	438886	IMAGE Consortium	AA008189
4460 G	4	021TA1GQ	IMAGE EST {IMAGE:657138}	657138	IMAGE Consortium	AA253627
4461 G	6	021TA1GQ	ESTs {IMAGE:662025}	662025	IMAGE Consortium	AA207462
4462 G	8	021TA1GQ	IMAGE EST {IMAGE:732593}	732593	IMAGE Consortium	AA286226
4463 G	10	021TA1GQ	IMAGE EST {IMAGE:633634}	633634	IMAGE Consortium	AA184529
4464 G	12	021TA1GQ	IMAGE EST {IMAGE:652282}	652282	IMAGE Consortium	AA215111
4465 A	2	0210A1GR	IMAGE EST {IMAGE:651986}	651986	IMAGE Consortium	AA209128
4466 A	4	0210A1GR	IMAGE EST {IMAGE:732669}	732669	IMAGE Consortium	AA286475
4467 A	6	0210A1GR	IMAGE EST {IMAGE:662909}	662909	IMAGE Consortium	AA208292
4468 A	8	0210A1GR	ESTs, Moderately similar to Tat-SF1 [H.sapiens] {IMAGE:651948}	651948	IMAGE Consortium	AA214786
4469 A	10	0210A1GR	IMAGE EST {IMAGE:917643}	917643	IMAGE Consortium	AA492969
4470 A	12	0210A1GR	IMAGE EST {IMAGE:917968}	917968	IMAGE Consortium	AA499758
4471 C	2	0210A1GR	ESTs {IMAGE:651648}	651648	IMAGE Consortium	AA212260
4472 C	4	0210A1GR	ESTs, Highly similar to ALPHA-ACTININ, SMOOTH MUSCLE ISOFORM [Gallus gallus] {IMAGE:733060}	733060	IMAGE Consortium	AA403949
4473 C	6	0210A1GR	IMAGE EST {IMAGE:656089}	656089	IMAGE Consortium	AA239554
4474 C	8	0210A1GR	IMAGE EST {IMAGE:651870}	651870	IMAGE Consortium	AA212640

4475 C	10 0210A1GR	ESTs {IMAGE:651988}		651988 IMAGE Consortium	AA209129
4476 C	12 0210A1GR	Preproenkephalin 2 {IMAGE:733276}		733276 IMAGE Consortium	AA403770
4477 E	2 0210A1GR	ESTs, Moderately similar to cell cycle progression 2 protein [H.sapiens] {IMAGE:656956}		656956 IMAGE Consortium	AA240480
4478 E	4 0210A1GR	ESTs, Moderately similar to E1B 19K/Bcl-2-interacting protein Nip3 [H.sapiens] {IMAGE:656945}		656945 IMAGE Consortium	AA240743
4479 E	6 0210A1GR	IMAGE EST {IMAGE:651304}		651304 IMAGE Consortium	AA214855
4480 E	8 0210A1GR	IG ALPHA CHAIN C REGION {IMAGE:551003}		551003 IMAGE Consortium	AA098196
4481 E	10 0210A1GR	IMAGE EST {IMAGE:652181}		652181 IMAGE Consortium	AA215158
4482 E	12 0210A1GR	Eukaryotic translation elongation factor 1 alpha 2 {IMAGE:330170}		330170 IMAGE Consortium	W11762
4483 G	2 0210A1GR	IMAGE EST {IMAGE:656667}		656667 IMAGE Consortium	AA240082
4484 G	4 0210A1GR	ESTs, Weakly similar to ORF YOR006c [S.cerevisiae] {IMAGE:661504}		661504 IMAGE Consortium	AA207598
4485 G	6 0210A1GR	ESTs, Moderately similar to H-2 class I histocompatibility antigen T15 [M.musculus] {IMAGE:651922}		651922 IMAGE Consortium	AA214771
4486 G	8 0210A1GR	IMAGE EST {IMAGE:917265}		917265 IMAGE Consortium	AA492934
4487 G	10 0210A1GR				
4488 G	12 0210A1GR				
4489 A	2 021Q9974	Control: Ratio 1:3 (Cy3:Cy5)	Ratio_A	Controls	
4490 A	4 021Q9974	Control: Ratio 3:1 (Cy3:Cy5)	Ratio_B	Controls	
4491 A	6 021Q9974	Control: Ratio 1:10 (Cy3:Cy5)	Ratio_C	Controls	
4492 A	8 021Q9974	Control: Ratio 10:1 (Cy3:Cy5)	Ratio_D	Controls	
4493 A	10 021Q9974	Control: Ratio 1:25 (Cy3:Cy5)	Ratio_E	Controls	
4494 A	12 021Q9974	Control: Ratio 25:1 (Cy3:Cy5)	Ratio_F	Controls	
4495 C	2 021Q9974	Control: Sensitivity 2pg	Sensitivity_A	Controls	
4496 C	4 021Q9974	Control: Sensitivity 20pg	Sensitivity_B	Controls	
4497 C	6 021Q9974	Control: Sensitivity 200pg	Sensitivity_C	Controls	
4498 C	8 021Q9974	Control: Sensitivity 2000pg	Sensitivity_D	Controls	
4499 C	10 021Q9974	Control: Buffer Only	Buffer_Only_Control	Controls	
4500 C	12 021Q9974	Control: Buffer Only	Buffer_Only_Control	Controls	
4501 E	2 021Q9974	Control: Internal	Internal_Control_A	Controls	
4502 E	4 021Q9974	Control: Buffer Only	Buffer_Only_Control	Controls	
4503 E	6 021Q9974	Control: Internal	Internal_Control_B	Controls	
4504 E	8 021Q9974	Control: Buffer Only	Buffer_Only_Control	Controls	
4505 E	10 021Q9974	Control: Internal	Internal_Control_C	Controls	
4506 E	12 021Q9974	Control: Internal	Internal_Control_D	Controls	
4507 G	2 021Q9974	Control: Complex Target (Mus musculus)	RNA_Quality_MM	Controls	
4508 G	4 021Q9974	Control: Complex Target (Mus musculus)	RNA_Quality_MM	Controls	
4509 G	6 021Q9974	Control: Complex Target (Mus musculus)	RNA_Quality_MM	Controls	
4510 G	8 021Q9974	Control: Housekeeping (Alpha Tubulin)	Housekeeping_B	Controls	
4511 G	10 021Q9974	Control: Housekeeping (Ribosomal S9)	Housekeeping_A	Controls	
4512 G	12 021Q9974	Control: Housekeeping (23kD HBP)	Housekeeping_C	Controls	

4513 B	1	2159971 Internal_Control_E	Internal_Control_E	Controls	
4514 B	3	2159971 Internal_Control_F	Internal_Control_F	Controls	
4515 B	5	2159971 Internal_Control_G	Internal_Control_G	Controls	
4516 B	7	2159971 Internal_Control_H	Internal_Control_H	Controls	
4517 B	9	2159971 Internal_Control_I	Internal_Control_I	Controls	
4518 B	11	2159971 Internal_Control_J	Internal_Control_J	Controls	
4519 D	1	2159971 Internal_Control_K	Internal_Control_K	Controls	
4520 D	3	2159971 Internal_Control_L	Internal_Control_L	Controls	
4521 D	5	2159971 Internal_Control_M	Internal_Control_M	Controls	
4522 D	7	2159971 Internal_Control_N	Internal_Control_N	Controls	
4523 D	9	2159971 Internal_Control_O	Internal_Control_O	Controls	
4524 D	11	2159971 Internal_Control_P	Internal_Control_P	Controls	
4525 F	1	2159971 Internal_Control_Q	Internal_Control_Q	Controls	
4526 F	3	2159971 Internal_Control_R	Internal_Control_R	Controls	
4527 F	5	2159971 Internal_Control_S	Internal_Control_S	Controls	
4528 F	7	2159971 Internal_Control_T	Internal_Control_T	Controls	
4529 F	9	2159971 Internal_Control_U	Internal_Control_U	Controls	
4530 F	11	2159971 Internal_Control_V	Internal_Control_V	Controls	
4531 H	1	2159971 Internal_Control_W	Internal_Control_W	Controls	
4532 H	3	2159971 Internal_Control_X	Internal_Control_X	Controls	
4533 H	5	2159971 Internal_Control_Y	Internal_Control_Y	Controls	
4534 H	7	2159971 Internal_Control_Z	Internal_Control_Z	Controls	
4535 H	9	2159971 Internal_Control_AA	Internal_Control_AA	Controls	
4536 H	11	2159971 Internal_Control_AB	Internal_Control_AB	Controls	
4537 B	1	021MA1E8 IMAGE EST {IMAGE:760594}	760594 IMAGE Consortium	AA386509	
4538 B	3	021MA1E8 ESTs {IMAGE:760769}	760769 IMAGE Consortium	AA386762	
4539 B	5	021MA1E8 Centromere autoantigen A {IMAGE:894092}	894092 IMAGE Consortium	AA516862	
4540 B	7	021MA1E8 IMAGE EST {IMAGE:894275}	894275 IMAGE Consortium	AA517335	
4541 B	9	021MA1E8 IMAGE EST {IMAGE:780202}	780202 IMAGE Consortium	AA386460	
4542 B	11	021MA1E8 ESTs, Weakly similar to cDNA EST EMBL:Z14593 comes from this gene [C.elegans] {IMAGE:934111}	934111 IMAGE Consortium	AA543829	
4543 D	1	021MA1E8 ESTs, Weakly similar to GSC-2 [H.sapiens] {IMAGE:804404}	804404 IMAGE Consortium	AA474743	
4544 D	3	021MA1E8 ESTs {IMAGE:934615}	934615 IMAGE Consortium	AA536786	
4545 D	5	021MA1E8 Myeloblastosis oncogene-like 2 {IMAGE:833557}	833557 IMAGE Consortium	AA432951	
4546 D	7	021MA1E8 ESTs, Weakly similar to contains strong similarity to a DNAJ-like domain [C.elegans] {IMAGE:834891}	834891 IMAGE Consortium	AA467252	
4547 D	9	021MA1E8 ESTs {IMAGE:894169}	894169 IMAGE Consortium	AA517043	
4548 D	11	021MA1E8 Cytochrome P450, 2e1, ethanol inducible {IMAGE:760871}	760871 IMAGE Consortium	AA386802	
4549 F	1	021MA1E8 IMAGE EST {IMAGE:894302}	894302 IMAGE Consortium	AA517346	
4550 F	3	021MA1E8 IMAGE EST {IMAGE:892366}	892366 IMAGE Consortium	AA501255	
4551 F	5	021MA1E8 ESTs {IMAGE:934500}	934500 IMAGE Consortium	AA536624	
4552 F	7	021MA1E8 ESTs {IMAGE:805893}	805893 IMAGE Consortium	AA399827	

4553 F	9	021MA1E8	ESTs {IMAGE:833919}	833919	IMAGE Consortium	AA413174
4554 F	11	021MA1E8	ESTs, Moderately similar to TFE3 transcription factor [H.sapiens] {IMAGE:834917}	834917	IMAGE Consortium	AA467269
4555 H	1	021MA1E8	ESTs, Moderately similar to RNA polymerase II transcription factor SIII p18 subunit [R.norvegicus] {IMAGE:933915}	933915	IMAGE Consortium	AA544428
4556 H	3	021MA1E8	ESTs {IMAGE:934337}	934337	IMAGE Consortium	AA544844
4557 H	5	021MA1E8	ESTs {IMAGE:833899}	833899	IMAGE Consortium	AA413020
4558 H	7	021MA1E8	IMAGE EST {IMAGE:835405}	835405	IMAGE Consortium	AA445803
4559 H	9	021MA1E8	RAD51 associated protein 1 {IMAGE:760795}	760795	IMAGE Consortium	AA386769
4560 H	11	021MA1E8	ESTs {IMAGE:761127}	761127	IMAGE Consortium	AA387679
4561 B	1	021TA1E9	ESTs, Weakly similar to cDNA EST EMBL:D72691 comes from this gene [C.elegans] {IMAGE:761319}	761319	IMAGE Consortium	AA388260
4562 B	3	021TA1E9	IMAGE EST {IMAGE:804726}	804726	IMAGE Consortium	AA474390
4563 B	5	021TA1E9	IMAGE EST {IMAGE:804793}	804793	IMAGE Consortium	AA472833
4564 B	7	021TA1E9	ESTs {IMAGE:805455}	805455	IMAGE Consortium	AA474587
4565 B	9	021TA1E9	ESTs {IMAGE:945014}	945014	IMAGE Consortium	AA537234
4566 B	11	021TA1E9	IMAGE EST {IMAGE:834617}	834617	IMAGE Consortium	AA467546
4567 D	1	021TA1E9	ESTs {IMAGE:805525}	805525	IMAGE Consortium	AA397161
4568 D	3	021TA1E9	Mus musculus RalBP1-associated EH domain protein Reps1 (reps1) mRNA, complete cds {IMAGE:834192}	834192	IMAGE Consortium	AA413178
4569 D	5	021TA1E9	ESTs, Weakly similar to contains strong similarity to a DNAJ-like domain [C.elegans] {IMAGE:760832}	760832	IMAGE Consortium	AA386630
4570 D	7	021TA1E9	ESTs, Weakly similar to similar to S. cerevisiae hypothetical protein YKL166 [C.elegans] {IMAGE:894102}	894102	IMAGE Consortium	AA516875
4571 D	9	021TA1E9	Breast cancer 2 {IMAGE:761119}	761119	IMAGE Consortium	AA387675
4572 D	11	021TA1E9	ESTs {IMAGE:933205}	933205	IMAGE Consortium	AA541928
4573 F	1	021TA1E9	ESTs {IMAGE:933229}	933229	IMAGE Consortium	AA541946
4574 F	3	021TA1E9	ESTs, Weakly similar to PROBABLE LEUCYL-TRNA SYNTHETASE [C.elegans] {IMAGE:804963}	804963	IMAGE Consortium	AA397172
4575 F	5	021TA1E9	IMAGE EST {IMAGE:805526}	805526	IMAGE Consortium	AA397153
4576 F	7	021TA1E9	ESTs {IMAGE:834291}	834291	IMAGE Consortium	AA413296
4577 F	9	021TA1E9	IMAGE EST {IMAGE:760824}	760824	IMAGE Consortium	AA386627
4578 F	11	021TA1E9	ESTs {IMAGE:760857}	760857	IMAGE Consortium	AA386550
4579 H	1	021TA1E9	IMAGE EST {IMAGE:774930}	774930	IMAGE Consortium	AA387184
4580 H	3	021TA1E9	IMAGE EST {IMAGE:775283}	775283	IMAGE Consortium	AA387141
4581 H	5	021TA1E9	IMAGE EST {IMAGE:789466}	789466	IMAGE Consortium	AA386980
4582 H	7	021TA1E9	IMAGE EST {IMAGE:774877}	774877	IMAGE Consortium	AA386758
4583 H	9	021TA1E9	ESTs, Weakly similar to RSP-1 PROTEIN [Mus musculus] {IMAGE:775307}	775307	IMAGE Consortium	AA387114
4584 H	11	021TA1E9	IMAGE EST {IMAGE:775093}	775093	IMAGE Consortium	AA386857
4585 B	1	021DA1EA	ESTs {IMAGE:793692}	793692	IMAGE Consortium	AA415691
4586 B	3	021DA1EA	ESTs {IMAGE:791051}	791051	IMAGE Consortium	AA415219
4587 B	5	021DA1EA	Mus musculus calmodulin-binding protein SHA1 (Sha1)	791460	IMAGE Consortium	AA416426



		mRNA, complete cds {IMAGE:791460}		
4588 B	7 021DA1EA	Selenophosphate synthetase 2 {IMAGE:791719}	791719 IMAGE Consortium	AA414662
4589 B	9 021DA1EA	ESTs {IMAGE:792921}	792921 IMAGE Consortium	AA415579
4590 B	11 021DA1EA	ESTs {IMAGE:779168}	779168 IMAGE Consortium	AA445722
4591 D	1 021DA1EA	ESTs {IMAGE:803594}	803594 IMAGE Consortium	AA422749
4592 D	3 021DA1EA	IMAGE EST {IMAGE:803629}	803629 IMAGE Consortium	AA422805
4593 D	5 021DA1EA	Cyclin E {IMAGE:833511}	833511 IMAGE Consortium	AA465987
4594 D	7 021DA1EA	ESTs, Highly similar to KIAA0095 gene is related to <i>S.cerevisiae</i> NIC96 gene. [H.sapiens] {IMAGE:792736}	792736 IMAGE Consortium	AA416435
4595 D	9 021DA1EA	IMAGE EST {IMAGE:779034}	779034 IMAGE Consortium	AA413764
4596 D	11 021DA1EA	ESTs, Weakly similar to predicted using Genefinder [C.elegans] {IMAGE:793789}	793789 IMAGE Consortium	AA415746
4597 F	1 021DA1EA	ESTs {IMAGE:833477}	833477 IMAGE Consortium	AA465972
4598 F	3 021DA1EA	IMAGE EST {IMAGE:836369}	836369 IMAGE Consortium	AA426860
4599 F	5 021DA1EA	ESTs {IMAGE:779373}	779373 IMAGE Consortium	AA414831
4600 F	7 021DA1EA	ESTs {IMAGE:792051}	792051 IMAGE Consortium	AA414507
4601 F	9 021DA1EA	IMAGE EST {IMAGE:791925}	791925 IMAGE Consortium	AA414432
4602 F	11 021DA1EA	IMAGE EST {IMAGE:779362}	779362 IMAGE Consortium	AA414108
4603 H	1 021DA1EA	ESTs {IMAGE:961036}	961036 IMAGE Consortium	AA547162
4604 H	3 021DA1EA	IMAGE EST {IMAGE:944469}	944469 IMAGE Consortium	AA545091
4605 H	5 021DA1EA	ESTs {IMAGE:961271}	961271 IMAGE Consortium	AA547550
4606 H	7 021DA1EA	ESTs, Highly similar to CYCLIN-DEPENDENT KINASES REGULATORY SUBUNIT 1 [Homo sapiens] {IMAGE:961282}	961282 IMAGE Consortium	AA547555
4607 H	9 021DA1EA	IMAGE EST {IMAGE:959853}	959853 IMAGE Consortium	AA547029
4608 H	11 021DA1EA	IMAGE EST {IMAGE:961094}	961094 IMAGE Consortium	AA547202
4609 B	1 021KA1EB	IMAGE EST {IMAGE:775342}	775342 IMAGE Consortium	AA386941
4610 B	3 021KA1EB	Tetratricopeptide repeat domain {IMAGE:775150}	775150 IMAGE Consortium	AA386895
4611 B	5 021KA1EB	IMAGE EST {IMAGE:775228}	775228 IMAGE Consortium	AA386847
4612 B	7 021KA1EB	ESTs {IMAGE:775188}	775188 IMAGE Consortium	AA387164
4613 B	9 021KA1EB	IMAGE EST {IMAGE:775296}	775296 IMAGE Consortium	AA387144
4614 B	11 021KA1EB	IMAGE EST {IMAGE:775504}	775504 IMAGE Consortium	AA387267
4615 D	1 021KA1EB	ESTs {IMAGE:778268}	778268 IMAGE Consortium	AA413761
4616 D	3 021KA1EB	Mus musculus p53 responsive (E124) mRNA, complete cds {IMAGE:778590}	778590 IMAGE Consortium	AA413622
4617 D	5 021KA1EB	IMAGE EST {IMAGE:778447}	778447 IMAGE Consortium	AA413717
4618 D	7 021KA1EB	ESTs {IMAGE:778602}	778602 IMAGE Consortium	AA413568
4619 D	9 021KA1EB	IMAGE EST {IMAGE:778923}	778923 IMAGE Consortium	AA414282
4620 D	11 021KA1EB	ESTs, Highly similar to N33 protein form 2 [H.sapiens] {IMAGE:779020}	779020 IMAGE Consortium	AA413956
4621 F	1 021KA1EB	IMAGE EST {IMAGE:803681}	803681 IMAGE Consortium	AA422887
4622 F	3 021KA1EB	ESTs {IMAGE:791433}	791433 IMAGE Consortium	AA416417
4623 F	5 021KA1EB	ESTs, Weakly similar to cyclin E [M.musculus] {IMAGE:791957}	791957 IMAGE Consortium	AA414293

4624 F	7 021KA1EB	IMAGE EST {IMAGE:791927}	791927 IMAGE Consortium	AA414433
4625 F	9 021KA1EB	IMAGE EST {IMAGE:778819}	778819 IMAGE Consortium	AA413678
4626 F	11 021KA1EB	ESTs, Weakly similar to LONG-CHAIN-FATTY-ACID--COA LIGASE [M.musculus] {IMAGE:780111}	780111 IMAGE Consortium	AA414751
4627 H	1 021KA1EB	ESTs {IMAGE:803897}	803897 IMAGE Consortium	AA435252
4628 H	3 021KA1EB	ESTs, Moderately similar to growth factor-responsive protein, vascular smooth muscle [R.norvegicus] {IMAGE:804313}	804313 IMAGE Consortium	AA474681
4629 H	5 021KA1EB	IMAGE EST {IMAGE:833403}	833403 IMAGE Consortium	AA465936
4630 H	7 021KA1EB	ESTs {IMAGE:835488}	835488 IMAGE Consortium	AA422455
4631 H	9 021KA1EB	IMAGE EST {IMAGE:779166}	779166 IMAGE Consortium	AA467575
4632 H	11 021KA1EB	IMAGE EST {IMAGE:779453}	779453 IMAGE Consortium	AA414212
4633 B	1 021RA1EC	ESTs {IMAGE:315890}	315890 IMAGE Consortium	W12793
4634 B	3 021RA1EC	IMAGE EST {IMAGE:315790}	315790 IMAGE Consortium	W12390
4635 B	5 021RA1EC	Laminin, gamma 2 {IMAGE:315993}	315993 IMAGE Consortium	W09048
4636 B	7 021RA1EC	IMAGE EST {IMAGE:330150}	330150 IMAGE Consortium	W11609
4637 B	9 021RA1EC	ESTs {IMAGE:330233}	330233 IMAGE Consortium	W14737
4638 B	11 021RA1EC	IMAGE EST {IMAGE:316933}	316933 IMAGE Consortium	W11395
4639 D	1 021RA1EC	IMAGE EST {IMAGE:315623}	315623 IMAGE Consortium	W11857
4640 D	3 021RA1EC	Histocompatibility 2, K region {IMAGE:330336}	330336 IMAGE Consortium	W14540
4641 D	5 021RA1EC	IMAGE EST {IMAGE:316482}	316482 IMAGE Consortium	W12977
4642 D	7 021RA1EC	ESTs {IMAGE:316804}	316804 IMAGE Consortium	W11317
4643 D	9 021RA1EC	ESTs {IMAGE:331126}	331126 IMAGE Consortium	W13745
4644 D	11 021RA1EC	IMAGE EST {IMAGE:317368}	317368 IMAGE Consortium	W34018
4645 F	1 021RA1EC	ESTs {IMAGE:330408}	330408 IMAGE Consortium	W14841
4646 F	3 021RA1EC	Caveolin, caveolae protein, 22 kDa {IMAGE:331186}	331186 IMAGE Consortium	W13196
4647 F	5 021RA1EC	IMAGE EST {IMAGE:317156}	317156 IMAGE Consortium	W35841
4648 F	7 021RA1EC	IMAGE EST {IMAGE:313840}	313840 IMAGE Consortium	W11016
4649 F	9 021RA1EC	ESTs, Moderately similar to MITOCHONDRIAL IMPORT INNER MEMBRANE TRANSLOCASE SUBUNIT TIM17 [Saccharomyces cerevisiae] {IMAGE:318134}	318134 IMAGE Consortium	W11644
4650 F	11 021RA1EC	IMAGE EST {IMAGE:318263}	318263 IMAGE Consortium	W13243
4651 H	1 021RA1EC	ESTs {IMAGE:317515}	317515 IMAGE Consortium	W34080
4652 H	3 021RA1EC	ESTs, Highly similar to SERINE CARBOXYPEPTIDASE PRECURSOR [Arabidopsis thaliana] {IMAGE:314456}	314456 IMAGE Consortium	W10703
4653 H	5 021RA1EC	ESTs {IMAGE:318817}	318817 IMAGE Consortium	W34216
4654 H	7 021RA1EC	IMAGE EST {IMAGE:329705}	329705 IMAGE Consortium	W13316
4655 H	9 021RA1EC	Mus musculus mRNA for protein-tyrosine-phosphatase IF1 {IMAGE:316284}	316284 IMAGE Consortium	W12121
4656 H	11 021RA1EC	IMAGE EST {IMAGE:330298}	330298 IMAGE Consortium	W13940
4657 B	1 021YA1ED	ALPHA-MANNOSIDASE II {IMAGE:315962}	315962 IMAGE Consortium	W09023
4658 B	3 021YA1ED	Dipeptidase 1 (renal) {IMAGE:316237}	316237 IMAGE Consortium	W08886
4659 B	5 021YA1ED	ESTs {IMAGE:330502}	330502 IMAGE Consortium	W15031
4660 B	7 021YA1ED	ESTs {IMAGE:317148}	317148 IMAGE Consortium	W13658

4661 B	9	021YA1ED	CTP phosphocholine cytidyl transferase {IMAGE:317421}	317421	IMAGE Consortium	W34054
4662 B	11	021YA1ED	ESTs {IMAGE:313652}	313652	IMAGE Consortium	W10662
4663 D	1	021YA1ED	ESTs {IMAGE:330672}	330672	IMAGE Consortium	W14237
4664 D	3	021YA1ED	IMAGE EST {IMAGE:313254}	313254	IMAGE Consortium	W11906
4665 D	5	021YA1ED	ESTs {IMAGE:317676}	317676	IMAGE Consortium	W11217
4666 D	7	021YA1ED	ESTs {IMAGE:314156}	314156	IMAGE Consortium	W12425
4667 D	9	021YA1ED	ESTs {IMAGE:315125}	315125	IMAGE Consortium	W12131
4668 D	11	021YA1ED	Mus musculus ES2 protein (Es2) mRNA, complete cds {IMAGE:318785}	318785	IMAGE Consortium	W34498
4669 F	1	021YA1ED	IMAGE EST {IMAGE:318128}	318128	IMAGE Consortium	W11641
4670 F	3	021YA1ED	ESTs {IMAGE:315663}	315663	IMAGE Consortium	W09954
4671 F	5	021YA1ED	ESTs, Weakly similar to PERIPLASMIC DIVALENT CATION TOLERANCE PROTEIN CUTA [Escherichia coli] {IMAGE:316110}	316110	IMAGE Consortium	W12255
4672 F	7	021YA1ED	IMAGE EST {IMAGE:330147}	330147	IMAGE Consortium	W11614
4673 F	9	021YA1ED	IMAGE EST {IMAGE:330534}	330534	IMAGE Consortium	W15055
4674 F	11	021YA1ED	Twist gene homolog, (Drosophila) {IMAGE:331264}	331264	IMAGE Consortium	W14113
4675 H	1	021YA1ED	ESTs, Weakly similar to orf, hypothetical protein [E.coli] {IMAGE:314236}	314236	IMAGE Consortium	W09936
4676 H	3	021YA1ED	IMAGE EST {IMAGE:314524}	314524	IMAGE Consortium	W10780
4677 H	5	021YA1ED	Mus musculus ADP-ribosylarginine hydrolase mRNA, complete cds {IMAGE:314462}	314462	IMAGE Consortium	W10726
4678 H	7	021YA1ED	Cathepsin L {IMAGE:315082}	315082	IMAGE Consortium	AA174215
4679 H	9	021YA1ED	ESTs {IMAGE:315787}	315787	IMAGE Consortium	W12392
4680 H	11	021YA1ED	ESTs, Moderately similar to INTEGRAL MEMBRANE PROTEIN SED5 [Saccharomyces cerevisiae] {IMAGE:316087}	316087	IMAGE Consortium	W09139
4681 B	1	0215A1EE	IMAGE EST {IMAGE:332396}	332396	IMAGE Consortium	W08720
4682 B	3	0215A1EE	ESTs {IMAGE:332791}	332791	IMAGE Consortium	W07997
4683 B	5	0215A1EE	ESTs {IMAGE:336334}	336334	IMAGE Consortium	W18457
4684 B	7	0215A1EE	IMAGE EST {IMAGE:333274}	333274	IMAGE Consortium	W33809
4685 B	9	0215A1EE	IMAGE EST {IMAGE:334450}	334450	IMAGE Consortium	AA052395
4686 B	11	0215A1EE	Pale ear {IMAGE:347985}	347985	IMAGE Consortium	W29855
4687 D	1	0215A1EE	ESTs {IMAGE:337964}	337964	IMAGE Consortium	W29642
4688 D	3	0215A1EE	ESTs {IMAGE:347890}	347890	IMAGE Consortium	W29589
4689 D	5	0215A1EE	Keratin complex 2, gene 6A {IMAGE:335736}	335736	IMAGE Consortium	W20986
4690 D	7	0215A1EE	Sphingosine phosphate lyase 1 {IMAGE:332091}	332091	IMAGE Consortium	W08172
4691 D	9	0215A1EE	IMAGE EST {IMAGE:332181}	332181	IMAGE Consortium	W08394
4692 D	11	0215A1EE	ESTs, Weakly similar to B0414.8 [C.elegans] {IMAGE:336399}	336399	IMAGE Consortium	W36692
4693 F	1	0215A1EE	IMAGE EST {IMAGE:335385}	335385	IMAGE Consortium	W17702
4694 F	3	0215A1EE	ESTs {IMAGE:335979}	335979	IMAGE Consortium	W18013
4695 F	5	0215A1EE	Cytochrome P450, 2e1, ethanol inducible {IMAGE:335899}	335899	IMAGE Consortium	W36926

4696 F	7 0215A1EE	3'-phosphoadenosine 5'-phosphosulfate synthase 2 {IMAGE:336536}	336536 IMAGE Consortium	W18519
4697 F	9 0215A1EE	ESTs, Weakly similar to MHC H-2K/t-w5-linked ORF precursor [M.musculus] {IMAGE:336497}	336497 IMAGE Consortium	W18585
4698 F	11 0215A1EE	TUBULIN BETA-5 CHAIN {IMAGE:333325}	333325 IMAGE Consortium	W16254
4699 H	1 0215A1EE	ESTs {IMAGE:332381}	332381 IMAGE Consortium	W07947
4700 H	3 0215A1EE	IMAGE EST {IMAGE:332625}	332625 IMAGE Consortium	W08703
4701 H	5 0215A1EE	ESTs, Weakly similar to plakophilin 1 [M.musculus] {IMAGE:332567}	332567 IMAGE Consortium	W08501
4702 H	7 0215A1EE	ESTs, Moderately similar to HYPOTHETICAL 16.7 KD PROTEIN MRP17-MET14 INTERGENIC REGION [Saccharomyces cerevisiae] {IMAGE:337278}	337278 IMAGE Consortium	W20752
4703 H	9 0215A1EE	ESTs, Moderately similar to ELONGATION FACTOR 1-ALPHA [Trichoderma reesei] {IMAGE:337147}	337147 IMAGE Consortium	W20903
4704 H	11 0215A1EE	Mus musculus asparagine synthetase mRNA, complete cds {IMAGE:337748}	337748 IMAGE Consortium	W29492
4705 B	1 021CA1EF	ESTs, Moderately similar to sarcolipin [H.sapiens] {IMAGE:318692}	318692 IMAGE Consortium	W11587
4706 B	3 021CA1EF	IMAGE EST {IMAGE:330076}	330076 IMAGE Consortium	W11846
4707 B	5 021CA1EF	ESTs {IMAGE:330495}	330495 IMAGE Consortium	W14898
4708 B	7 021CA1EF	IMAGE EST {IMAGE:330871}	330871 IMAGE Consortium	W14938
4709 B	9 021CA1EF	ESTs {IMAGE:331327}	331327 IMAGE Consortium	W14581
4710 B	11 021CA1EF	IMAGE EST {IMAGE:331531}	331531 IMAGE Consortium	W14371
4711 D	1 021CA1EF	IMAGE EST {IMAGE:334554}	334554 IMAGE Consortium	W18526
4712 D	3 021CA1EF	ESTs {IMAGE:334783}	334783 IMAGE Consortium	W17605
4713 D	5 021CA1EF	Arachidonate 12-lipoxygenase, pseudogene 2 {IMAGE:335369}	335369 IMAGE Consortium	W36511
4714 D	7 021CA1EF	ESTs {IMAGE:331822}	331822 IMAGE Consortium	W08071
4715 D	9 021CA1EF	IMAGE EST {IMAGE:336482}	336482 IMAGE Consortium	W18386
4716 D	11 021CA1EF	Fibroblast inducible secreted protein {IMAGE:336325}	336325 IMAGE Consortium	W36541
4717 F	1 021CA1EF	IMAGE EST {IMAGE:335210}	335210 IMAGE Consortium	W17578
4718 F	3 021CA1EF	Mus musculus ubiquitin conjugating enzyme UBC9 mRNA, complete cds {IMAGE:336206}	336206 IMAGE Consortium	W18065
4719 F	5 021CA1EF	Apolipoprotein C2 linked {IMAGE:336398}	336398 IMAGE Consortium	W36685
4720 F	7 021CA1EF	ESTs, Highly similar to ELECTRON TRANSFER FLAVOPROTEIN BETA-SUBUNIT [Paracoccus denitrificans] {IMAGE:333641}	333641 IMAGE Consortium	W18161
4721 F	9 021CA1EF	ESTs {IMAGE:334016}	334016 IMAGE Consortium	W16166
4722 F	11 021CA1EF	IMAGE EST {IMAGE:334052}	334052 IMAGE Consortium	W17543
4723 H	1 021CA1EF	IMAGE EST {IMAGE:332477}	332477 IMAGE Consortium	W08616
4724 H	3 021CA1EF	ESTs, Weakly similar to R07E5.13 protein [C.elegans] {IMAGE:332442}	332442 IMAGE Consortium	W08432
4725 H	5 021CA1EF	Mus musculus amino levulinate synthase (ALAS-H) mRNA, 3' end {IMAGE:332943}	332943 IMAGE Consortium	W15813

4726 H	7 021CA1EF	ESTs, Weakly similar to HYPOTHETICAL 103.9 KD PROTEIN ZK370.3 IN CHROMOSOME III [Caenorhabditis elegans] {IMAGE:333511}	333511 IMAGE Consortium	W33982
4727 H	9 021CA1EF	IMAGE EST {IMAGE:333239}	333239 IMAGE Consortium	W18382
4728 H	11 021CA1EF	ESTs, Moderately similar to KIAA0330 [H.sapiens] {IMAGE:333618}	333618 IMAGE Consortium	W18153
4729 B	1 021JA1EG	IMAGE EST {IMAGE:336045}	336045 IMAGE Consortium	W36407
4730 B	3 021JA1EG	IMAGE EST {IMAGE:336282}	336282 IMAGE Consortium	W18224
4731 B	5 021JA1EG	IMAGE EST {IMAGE:333094}	333094 IMAGE Consortium	W15714
4732 B	7 021JA1EG	IMAGE EST {IMAGE:337401}	337401 IMAGE Consortium	W18639
4733 B	9 021JA1EG	Somatostatin {IMAGE:337219}	337219 IMAGE Consortium	W20888
4734 B	11 021JA1EG	IMAGE EST {IMAGE:337772}	337772 IMAGE Consortium	W29499
4735 D	1 021JA1EG	Mus musculus A10 mRNA, partial cds {IMAGE:333376}	333376 IMAGE Consortium	W15888
4736 D	3 021JA1EG	FK506-BINDING PROTEIN PRECURSOR {IMAGE:333683}	333683 IMAGE Consortium	W15971
4737 D	5 021JA1EG	Mus musculus mel (MEL91) mRNA, complete cds {IMAGE:337618}	337618 IMAGE Consortium	W18484
4738 D	7 021JA1EG	ESTs, Moderately similar to nebulin, skeletal muscle [H.sapiens] {IMAGE:334648}	334648 IMAGE Consortium	W18356
4739 D	9 021JA1EG	Mus musculus mRNA for pyruvate dehydrogenase kinase-like protein {IMAGE:334876}	334876 IMAGE Consortium	W17814
4740 D	11 021JA1EG	ESTs {IMAGE:348245}	348245 IMAGE Consortium	W18841
4741 F	1 021JA1EG	IMAGE EST {IMAGE:333636}	333636 IMAGE Consortium	W15957
4742 F	3 021JA1EG	ESTs, Highly similar to HYPOTHETICAL 25.7 KD PROTEIN IN MSH1-EPT1 INTERGENIC REGION [Saccharomyces cerevisiae] {IMAGE:337654}	337654 IMAGE Consortium	W54688
4743 F	5 021JA1EG	ESTs {IMAGE:334410}	334410 IMAGE Consortium	W18278
4744 F	7 021JA1EG	ESTs {IMAGE:334918}	334918 IMAGE Consortium	W36492
4745 F	9 021JA1EG	ESTs {IMAGE:351968}	351968 IMAGE Consortium	W40838
4746 F	11 021JA1EG	Mus musculus timeless homolog mRNA, complete cds {IMAGE:352877}	352877 IMAGE Consortium	W34732
4747 H	1 021JA1EG	IMAGE EST {IMAGE:350904}	350904 IMAGE Consortium	W40998
4748 H	3 021JA1EG	IMAGE EST {IMAGE:351720}	351720 IMAGE Consortium	W41405
4749 H	5 021JA1EG	ESTs {IMAGE:352643}	352643 IMAGE Consortium	AA052540
4750 H	7 021JA1EG	IMAGE EST {IMAGE:353202}	353202 IMAGE Consortium	W41810
4751 H	9 021JA1EG	ESTs, Highly similar to SMALL NUCLEAR RIBONUCLEOPROTEIN SM D2 [Homo sapiens] {IMAGE:351154}	351154 IMAGE Consortium	W41213
4752 H	11 021JA1EG	ESTs, Highly similar to MITOCHONDRIAL RNA SPLICING PROTEIN MSR4 [Saccharomyces cerevisiae] {IMAGE:354607}	354607 IMAGE Consortium	W46092
4753 B	1 021QA1EH	ESTs {IMAGE:354655}	354655 IMAGE Consortium	W46125
4754 B	3 021QA1EH	ESTs, Moderately similar to weak similarity to Arabidopsis thaliana ubiquitin-like protein 8 [C.elegans] {IMAGE:351643}	351643 IMAGE Consortium	W41624
4755 B	5 021QA1EH	ESTs, Moderately similar to GLUTATHIONE PEROXIDASE	353017 IMAGE Consortium	W35981

		[Schistosoma mansoni] {IMAGE:353017}		
4756 B	7 021QA1EH	EST {IMAGE:352795}	352795 IMAGE Consortium	W36053
4757 B	9 021QA1EH	Early B-cell factor {IMAGE:353456}	353456 IMAGE Consortium	W40683
4758 B	11 021QA1EH	ESTs {IMAGE:351123}	351123 IMAGE Consortium	W40740
4759 D	1 021QA1EH	ESTs {IMAGE:352798}	352798 IMAGE Consortium	W36059
4760 D	3 021QA1EH	ESTs, Weakly similar to EF-7 [M.musculus] {IMAGE:353221}	353221 IMAGE Consortium	W41647
4761 D	5 021QA1EH	IMAGE EST {IMAGE:353445}	353445 IMAGE Consortium	W41928
4762 D	7 021QA1EH	ESTs, Weakly similar to coded for by C. elegans cDNA yk157f8.5 [C.elegans] {IMAGE:354155}	354155 IMAGE Consortium	W43938
4763 D	9 021QA1EH	ESTs, Moderately similar to FAST kinase [H.sapiens] {IMAGE:354394}	354394 IMAGE Consortium	W42367
4764 D	11 021QA1EH	ESTs {IMAGE:352231}	352231 IMAGE Consortium	W33596
4765 F	1 021QA1EH	Mus musculus Numblike (m-nbl) mRNA, complete cds {IMAGE:352882}	352882 IMAGE Consortium	W34802
4766 F	3 021QA1EH	IMAGE EST {IMAGE:353294}	353294 IMAGE Consortium	W41937
4767 F	5 021QA1EH	Glycerol phosphate dehydrogenase 1, mitochondrial {IMAGE:351221}	351221 IMAGE Consortium	W41175
4768 F	7 021QA1EH	ESTs {IMAGE:351116}	351116 IMAGE Consortium	W41107
4769 F	9 021QA1EH	IMAGE EST {IMAGE:352541}	352541 IMAGE Consortium	W33902
4770 F	11 021QA1EH	PROTEIN-GLUTAMINE GAMMA-GLUTAMYLTRANSFERASE {IMAGE:352804}	352804 IMAGE Consortium	W34612
4771 H	1 021QA1EH	ESTs {IMAGE:353521}	353521 IMAGE Consortium	W40904
4772 H	3 021QA1EH	ESTs {IMAGE:350252}	350252 IMAGE Consortium	W34669
4773 H	5 021QA1EH	IMAGE EST {IMAGE:350756}	350756 IMAGE Consortium	W44064
4774 H	7 021QA1EH	Pentaxin related gene {IMAGE:354318}	354318 IMAGE Consortium	W42321
4775 H	9 021QA1EH	Histone deacetylase 2 {IMAGE:355035}	355035 IMAGE Consortium	W46034
4776 H	11 021QA1EH	IMAGE EST {IMAGE:352663}	352663 IMAGE Consortium	AA052541
4777 B	1 021XA1EI	IMAGE EST {IMAGE:368432}	368432 IMAGE Consortium	W53698
4778 B	3 021XA1EI	IMAGE EST {IMAGE:354969}	354969 IMAGE Consortium	W46002
4779 B	5 021XA1EI	ESTs, Highly similar to INTEGRAL MEMBRANE GLYCOPROTEIN GP210 PRECURSOR [Rattus norvegicus] {IMAGE:355445}	355445 IMAGE Consortium	W48307
4780 B	7 021XA1EI	IMAGE EST {IMAGE:355808}	355808 IMAGE Consortium	W47799
4781 B	9 021XA1EI	ESTs {IMAGE:355932}	355932 IMAGE Consortium	W48111
4782 B	11 021XA1EI	COPROPORPHYRINOGEN III OXIDASE PRECURSOR {IMAGE:367358}	367358 IMAGE Consortium	W53951
4783 D	1 021XA1EI	Mus musculus mRNA for type II phosphatidylinositolphosphate kinase-alpha, complete cds {IMAGE:372957}	372957 IMAGE Consortium	W58969
4784 D	3 021XA1EI	ESTs {IMAGE:367532}	367532 IMAGE Consortium	W54420
4785 D	5 021XA1EI	ESTs, Highly similar to NADH-UBIQUINONE OXIDOREDUCTASE SUBUNIT B14.5B [Bos taurus] {IMAGE:367925}	367925 IMAGE Consortium	W54068
4786 D	7 021XA1EI	IMAGE EST {IMAGE:354815}	354815 IMAGE Consortium	W45812

4787 D	9 021XA1EI	IMAGE EST {IMAGE:355165}	355165 IMAGE Consortium	W48287
4788 D	11 021XA1EI	ESTs, Weakly similar to ladinin [H.sapiens] {IMAGE:372468}	372468 IMAGE Consortium	W61552
4789 F	1 021XA1EI	IMAGE EST {IMAGE:385725}	385725 IMAGE Consortium	W64239
4790 F	3 021XA1EI	IMAGE EST {IMAGE:386122}	386122 IMAGE Consortium	W62349
4791 F	5 021XA1EI	IMAGE EST {IMAGE:386109}	386109 IMAGE Consortium	W64945
4792 F	7 021XA1EI	IMAGE EST {IMAGE:390337}	390337 IMAGE Consortium	W70804
4793 F	9 021XA1EI	ESTs {IMAGE:390396}	390396 IMAGE Consortium	W70826
4794 F	11 021XA1EI	ESTs, Highly similar to nucleoporin Nup84 [R.norvegicus] {IMAGE:374564}	374564 IMAGE Consortium	W63828
4795 H	1 021XA1EI	ESTs {IMAGE:387284}	387284 IMAGE Consortium	W65621
4796 H	3 021XA1EI	ESTs {IMAGE:387936}	387936 IMAGE Consortium	W70626
4797 H	5 021XA1EI	IMAGE EST {IMAGE:388376}	388376 IMAGE Consortium	W70912
4798 H	7 021XA1EI	ESTs, Weakly similar to Oxoglutarate/malate carrier protein [C.elegans] {IMAGE:388288}	388288 IMAGE Consortium	W66635
4799 H	9 021XA1EI	ESTs {IMAGE:388580}	388580 IMAGE Consortium	W67079
4800 H	11 021XA1EI	IMAGE EST {IMAGE:385812}	385812 IMAGE Consortium	W62202
4801 B	1 0214A1EJ	ESTs {IMAGE:374179}	374179 IMAGE Consortium	W63930
4802 B	3 0214A1EJ	Histocompatibility 2, D region locus 1 {IMAGE:355765}	355765 IMAGE Consortium	W47808
4803 B	5 0214A1EJ	Mouse chromosome 6 BAC-284H12 (Research Genetics mouse BAC library) complete sequence {IMAGE:355935}	355935 IMAGE Consortium	W48119
4804 B	7 0214A1EJ	Neural cell adhesion molecule {IMAGE:367050}	367050 IMAGE Consortium	W54389
4805 B	9 0214A1EJ	Nuclear factor I/X {IMAGE:367660}	367660 IMAGE Consortium	W55827
4806 B	11 0214A1EJ	ESTs, Highly similar to PROTEIN TRANSPORT PROTEIN SEC61 ALPHA SUBUNIT [Canis familiaris] {IMAGE:373834}	373834 IMAGE Consortium	W62742
4807 D	1 0214A1EJ	ESTs, Moderately similar to OXA1L [H.sapiens] {IMAGE:373988}	373988 IMAGE Consortium	W62516
4808 D	3 0214A1EJ	IMAGE EST {IMAGE:355778}	355778 IMAGE Consortium	W47810
4809 D	5 0214A1EJ	IMAGE EST {IMAGE:372023}	372023 IMAGE Consortium	W58881
4810 D	7 0214A1EJ	IMAGE EST {IMAGE:372639}	372639 IMAGE Consortium	W62653
4811 D	9 0214A1EJ	ESTs {IMAGE:367565}	367565 IMAGE Consortium	W54715
4812 D	11 0214A1EJ	ESTs {IMAGE:367529}	367529 IMAGE Consortium	W54421
4813 F	1 0214A1EJ	ESTs, Highly similar to MITOCHONDRIAL 2-OXOGLUTARATE/MALATE CARRIER PROTEIN [Bos taurus] {IMAGE:367801}	367801 IMAGE Consortium	W54000
4814 F	3 0214A1EJ	IMAGE EST {IMAGE:374240}	374240 IMAGE Consortium	W59175
4815 F	5 0214A1EJ	Hypoxanthine guanine phosphoribosyl transferase {IMAGE:355084}	355084 IMAGE Consortium	W48168
4816 F	7 0214A1EJ	Mus musculus activin type IB receptor (ALK-1) mRNA, complete cds {IMAGE:355337}	355337 IMAGE Consortium	W48139
4817 F	9 0214A1EJ	Inhibitor of DNA binding 4 {IMAGE:355929}	355929 IMAGE Consortium	W48116
4818 F	11 0214A1EJ	IMAGE EST {IMAGE:372592}	372592 IMAGE Consortium	W58982
4819 H	1 0214A1EJ	ESTs {IMAGE:355157}	355157 IMAGE Consortium	W48283
4820 H	3 0214A1EJ	M.musculus mRNA for N-glycan alpha 2,8-sialyltransferase	355197 IMAGE Consortium	W47914

		{IMAGE:355197}		
4821 H	5 0214A1EJ	IMAGE EST {IMAGE:355507}	355507 IMAGE Consortium	W48190
4822 H	7 0214A1EJ	IMAGE EST {IMAGE:372366}	372366 IMAGE Consortium	W53543
4823 H	9 0214A1EJ	Mouse mRNA for nociceptin/orphanin FQ, complete cds {IMAGE:367399}	367399 IMAGE Consortium	W50641
4824 H	11 0214A1EJ	Glutathione-S-transferase, alpha 1 (Ya) {IMAGE:367627}	367627 IMAGE Consortium	W54349
4825 B	1 021BA1EK	IMAGE EST {IMAGE:385119}	385119 IMAGE Consortium	W62894
4826 B	3 021BA1EK	ESTs, Highly similar to OOCYTE ZINC FINGER PROTEIN XLCOF28 [ <i>Xenopus laevis</i> ] {IMAGE:385763}	385763 IMAGE Consortium	W62249
4827 B	5 021BA1EK	Keratin complex 1, acidic, gene 10 {IMAGE:388916}	388916 IMAGE Consortium	W67022
4828 B	7 021BA1EK	Imprinted polyspecific membrane transporter-like gene 1 {IMAGE:389325}	389325 IMAGE Consortium	W71351
4829 B	9 021BA1EK	ESTs, Weakly similar to MITOCHONDRIAL 60S RIBOSOMAL PROTEIN L2 PRECURSOR [ <i>S.cerevisiae</i> ] {IMAGE:389993}	389993 IMAGE Consortium	W71625
4830 B	11 021BA1EK	ESTs, Highly similar to NADH-UBIQUINONE OXIDOREDUCTASE SUBUNIT B14.5A [ <i>Bos taurus</i> ] {IMAGE:386766}	386766 IMAGE Consortium	W64426
4831 D	1 021BA1EK	ESTs {IMAGE:389205}	389205 IMAGE Consortium	W74845
4832 D	3 021BA1EK	Mouse mRNA for meltrin alpha, complete cds {IMAGE:386236}	386236 IMAGE Consortium	W65081
4833 D	5 021BA1EK	IMAGE EST {IMAGE:390150}	390150 IMAGE Consortium	W70677
4834 D	7 021BA1EK	ESTs {IMAGE:390329}	390329 IMAGE Consortium	W74931
4835 D	9 021BA1EK	IMAGE EST {IMAGE:386963}	386963 IMAGE Consortium	W65725
4836 D	11 021BA1EK	ESTs {IMAGE:388074}	388074 IMAGE Consortium	W66934
4837 F	1 021BA1EK	NEURONAL PROTEIN 3.1 {IMAGE:374970}	374970 IMAGE Consortium	W62819
4838 F	3 021BA1EK	IMAGE EST {IMAGE:387773}	387773 IMAGE Consortium	W71660
4839 F	5 021BA1EK	ESTs, Highly similar to NEUROPEPTIDE Y PRECURSOR [ <i>Gallus gallus</i> ] {IMAGE:385331}	385331 IMAGE Consortium	W70782
4840 F	7 021BA1EK	Fyn proto-oncogene {IMAGE:385072}	385072 IMAGE Consortium	W62969
4841 F	9 021BA1EK	<i>M. musculus</i> mRNA for MAP kinase-activated protein kinase 2 {IMAGE:388617}	388617 IMAGE Consortium	W66726
4842 F	11 021BA1EK	IMAGE EST {IMAGE:385626}	385626 IMAGE Consortium	W65201
4843 H	1 021BA1EK	IMAGE EST {IMAGE:385621}	385621 IMAGE Consortium	W61502
4844 H	3 021BA1EK	IMAGE EST {IMAGE:385901}	385901 IMAGE Consortium	W62273
4845 H	5 021BA1EK	IMAGE EST {IMAGE:385914}	385914 IMAGE Consortium	W64255
4846 H	7 021BA1EK	IMAGE EST {IMAGE:386384}	386384 IMAGE Consortium	W64296
4847 H	9 021BA1EK	ESTs, Highly similar to GTP-binding protein-associated protein [ <i>M.musculus</i> ] {IMAGE:374296}	374296 IMAGE Consortium	W65211
4848 H	11 021BA1EK	Cytochrome P450, 2e1, ethanol inducible {IMAGE:374886}	374886 IMAGE Consortium	W61665
4849 B	1 021IA1EL	ESTs {IMAGE:385949}	385949 IMAGE Consortium	W62311
4850 B	3 021IA1EL	IMAGE EST {IMAGE:389880}	389880 IMAGE Consortium	W70688
4851 B	5 021IA1EL	IMAGE EST {IMAGE:386417}	386417 IMAGE Consortium	W64285



4852 B	7 021IA1EL	ESTs {IMAGE:374369}	374369 IMAGE Consortium	W65236
4853 B	9 021IA1EL	ESTs, Highly similar to RTS1 PROTEIN [Saccharomyces cerevisiae] {IMAGE:387506}	387506 IMAGE Consortium	W65797
4854 B	11 021IA1EL	IMAGE EST {IMAGE:388091}	388091 IMAGE Consortium	W70420
4855 D	1 021IA1EL	IMAGE EST {IMAGE:374745}	374745 IMAGE Consortium	W59461
4856 D	3 021IA1EL	IMAGE EST {IMAGE:387619}	387619 IMAGE Consortium	W66857
4857 D	5 021IA1EL	ESTs {IMAGE:388190}	388190 IMAGE Consortium	W70536
4858 D	7 021IA1EL	ESTs, Highly similar to NEURONAL PROTEIN NP25 [Rattus norvegicus] {IMAGE:388461}	388461 IMAGE Consortium	W67105
4859 D	9 021IA1EL	IMAGE EST {IMAGE:388936}	388936 IMAGE Consortium	W67028
4860 D	11 021IA1EL	ESTs {IMAGE:390312}	390312 IMAGE Consortium	W70776
4861 F	1 021IA1EL	ESTs {IMAGE:386349}	386349 IMAGE Consortium	W64937
4862 F	3 021IA1EL	IMAGE EST {IMAGE:402221}	402221 IMAGE Consortium	W76872
4863 F	5 021IA1EL	IMAGE EST {IMAGE:402963}	402963 IMAGE Consortium	W80160
4864 F	7 021IA1EL	ESTs, Moderately similar to P21 activated kinase-3 [M.musculus] {IMAGE:391871}	391871 IMAGE Consortium	W77082
4865 F	9 021IA1EL	IMAGE EST {IMAGE:400219}	400219 IMAGE Consortium	W77046
4866 F	11 021IA1EL	ESTs {IMAGE:403770}	403770 IMAGE Consortium	W82732
4867 H	1 021IA1EL	IMAGE EST {IMAGE:400200}	400200 IMAGE Consortium	W77521
4868 H	3 021IA1EL	Fc receptor, IgG, low affinity III {IMAGE:403499}	403499 IMAGE Consortium	W82159
4869 H	5 021IA1EL	Cytochrome P450, 2e1, ethanol inducible {IMAGE:403678}	403678 IMAGE Consortium	W82358
4870 H	7 021IA1EL	M.musculus rab3A gene {IMAGE:403856}	403856 IMAGE Consortium	W82220
4871 H	9 021IA1EL	IMAGE EST {IMAGE:403961}	403961 IMAGE Consortium	W82923
4872 H	11 021IA1EL	PS2 PROTEIN PRECURSOR {IMAGE:404550}	404550 IMAGE Consortium	W83072
4873 B	1 021PA1EM	IMAGE EST {IMAGE:402985}	402985 IMAGE Consortium	W81895
4874 B	3 021PA1EM	ESTs {IMAGE:402904}	402904 IMAGE Consortium	W80179
4875 B	5 021PA1EM	IMAGE EST {IMAGE:403189}	403189 IMAGE Consortium	W81737
4876 B	7 021PA1EM	ESTs {IMAGE:400286}	400286 IMAGE Consortium	W77536
4877 B	9 021PA1EM	ESTs, Weakly similar to zinc finger protein [M.musculus] {IMAGE:403670}	403670 IMAGE Consortium	W82276
4878 B	11 021PA1EM	Lysosomal membrane glycoprotein 1 {IMAGE:403851}	403851 IMAGE Consortium	W82141
4879 D	1 021PA1EM	ESTs {IMAGE:401441}	401441 IMAGE Consortium	AA000556
4880 D	3 021PA1EM	ESTs {IMAGE:401741}	401741 IMAGE Consortium	W78474
4881 D	5 021PA1EM	IMAGE EST {IMAGE:405888}	405888 IMAGE Consortium	W83214
4882 D	7 021PA1EM	ESTs, Highly similar to PROBABLE ATP-DEPENDENT TRANSPORTER YER036C [Saccharomyces cerevisiae] {IMAGE:406088}	406088 IMAGE Consortium	W83271
4883 D	9 021PA1EM	ESTs, Highly similar to HYPOTHETICAL 27.6 KD PROTEIN IN PDX1-SNG1 INTERGENIC REGION [Saccharomyces cerevisiae] {IMAGE:402449}	402449 IMAGE Consortium	W80023
4884 D	11 021PA1EM	ESTs, Highly similar to PYRROLINE-5-CARBOXYLATE REDUCTASE [Homo sapiens] {IMAGE:402730}	402730 IMAGE Consortium	W80133
4885 F	1 021PA1EM	Cytochrome P450, 2d10 {IMAGE:403066}	403066 IMAGE Consortium	W81792

4886 F	3 021PA1EM	ESTs {IMAGE:403166}	403166 IMAGE Consortium	W82868
4887 F	5 021PA1EM	ESTs {IMAGE:403738}	403738 IMAGE Consortium	W82294
4888 F	7 021PA1EM	Mus musculus DAP-kinase related protein 1 mRNA, partial cds {IMAGE:403602}	403602 IMAGE Consortium	W82116
4889 F	9 021PA1EM	Plexin 3 {IMAGE:401253}	401253 IMAGE Consortium	W76838
4890 F	11 021PA1EM	Laminin, gamma 1 {IMAGE:401838}	401838 IMAGE Consortium	W82406
4891 H	1 021PA1EM	ESTs {IMAGE:419979}	419979 IMAGE Consortium	AA052668
4892 H	3 021PA1EM	Procollagen, type III, alpha 1 {IMAGE:420322}	420322 IMAGE Consortium	W89883
4893 H	5 021PA1EM	Replication factor C, 140 kDa {IMAGE:420302}	420302 IMAGE Consortium	W89873
4894 H	7 021PA1EM	IMAGE EST {IMAGE:420558}	420558 IMAGE Consortium	W89417
4895 H	9 021PA1EM	Mouse mRNA for small GTP-binding protein S10, complete cds {IMAGE:406177}	406177 IMAGE Consortium	W83866
4896 H	11 021PA1EM	ESTs {IMAGE:406561}	406561 IMAGE Consortium	W83394
4897 B	1 021WA1EN	IMAGE EST {IMAGE:400578}	400578 IMAGE Consortium	W76696
4898 B	3 021WA1EN	Proprotein convertase subtilisin/kexin type 5 {IMAGE:401263}	401263 IMAGE Consortium	W98241
4899 B	5 021WA1EN	ESTs, Highly similar to GLYCINE N-METHYLTRANSFERASE [Rattus norvegicus] {IMAGE:404574}	404574 IMAGE Consortium	W83078
4900 B	7 021WA1EN	Scavenger receptor class B1 {IMAGE:404540}	404540 IMAGE Consortium	W82738
4901 B	9 021WA1EN	IMAGE EST {IMAGE:405915}	405915 IMAGE Consortium	W83222
4902 B	11 021WA1EN	IMAGE EST {IMAGE:402559}	402559 IMAGE Consortium	W80120
4903 D	1 021WA1EN	IMAGE EST {IMAGE:405820}	405820 IMAGE Consortium	W83006
4904 D	3 021WA1EN	Mus musculus A6 protein tyrosine kinase homolog mRNA, complete cds {IMAGE:402345}	402345 IMAGE Consortium	W78591
4905 D	5 021WA1EN	Mus musculus mRNA for MD-1, complete cds {IMAGE:402964}	402964 IMAGE Consortium	W80147
4906 D	7 021WA1EN	Mouse mRNA for RPA16, complete cds {IMAGE:403327}	403327 IMAGE Consortium	W82822
4907 D	9 021WA1EN	IMAGE EST {IMAGE:400632}	400632 IMAGE Consortium	W75674
4908 D	11 021WA1EN	IMAGE EST {IMAGE:400571}	400571 IMAGE Consortium	W77518
4909 F	1 021WA1EN	Bone morphogenetic protein 1 {IMAGE:403597}	403597 IMAGE Consortium	W82677
4910 F	3 021WA1EN	IMAGE EST {IMAGE:403922}	403922 IMAGE Consortium	W82036
4911 F	5 021WA1EN	ESTs {IMAGE:401456}	401456 IMAGE Consortium	AA000873
4912 F	7 021WA1EN	ESTs {IMAGE:401906}	401906 IMAGE Consortium	W82161
4913 F	9 021WA1EN	Ufo oncogene homolog {IMAGE:401608}	401608 IMAGE Consortium	W78330
4914 F	11 021WA1EN	IMAGE EST {IMAGE:402072}	402072 IMAGE Consortium	W75898
4915 H	1 021WA1EN	IMAGE EST {IMAGE:404460}	404460 IMAGE Consortium	W83159
4916 H	3 021WA1EN	IMAGE EST {IMAGE:406085}	406085 IMAGE Consortium	W84046
4917 H	5 021WA1EN	IMAGE EST {IMAGE:402447}	402447 IMAGE Consortium	W79946
4918 H	7 021WA1EN	Mouse mRNA for osteoblast specific factor 2 (OSF-2) {IMAGE:403071}	403071 IMAGE Consortium	W81878
4919 H	9 021WA1EN	ESTs, Highly similar to The KIAA0144 gene product is novel. [H.sapiens] {IMAGE:403354}	403354 IMAGE Consortium	W81766
4920 H	11 021WA1EN	ESTs, Weakly similar to Gene product with similarity to	400308 IMAGE Consortium	W77405

4921 B	1 0213A1EO	KIAA0154 [H.sapiens] {IMAGE:400308} ESTs, Weakly similar to F39H11.3 [C.elegans] {IMAGE:409114}	409114 IMAGE Consortium	W87023
4922 B	3 0213A1EO	IMAGE EST {IMAGE:408959}	408959 IMAGE Consortium	AA052134
4923 B	5 0213A1EO	ESTs {IMAGE:418733}	418733 IMAGE Consortium	W96983
4924 B	7 0213A1EO	ESTs {IMAGE:406345}	406345 IMAGE Consortium	W83335
4925 B	9 0213A1EO	IMAGE EST {IMAGE:406239}	406239 IMAGE Consortium	W84023
4926 B	11 0213A1EO	ESTs, Moderately similar to KIAA0181 [H.sapiens] {IMAGE:406671}	406671 IMAGE Consortium	W83655
4927 D	1 0213A1EO	Mouse 19.5 mRNA, complete cds {IMAGE:420323}	420323 IMAGE Consortium	W89888
4928 D	3 0213A1EO	Cytochrome P450, 2e1, ethanol inducible {IMAGE:420040}	420040 IMAGE Consortium	W90893
4929 D	5 0213A1EO	ESTs {IMAGE:420528}	420528 IMAGE Consortium	W91443
4930 D	7 0213A1EO	Cytochrome P450, 2e1, ethanol inducible {IMAGE:406489}	406489 IMAGE Consortium	W83512
4931 D	9 0213A1EO	IMAGE EST {IMAGE:406193}	406193 IMAGE Consortium	W84058
4932 D	11 0213A1EO	Retinol binding protein 1, cellular {IMAGE:406897}	406897 IMAGE Consortium	W83609
4933 F	1 0213A1EO	ESTs, Highly similar to open reading frame [M.musculus] {IMAGE:420108}	420108 IMAGE Consortium	W89644
4934 F	3 0213A1EO	ESTs {IMAGE:420361}	420361 IMAGE Consortium	W91516
4935 F	5 0213A1EO	Avian erythroblastic leukemia viral (v-erb-a) oncogene homolog-like 3 {IMAGE:420535}	420535 IMAGE Consortium	W90859
4936 F	7 0213A1EO	ESTs {IMAGE:418606}	418606 IMAGE Consortium	W99195
4937 F	9 0213A1EO	IMAGE EST {IMAGE:406706}	406706 IMAGE Consortium	W83670
4938 F	11 0213A1EO	ESTs, Highly similar to N2,N2-DIMETHYLGUANOSINE TRNA METHYLTRANSFERASE PRECURSOR [Saccharomyces cerevisiae] {IMAGE:406889}	406889 IMAGE Consortium	W83899
4939 H	1 0213A1EO	IMAGE EST {IMAGE:418664}	418664 IMAGE Consortium	W96916
4940 H	3 0213A1EO	IMAGE EST {IMAGE:420631}	420631 IMAGE Consortium	W89301
4941 H	5 0213A1EO	ESTs {IMAGE:403263}	403263 IMAGE Consortium	W81972
4942 H	7 0213A1EO	RAD52 homolog, (S. cerevisiae) {IMAGE:419142}	419142 IMAGE Consortium	W88003
4943 H	9 0213A1EO	IMAGE EST {IMAGE:418864}	418864 IMAGE Consortium	AA052173
4944 H	11 0213A1EO	DIK1-like homolog (Drosophila) {IMAGE:407072}	407072 IMAGE Consortium	W83976
4945 B	1 021AA1EP	ESTs, Moderately similar to 13kD differentiation-associated protein [H.sapiens] {IMAGE:420017}	420017 IMAGE Consortium	W90880
4946 B	3 021AA1EP	IMAGE EST {IMAGE:420553}	420553 IMAGE Consortium	W89418
4947 B	5 021AA1EP	ESTs, Highly similar to PHOSPHORYLASE B KINASE BETA REGULATORY CHAIN [Oryctolagus cuniculus] {IMAGE:405880}	405880 IMAGE Consortium	W83326
4948 B	7 021AA1EP	ESTs {IMAGE:406171}	406171 IMAGE Consortium	W84055
4949 B	9 021AA1EP	ESTs {IMAGE:406805}	406805 IMAGE Consortium	W83777
4950 B	11 021AA1EP	IMAGE EST {IMAGE:406628}	406628 IMAGE Consortium	W83702
4951 D	1 021AA1EP	ESTs {IMAGE:420257}	420257 IMAGE Consortium	W89738
4952 D	3 021AA1EP	Mus musculus putative transcription factor mRNA, complete cds {IMAGE:420673}	420673 IMAGE Consortium	W89449

4953 D	5 021AA1EP	Cytochrome P450, 2e1, ethanol inducible {IMAGE:420656}	420656 IMAGE Consortium	W89701
4954 D	7 021AA1EP	ESTs {IMAGE:406351}	406351 IMAGE Consortium	W83381
4955 D	9 021AA1EP	Importin beta {IMAGE:419193}	419193 IMAGE Consortium	W87958
4956 D	11 021AA1EP	Mus musculus mRNA for DPM2, complete cds {IMAGE:406701}	406701 IMAGE Consortium	W83728
4957 F	1 021AA1EP	ESTs {IMAGE:418766}	418766 IMAGE Consortium	W97084
4958 F	3 021AA1EP	ESTs {IMAGE:406186}	406186 IMAGE Consortium	W83837
4959 F	5 021AA1EP	ESTs, Weakly similar to protein co-factor [M.musculus] {IMAGE:406173}	406173 IMAGE Consortium	W84056
4960 F	7 021AA1EP	Mus musculus tuftelin-interacting protein 10 mRNA, complete cds {IMAGE:406880}	406880 IMAGE Consortium	W83808
4961 F	9 021AA1EP	IMAGE EST {IMAGE:419478}	419478 IMAGE Consortium	W88090
4962 F	11 021AA1EP	ESTs {IMAGE:419440}	419440 IMAGE Consortium	W88188
4963 H	1 021AA1EP	ESTs, Moderately similar to PRE-MRNA SPLICING FACTOR SRP20 [Homo sapiens; Mus musculus] {IMAGE:406282}	406282 IMAGE Consortium	W83358
4964 H	3 021AA1EP	ESTs, Highly similar to NADH-UBIQUINONE OXIDOREDUCTASE B9 SUBUNIT [Bos taurus] {IMAGE:406509}	406509 IMAGE Consortium	W83574
4965 H	5 021AA1EP	Mus musculus metalloprotease-disintegrin MDC15 mRNA, complete cds {IMAGE:406689}	406689 IMAGE Consortium	W83666
4966 H	7 021AA1EP	ESTs, Highly similar to transcription factor IIIC alpha chain [R.norvegicus] {IMAGE:419263}	419263 IMAGE Consortium	W89677
4967 H	9 021AA1EP	ESTs, Highly similar to SH3 domain-containing protein SH3P17 [H.sapiens] {IMAGE:419417}	419417 IMAGE Consortium	W88185
4968 H	11 021AA1EP	IMAGE EST {IMAGE:407727}	407727 IMAGE Consortium	W85273
4969 B	1 021HA1EQ	CD9 antigen {IMAGE:421714}	421714 IMAGE Consortium	W98963
4970 B	3 021HA1EQ	EST {IMAGE:423531}	423531 IMAGE Consortium	W98905
4971 B	5 021HA1EQ	ESTs {IMAGE:422171}	422171 IMAGE Consortium	W96830
4972 B	7 021HA1EQ	ESTs {IMAGE:420975}	420975 IMAGE Consortium	W91135
4973 B	9 021HA1EQ	ESTs {IMAGE:422761}	422761 IMAGE Consortium	W97140
4974 B	11 021HA1EQ	IMAGE EST {IMAGE:422965}	422965 IMAGE Consortium	W97823
4975 D	1 021HA1EQ	ESTs {IMAGE:422953}	422953 IMAGE Consortium	W97980
4976 D	3 021HA1EQ	ESTs, Moderately similar to ZINC FINGER PROTEIN ZFP-36 [Homo sapiens] {IMAGE:421337}	421337 IMAGE Consortium	W97025
4977 D	5 021HA1EQ	ESTs {IMAGE:423170}	423170 IMAGE Consortium	W97210
4978 D	7 021HA1EQ	IMAGE EST {IMAGE:421811}	421811 IMAGE Consortium	W91395
4979 D	9 021HA1EQ	ESTs, Weakly similar to DNA-directed RNA polymerase [D.melanogaster] {IMAGE:423303}	423303 IMAGE Consortium	W98752
4980 D	11 021HA1EQ	IMAGE EST {IMAGE:422018}	422018 IMAGE Consortium	W96860
4981 F	1 021HA1EQ	IMAGE EST {IMAGE:421391}	421391 IMAGE Consortium	W97049
4982 F	3 021HA1EQ	IMAGE EST {IMAGE:423224}	423224 IMAGE Consortium	W97314
4983 F	5 021HA1EQ	ESTs {IMAGE:421766}	421766 IMAGE Consortium	W99015
4984 F	7 021HA1EQ	Mus musculus ion channel homolog RIC mRNA, complete cds {IMAGE:423334}	423334 IMAGE Consortium	W98807

4985 F	9 021HA1EQ	Erythroid Kruppel-like factor {IMAGE:422190}	422190 IMAGE Consortium	W97446
4986 F	11 021HA1EQ	ESTs {IMAGE:420520}	420520 IMAGE Consortium	W89688
4987 H	1 021HA1EQ	ESTs, Moderately similar to predicted using Genefinder [C.elegans] {IMAGE:423533}	423533 IMAGE Consortium	W98906
4988 H	3 021HA1EQ	ESTs, Highly similar to N2,N2-DIMETHYLGUANOSINE TRNA METHYLTRANSFERASE PRECURSOR [Saccharomyces cerevisiae] {IMAGE:423713}	423713 IMAGE Consortium	W98219
4989 H	5 021HA1EQ	IMAGE EST {IMAGE:420548}	420548 IMAGE Consortium	W89697
4990 H	7 021HA1EQ	ESTs {IMAGE:420905}	420905 IMAGE Consortium	AA052759
4991 H	9 021HA1EQ	ESTs, Highly similar to DIHYDROLIPOAMIDE ACETYLTRANSFERASE COMPONENT [Rattus norvegicus] {IMAGE:422907}	422907 IMAGE Consortium	W97904
4992 H	11 021HA1EQ	ESTs {IMAGE:423219}	423219 IMAGE Consortium	W97274
4993 B	1 021OA1ER	ESTs {IMAGE:418891}	418891 IMAGE Consortium	W89396
4994 B	3 021OA1ER	IMAGE EST {IMAGE:406983}	406983 IMAGE Consortium	W83952
4995 B	5 021OA1ER	IMAGE EST {IMAGE:419832}	419832 IMAGE Consortium	W89583
4996 B	7 021OA1ER	IMAGE EST {IMAGE:407704}	407704 IMAGE Consortium	W85662
4997 B	9 021OA1ER	IMAGE EST {IMAGE:408400}	408400 IMAGE Consortium	W85376
4998 B	11 021OA1ER	IMAGE EST {IMAGE:408679}	408679 IMAGE Consortium	W85015
4999 D	1 021OA1ER	Mouse spi2 proteinase inhibitor (spi2/eb1) mRNA, 3' end {IMAGE:406295}	406295 IMAGE Consortium	W83447
5000 D	3 021OA1ER	ESTs {IMAGE:406549}	406549 IMAGE Consortium	W83593
5001 D	5 021OA1ER	COPROPORPHYRINOGEN III OXIDASE PRECURSOR {IMAGE:419503}	419503 IMAGE Consortium	W87981
5002 D	7 021OA1ER	ESTs, Moderately similar to CYTOCHROME C OXIDASE POLYPEPTIDE VIB [Homo sapiens] {IMAGE:407551}	407551 IMAGE Consortium	W88135
5003 D	9 021OA1ER	IMAGE EST {IMAGE:408775}	408775 IMAGE Consortium	W85481
5004 D	11 021OA1ER	ESTs {IMAGE:409109}	409109 IMAGE Consortium	W87081
5005 F	1 021OA1ER	IMAGE EST {IMAGE:423177}	423177 IMAGE Consortium	W97303
5006 F	3 021OA1ER	Polypyrimidine tract binding protein {IMAGE:423349}	423349 IMAGE Consortium	W98869
5007 F	5 021OA1ER	M.musculus mRNA for semaphorin D {IMAGE:423697}	423697 IMAGE Consortium	W98303
5008 F	7 021OA1ER	EST {IMAGE:421977}	421977 IMAGE Consortium	W98958
5009 F	9 021OA1ER	IMAGE EST {IMAGE:421965}	421965 IMAGE Consortium	W97408
5010 F	11 021OA1ER	IMAGE EST {IMAGE:423010}	423010 IMAGE Consortium	W97985
5011 H	1 021OA1ER	ESTs {IMAGE:422434}	422434 IMAGE Consortium	W97660
5012 H	3 021OA1ER	IMAGE EST {IMAGE:421449}	421449 IMAGE Consortium	W99090
5013 H	5 021OA1ER	ESTs, Moderately similar to HYPOTHETICAL PROTEIN KIAA0076 [H.sapiens] {IMAGE:423035}	423035 IMAGE Consortium	W98022
5014 H	7 021OA1ER	Mus musculus mRNA for actin-related protein 1 alpha-isoform, complete cds {IMAGE:423120}	423120 IMAGE Consortium	W97866
5015 H	9 021OA1ER	Recombining binding protein suppressor of hairless (Drosophila) {IMAGE:421546}	421546 IMAGE Consortium	W97158
5016 H	11 021OA1ER	Amyloid beta (A4) precursor-like protein 1 {IMAGE:421741}	421741 IMAGE Consortium	W98956

5017 B	1	021VA1ES	Axin {IMAGE:422623}	422623	IMAGE Consortium	W97777
5018 B	3	021VA1ES	IMAGE EST {IMAGE:422912}	422912	IMAGE Consortium	W97901
5019 B	5	021VA1ES	IMAGE EST {IMAGE:421712}	421712	IMAGE Consortium	W98962
5020 B	7	021VA1ES	ESTs, Weakly similar to Yer134cp [S.cerevisiae] {IMAGE:421884}	421884	IMAGE Consortium	W91412
5021 B	9	021VA1ES	ESTs {IMAGE:423233}	423233	IMAGE Consortium	W97169
5022 B	11	021VA1ES	ESTs {IMAGE:423383}	423383	IMAGE Consortium	W98675
5023 D	1	021VA1ES	MAJOR PRION PROTEIN PRECURSOR {IMAGE:421749}	421749	IMAGE Consortium	W99102
5024 D	3	021VA1ES	IMAGE EST {IMAGE:422167}	422167	IMAGE Consortium	W97387
5025 D	5	021VA1ES	ESTs {IMAGE:420741}	420741	IMAGE Consortium	W89437
5026 D	7	021VA1ES	ESTs, Moderately similar to APC-binding protein EB1 homolog [M.musculus] {IMAGE:422813}	422813	IMAGE Consortium	W97177
5027 D	9	021VA1ES	Mus musculus mRNA for ganglioside-induced differentiation associated protein 10 {IMAGE:423021}	423021	IMAGE Consortium	W97235
5028 D	11	021VA1ES	ESTs, Highly similar to putative hydrophobic domain in amino acid positions 373-390. [H.sapiens] {IMAGE:423153}	423153	IMAGE Consortium	W97298
5029 F	1	021VA1ES	Cytochrome P450, 2e1, ethanol inducible {IMAGE:426537}	426537	IMAGE Consortium	AA002396
5030 F	3	021VA1ES	IMAGE EST {IMAGE:425664}	425664	IMAGE Consortium	AA000282
5031 F	5	021VA1ES	IMAGE EST {IMAGE:425409}	425409	IMAGE Consortium	AA000134
5032 F	7	021VA1ES	IMAGE EST {IMAGE:425491}	425491	IMAGE Consortium	AA000265
5033 F	9	021VA1ES	ESTs {IMAGE:426059}	426059	IMAGE Consortium	AA002845
5034 F	11	021VA1ES	ESTs {IMAGE:424921}	424921	IMAGE Consortium	AA000671
5035 H	1	021VA1ES	IMAGE EST {IMAGE:427002}	427002	IMAGE Consortium	AA003275
5036 H	3	021VA1ES	EST {IMAGE:427345}	427345	IMAGE Consortium	AA003152
5037 H	5	021VA1ES	Homeo box B9 {IMAGE:423995}	423995	IMAGE Consortium	W91526
5038 H	7	021VA1ES	IMAGE EST {IMAGE:425742}	425742	IMAGE Consortium	AA000618
5039 H	9	021VA1ES	IMAGE EST {IMAGE:425916}	425916	IMAGE Consortium	AA000764
5040 H	11	021VA1ES	Beta-glucuronidase structural {IMAGE:425969}	425969	IMAGE Consortium	AA000856
5041 B	1	0212A1ET	IMAGE EST {IMAGE:423948}	423948	IMAGE Consortium	W91510
5042 B	3	0212A1ET	ESTs {IMAGE:425934}	425934	IMAGE Consortium	AA000626
5043 B	5	0212A1ET	ESTs {IMAGE:424384}	424384	IMAGE Consortium	W98487
5044 B	7	0212A1ET	FBJ osteosarcoma oncogene {IMAGE:426070}	426070	IMAGE Consortium	AA002910
5045 B	9	0212A1ET	IMAGE EST {IMAGE:426211}	426211	IMAGE Consortium	AA003021
5046 B	11	0212A1ET	Troponin I, skeletal, fast 2 {IMAGE:426504}	426504	IMAGE Consortium	AA002376
5047 D	1	0212A1ET	Wingless-related MMTV integration site 3A {IMAGE:426103}	426103	IMAGE Consortium	AA000971
5048 D	3	0212A1ET	ESTs {IMAGE:425803}	425803	IMAGE Consortium	AA000807
5049 D	5	0212A1ET	EST {IMAGE:425807}	425807	IMAGE Consortium	AA000809
5050 D	7	0212A1ET	ESTs {IMAGE:426297}	426297	IMAGE Consortium	AA002955
5051 D	9	0212A1ET	ESTs {IMAGE:424677}	424677	IMAGE Consortium	AA000655
5052 D	11	0212A1ET	IMAGE EST {IMAGE:425342}	425342	IMAGE Consortium	W99957
5053 F	1	0212A1ET	ESTs, Highly similar to KIAA0103 [H.sapiens] {IMAGE:426345}	426345	IMAGE Consortium	AA003181
5054 F	3	0212A1ET	ESTs {IMAGE:426140}	426140	IMAGE Consortium	AA003120

5055 F	5 0212A1ET	ESTs {IMAGE:426712}	426712 IMAGE Consortium	AA002757
5056 F	7 0212A1ET	Complement component 1 inhibitor {IMAGE:426886}	426886 IMAGE Consortium	AA003214
5057 F	9 0212A1ET	ESTs {IMAGE:427205}	427205 IMAGE Consortium	AA002968
5058 F	11 0212A1ET	IMAGE EST {IMAGE:427094}	427094 IMAGE Consortium	AA003377
5059 H	1 0212A1ET	ESTs, Moderately similar to PROBABLE PEPTIDYL-PROLYL CIS-TRANS ISOMERASE C21E11.05C [Schizosaccharomyces pombe] {IMAGE:424649}	424649 IMAGE Consortium	AA000646
5060 H	3 0212A1ET	ESTs {IMAGE:426212}	426212 IMAGE Consortium	AA003144
5061 H	5 0212A1ET	ESTs {IMAGE:425200}	425200 IMAGE Consortium	W99975
5062 H	7 0212A1ET	IMAGE EST {IMAGE:425013}	425013 IMAGE Consortium	AA000059
5063 H	9 0212A1ET	ESTs {IMAGE:425619}	425619 IMAGE Consortium	AA000356
5064 H	11 0212A1ET	IMAGE EST {IMAGE:425578}	425578 IMAGE Consortium	AA000018
5065 B	1 0219A1EU	ESTs, Weakly similar to HYPOTHETICAL 117.9 KD PROTEIN IN FKH1-STH1 INTERGENIC REGION [S.cerevisiae] {IMAGE:427295}	427295 IMAGE Consortium	AA003260
5066 B	3 0219A1EU	IMAGE EST {IMAGE:439033}	439033 IMAGE Consortium	AA008240
5067 B	5 0219A1EU	IMAGE EST {IMAGE:436967}	436967 IMAGE Consortium	AA002716
5068 B	7 0219A1EU	IMAGE EST {IMAGE:437335}	437335 IMAGE Consortium	AA003687
5069 B	9 0219A1EU	Protein tyrosine phosphatase, non-receptor type 2 {IMAGE:437517}	437517 IMAGE Consortium	AA003599
5070 B	11 0219A1EU	ESTs {IMAGE:437673}	437673 IMAGE Consortium	AA007771
5071 D	1 0219A1EU	IMAGE EST {IMAGE:438478}	438478 IMAGE Consortium	AA008038
5072 D	3 0219A1EU	ESTs {IMAGE:436886}	436886 IMAGE Consortium	AA002611
5073 D	5 0219A1EU	ESTs, Moderately similar to rA9 [R.norvegicus] {IMAGE:437503}	437503 IMAGE Consortium	AA003764
5074 D	7 0219A1EU	IMAGE EST {IMAGE:439445}	439445 IMAGE Consortium	AA004095
5075 D	9 0219A1EU	IMAGE EST {IMAGE:439210}	439210 IMAGE Consortium	AA003881
5076 D	11 0219A1EU	IMAGE EST {IMAGE:439429}	439429 IMAGE Consortium	AA004075
5077 F	1 0219A1EU	IMAGE EST {IMAGE:437030}	437030 IMAGE Consortium	AA002812
5078 F	3 0219A1EU	ESTs {IMAGE:437288}	437288 IMAGE Consortium	AA003739
5079 F	5 0219A1EU	IMAGE EST {IMAGE:439517}	439517 IMAGE Consortium	AA004129
5080 F	7 0219A1EU	Mus musculus mitogen- and stress-activated protein kinase-2 (mMSK2) mRNA, complete cds {IMAGE:439453}	439453 IMAGE Consortium	AA004099
5081 F	9 0219A1EU	Mus musculus transcription factor PBX2 (PBX2) mRNA, complete cds {IMAGE:439268}	439268 IMAGE Consortium	AA003969
5082 F	11 0219A1EU	ESTs {IMAGE:439925}	439925 IMAGE Consortium	AA008734
5083 H	1 0219A1EU	Mus musculus ETO/MTG8-related protein ETO-2 mRNA, complete cds {IMAGE:439477}	439477 IMAGE Consortium	AA004178
5084 H	3 0219A1EU	Neurotrophic tyrosine kinase, receptor, type 3 {IMAGE:438183}	438183 IMAGE Consortium	AA008069
5085 H	5 0219A1EU	Mus musculus COP9 complex subunit 4 (COPS4) mRNA, complete cds {IMAGE:438210}	438210 IMAGE Consortium	AA008135
5086 H	7 0219A1EU	IMAGE EST {IMAGE:438747}	438747 IMAGE Consortium	AA050945

5087 H	9 0219A1EU	Mus musculus protein-tyrosine phosphatase mRNA, complete cds {IMAGE:427642}	427642 IMAGE Consortium	AA002437
5088 H	11 0219A1EU	IMAGE EST {IMAGE:438569}	438569 IMAGE Consortium	AA051060
5089 B	1 021GA1EV	IMAGE EST {IMAGE:425698}	425698 IMAGE Consortium	AA000313
5090 B	3 021GA1EV	ESTs {IMAGE:427563}	427563 IMAGE Consortium	AA002532
5091 B	5 021GA1EV	ESTs {IMAGE:425786}	425786 IMAGE Consortium	AA000543
5092 B	7 021GA1EV	IMAGE EST {IMAGE:425799}	425799 IMAGE Consortium	AA000696
5093 B	9 021GA1EV	IMAGE EST {IMAGE:425757}	425757 IMAGE Consortium	AA000794
5094 B	11 021GA1EV	IMAGE EST {IMAGE:426179}	426179 IMAGE Consortium	AA003011
5095 D	1 021GA1EV	Vasoactive intestinal peptide receptor 2 {IMAGE:425925}	425925 IMAGE Consortium	AA000726
5096 D	3 021GA1EV	ESTs {IMAGE:426395}	426395 IMAGE Consortium	AA003191
5097 D	5 021GA1EV	Mus musculus type 6 nucleoside diphosphate kinase NM23-M6 (Nm23-M6) mRNA, complete cds {IMAGE:424871}	424871 IMAGE Consortium	AA000602
5098 D	7 021GA1EV	IMAGE EST {IMAGE:425194}	425194 IMAGE Consortium	AA000073
5099 D	9 021GA1EV	Mouse mRNA for RPA16, complete cds {IMAGE:425081}	425081 IMAGE Consortium	W99934
5100 D	11 021GA1EV	ESTs, Highly similar to 40S RIBOSOMAL PROTEIN S4, X ISOFORM [Homo sapiens; Rattus norvegicus; Mus musculus; Mesocricetus auratus; Felis catus] {IMAGE:425352}	425352 IMAGE Consortium	AA000082
5101 F	1 021GA1EV	ESTs, Highly similar to MITOTIC MAD2 PROTEIN [Saccharomyces cerevisiae] {IMAGE:426406}	426406 IMAGE Consortium	AA002895
5102 F	3 021GA1EV	IMAGE EST {IMAGE:426511}	426511 IMAGE Consortium	AA002385
5103 F	5 021GA1EV	ESTs {IMAGE:425525}	425525 IMAGE Consortium	AA000224
5104 F	7 021GA1EV	EST {IMAGE:425442}	425442 IMAGE Consortium	AA000167
5105 F	9 021GA1EV	Wingless-related MMTV integration site 5B {IMAGE:423942}	423942 IMAGE Consortium	W91468
5106 F	11 021GA1EV	ESTs {IMAGE:424123}	424123 IMAGE Consortium	W98084
5107 H	1 021GA1EV	IMAGE EST {IMAGE:426890}	426890 IMAGE Consortium	AA003252
5108 H	3 021GA1EV	Mus musculus major histocompatibility locus class II region; Fas-binding protein Daxx (DAXX) gene, partial cds; Bing1 (BING1), tapasin (tapasin), RalGDS-like factor (RLF), KE2 (KE2), BING4 (BING4), beta1, 3-galactosyl transferase {IMAGE:425385}	425385 IMAGE Consortium	AA000107
5109 H	5 021GA1EV	IMAGE EST {IMAGE:425562}	425562 IMAGE Consortium	AA000227
5110 H	7 021GA1EV	IMAGE EST {IMAGE:427312}	427312 IMAGE Consortium	AA003261
5111 H	9 021GA1EV	ESTs {IMAGE:437235}	437235 IMAGE Consortium	AA003456
5112 H	11 021GA1EV	ESTs, Moderately similar to hypothetical human serine-threonine protein kinase R31240_1 [H.sapiens] {IMAGE:436939}	436939 IMAGE Consortium	AA002708
5113 B	1 021NA1EW	ESTs {IMAGE:437755}	437755 IMAGE Consortium	AA008001
5114 B	3 021NA1EW	ESTs {IMAGE:438130}	438130 IMAGE Consortium	AA008052
5115 B	5 021NA1EW	Mus musculus docking protein Dok-R mRNA, complete cds {IMAGE:438282}	438282 IMAGE Consortium	AA008417
5116 B	7 021NA1EW	ESTs {IMAGE:438412}	438412 IMAGE Consortium	AA008296
5117 B	9 021NA1EW	ESTs {IMAGE:427283}	427283 IMAGE Consortium	AA003094



5118 B	11 021NA1EW	ESTs, Weakly similar to HYPOTHETICAL 15.1 KD PROTEIN IN PET494-MSO1 INTERGENIC REGION [ <i>S.cerevisiae</i> ] {IMAGE:437209}	437209 IMAGE Consortium	AA003451
5119 D	1 021NA1EW	IMAGE EST {IMAGE:438154}	438154 IMAGE Consortium	AA008057
5120 D	3 021NA1EW	IMAGE EST {IMAGE:438351}	438351 IMAGE Consortium	AA008154
5121 D	5 021NA1EW	ESTs {IMAGE:439665}	439665 IMAGE Consortium	AA014509
5122 D	7 021NA1EW	IMAGE EST {IMAGE:439621}	439621 IMAGE Consortium	AA008693
5123 D	9 021NA1EW	Dlk1-like homolog ( <i>Drosophila</i> ) {IMAGE:427339}	427339 IMAGE Consortium	AA003303
5124 D	11 021NA1EW	IMAGE EST {IMAGE:437104}	437104 IMAGE Consortium	AA003402
5125 F	1 021NA1EW	ESTs, Weakly similar to retinal-specific heterotrimeric GTP-binding protein beta subunit, G beta5L [ <i>M.musculus</i> ] {IMAGE:436956}	436956 IMAGE Consortium	AA002632
5126 F	3 021NA1EW	IMAGE EST {IMAGE:437594}	437594 IMAGE Consortium	AA003561
5127 F	5 021NA1EW	IMAGE EST {IMAGE:439542}	439542 IMAGE Consortium	AA004173
5128 F	7 021NA1EW	IMAGE EST {IMAGE:439379}	439379 IMAGE Consortium	AA004054
5129 F	9 021NA1EW	ESTs, Highly similar to HYPOTHETICAL 39.5 KD PROTEIN C12G12.06C IN CHROMOSOME I [ <i>Schizosaccharomyces pombe</i> ] {IMAGE:437831}	437831 IMAGE Consortium	AA007803
5130 F	11 021NA1EW	IMAGE EST {IMAGE:438232}	438232 IMAGE Consortium	AA008381
5131 H	1 021NA1EW	IMAGE EST {IMAGE:437564}	437564 IMAGE Consortium	AA003773
5132 H	3 021NA1EW	IMAGE EST {IMAGE:439427}	439427 IMAGE Consortium	AA004074
5133 H	5 021NA1EW	IMAGE EST {IMAGE:439389}	439389 IMAGE Consortium	AA004105
5134 H	7 021NA1EW	IMAGE EST {IMAGE:438304}	438304 IMAGE Consortium	AA008424
5135 H	9 021NA1EW	<i>Mus musculus</i> AP-3 complex sigma3B subunit mRNA, complete cds {IMAGE:438481}	438481 IMAGE Consortium	AA008255
5136 H	11 021NA1EW	IMAGE EST {IMAGE:438725}	438725 IMAGE Consortium	AA051124
5137 B	1 021UA1EX	Retinoic acid receptor, gamma {IMAGE:439731}	439731 IMAGE Consortium	AA014382
5138 B	3 021UA1EX	ESTs {IMAGE:438745}	438745 IMAGE Consortium	AA050944
5139 B	5 021UA1EX	IMAGE EST {IMAGE:438749}	438749 IMAGE Consortium	AA051080
5140 B	7 021UA1EX	IMAGE EST {IMAGE:427574}	427574 IMAGE Consortium	AA002530
5141 B	9 021UA1EX	ESTs {IMAGE:427652}	427652 IMAGE Consortium	AA002704
5142 B	11 021UA1EX	ESTs {IMAGE:437206}	437206 IMAGE Consortium	AA003605
5143 D	1 021UA1EX	IMAGE EST {IMAGE:441293}	441293 IMAGE Consortium	AA009195
5144 D	3 021UA1EX	IMAGE EST {IMAGE:441323}	441323 IMAGE Consortium	AA009265
5145 D	5 021UA1EX	IMAGE EST {IMAGE:441807}	441807 IMAGE Consortium	AA014053
5146 D	7 021UA1EX	ESTs, Highly similar to PLATELET FACTOR 4 PRECURSOR [ <i>Rattus norvegicus</i> ] {IMAGE:440563}	440563 IMAGE Consortium	AA014819
5147 D	9 021UA1EX	ESTs, Moderately similar to HYPOTHETICAL 27.1 KD PROTEIN CCE1-CAP1 INTERGENIC REGION [ <i>Saccharomyces cerevisiae</i> ] {IMAGE:440795}	440795 IMAGE Consortium	AA009187
5148 D	11 021UA1EX	IMAGE EST {IMAGE:440879}	440879 IMAGE Consortium	AA008851
5149 F	1 021UA1EX	IMAGE EST {IMAGE:441596}	441596 IMAGE Consortium	AA014048
5150 F	3 021UA1EX	<i>Mus musculus</i> major histocompatibility locus class III	440840 IMAGE Consortium	AA008747

		region:butyrophilin-like protein gene, partial cds; Notch4, PBX2, RAGE, lysophatidic acid acyl transferase-alpha, palmitoyl-protein thioesterase 2 (PPT2), CREB-RP, and tenascin {IMAGE:440840}		
5151 F	5 021UA1EX	3,2-TRANS-ENOYL-COA ISOMERASE, MITOCHONDRIAL PRECURSOR {IMAGE:440895}	440895 IMAGE Consortium	AA008767
5152 F	7 021UA1EX	ESTs, Moderately similar to FERRITIN HEAVY CHAIN [Cricetulus griseus] {IMAGE:442277}	442277 IMAGE Consortium	AA013495
5153 F	9 021UA1EX	IMAGE EST {IMAGE:439671}	439671 IMAGE Consortium	AA014305
5154 F	11 021UA1EX	IMAGE EST {IMAGE:440100}	440100 IMAGE Consortium	AA016346
5155 H	1 021UA1EX	ESTs, Moderately similar to PROTEIN PHOSPHATASE 2C ALPHA ISOFORM [Mus musculus] {IMAGE:441153}	441153 IMAGE Consortium	AA009055
5156 H	3 021UA1EX	ESTs {IMAGE:441602}	441602 IMAGE Consortium	AA013940
5157 H	5 021UA1EX	ESTs, Moderately similar to PLASMA-CELL MEMBRANE GLYCOPROTEIN PC-1 [Homo sapiens] {IMAGE:440103}	440103 IMAGE Consortium	AA016397
5158 H	7 021UA1EX	GLUTAMINE SYNTHETASE {IMAGE:440344}	440344 IMAGE Consortium	AA011759
5159 H	9 021UA1EX	IMAGE EST {IMAGE:440715}	440715 IMAGE Consortium	AA009123
5160 H	11 021UA1EX	HU-antigen D {IMAGE:441193}	441193 IMAGE Consortium	AA011823
5161 B	1 0211A1EY	Mus musculus elongation factor 1-beta homolog mRNA, complete cds {IMAGE:442662}	442662 IMAGE Consortium	AA015145
5162 B	3 0211A1EY	M.musculus rhoC mRNA {IMAGE:442818}	442818 IMAGE Consortium	AA036489
5163 B	5 0211A1EY	IMAGE EST {IMAGE:445310}	445310 IMAGE Consortium	AA017768
5164 B	7 0211A1EY	IMAGE EST {IMAGE:445688}	445688 IMAGE Consortium	AA017868
5165 B	9 0211A1EY	IMAGE EST {IMAGE:445573}	445573 IMAGE Consortium	AA017931
5166 B	11 0211A1EY	IMAGE EST {IMAGE:446075}	446075 IMAGE Consortium	AA020208
5167 D	1 0211A1EY	Myosin light chain, alkali, cardiac atria {IMAGE:444059}	444059 IMAGE Consortium	AA014929
5168 D	3 0211A1EY	ESTs {IMAGE:444087}	444087 IMAGE Consortium	AA014939
5169 D	5 0211A1EY	ESTs {IMAGE:442783}	442783 IMAGE Consortium	AA035848
5170 D	7 0211A1EY	Mouse mRNA for mMIS5, complete cds {IMAGE:444918}	444918 IMAGE Consortium	AA016759
5171 D	9 0211A1EY	IMAGE EST {IMAGE:444944}	444944 IMAGE Consortium	AA016408
5172 D	11 0211A1EY	Tumor necrosis factor induced protein 2 {IMAGE:443884}	443884 IMAGE Consortium	AA014136
5173 F	1 0211A1EY	Mus musculus deubiquitinating enzyme (UBH1) mRNA, partial cds {IMAGE:443554}	443554 IMAGE Consortium	AA013513
5174 F	3 0211A1EY	ESTs {IMAGE:445450}	445450 IMAGE Consortium	AA018015
5175 F	5 0211A1EY	EST {IMAGE:443820}	443820 IMAGE Consortium	AA014086
5176 F	7 0211A1EY	Delta-like 1 homolog (Drosophila) {IMAGE:444023}	444023 IMAGE Consortium	AA015264
5177 F	9 0211A1EY	Ferritin heavy chain {IMAGE:443046}	443046 IMAGE Consortium	AA014230
5178 F	11 0211A1EY	IMAGE EST {IMAGE:443916}	443916 IMAGE Consortium	AA014102
5179 H	1 0211A1EY	Mus musculus mRNA for Mad-related protein Smad7 {IMAGE:457221}	457221 IMAGE Consortium	AA022262
5180 H	3 0211A1EY	IMAGE EST {IMAGE:455385}	455385 IMAGE Consortium	AA020436
5181 H	5 0211A1EY	ESTs {IMAGE:457533}	457533 IMAGE Consortium	AA021792
5182 H	7 0211A1EY	ESTs, Highly similar to ANKYRIN, BRAIN VARIANT 2 [Homo	457954 IMAGE Consortium	AA030190

		sapiens] {IMAGE:457954}		
5183 H	9 0211A1EY	IMAGE EST {IMAGE:457892}	457892 IMAGE Consortium	AA030169
5184 H	11 0211A1EY	Very low density lipoprotein receptor {IMAGE:446084}	446084 IMAGE Consortium	AA020307
5185 B	1 0218A1EZ	IMAGE EST {IMAGE:442201}	442201 IMAGE Consortium	AA013653
5186 B	3 0218A1EZ	Cytochrome P450, 2e1, ethanol inducible {IMAGE:442004}	442004 IMAGE Consortium	AA015423
5187 B	5 0218A1EZ	IMAGE EST {IMAGE:440240}	440240 IMAGE Consortium	AA016474
5188 B	7 0218A1EZ	IMAGE EST {IMAGE:441880}	441880 IMAGE Consortium	AA013766
5189 B	9 0218A1EZ	IMAGE EST {IMAGE:440733}	440733 IMAGE Consortium	AA009223
5190 B	11 0218A1EZ	ESTs {IMAGE:439941}	439941 IMAGE Consortium	AA008573
5191 D	1 0218A1EZ	IMAGE EST {IMAGE:441766}	441766 IMAGE Consortium	AA013977
5192 D	3 0218A1EZ	ESTs, Highly similar to SPLICING FACTOR U2AF 35 KD SUBUNIT [H.sapiens] {IMAGE:440639}	440639 IMAGE Consortium	AA015221
5193 D	5 0218A1EZ	Stimulated by retinoic acid gene 6 {IMAGE:442048}	442048 IMAGE Consortium	AA015434
5194 D	7 0218A1EZ	IMAGE EST {IMAGE:441227}	441227 IMAGE Consortium	AA011838
5195 D	9 0218A1EZ	IMAGE EST {IMAGE:440132}	440132 IMAGE Consortium	AA016627
5196 D	11 0218A1EZ	ESTs {IMAGE:440634}	440634 IMAGE Consortium	AA015373
5197 F	1 0218A1EZ	ESTs {IMAGE:443857}	443857 IMAGE Consortium	AA014128
5198 F	3 0218A1EZ	ESTs, Highly similar to checkpoint suppressor 1 [H.sapiens] {IMAGE:444383}	444383 IMAGE Consortium	AA016382
5199 F	5 0218A1EZ	ESTs {IMAGE:443413}	443413 IMAGE Consortium	AA013835
5200 F	7 0218A1EZ	IMAGE EST {IMAGE:445667}	445667 IMAGE Consortium	AA017867
5201 F	9 0218A1EZ	IMAGE EST {IMAGE:445874}	445874 IMAGE Consortium	AA020125
5202 F	11 0218A1EZ	ESTs {IMAGE:445809}	445809 IMAGE Consortium	AA020151
5203 H	1 0218A1EZ	Erythrocyte protein band 4.1 {IMAGE:444037}	444037 IMAGE Consortium	AA014918
5204 H	3 0218A1EZ	IMAGE EST {IMAGE:444449}	444449 IMAGE Consortium	AA016923
5205 H	5 0218A1EZ	ESTs {IMAGE:443276}	443276 IMAGE Consortium	AA015504
5206 H	7 0218A1EZ	IMAGE EST {IMAGE:444007}	444007 IMAGE Consortium	AA014889
5207 H	9 0218A1EZ	ESTs, Weakly similar to CYSTATIN C PRECURSOR [M.musculus] {IMAGE:443798}	443798 IMAGE Consortium	AA014980
5208 H	11 0218A1EZ	IMAGE EST {IMAGE:442460}	442460 IMAGE Consortium	AA015288
5209 B	1 0218A1F0	IMAGE EST {IMAGE:455836}	455836 IMAGE Consortium	AA023797
5210 B	3 0218A1F0	ESTs {IMAGE:457703}	457703 IMAGE Consortium	AA021959
5211 B	5 0218A1F0	ESTs, Weakly similar to Similarity to Yeast hypothetical 65.2 KD protein [C.elegans] {IMAGE:456398}	456398 IMAGE Consortium	AA023316
5212 B	7 0218A1F0	IMAGE EST {IMAGE:459002}	459002 IMAGE Consortium	AA024072
5213 B	9 0218A1F0	IMAGE EST {IMAGE:455522}	455522 IMAGE Consortium	AA020522
5214 B	11 0218A1F0	Polycystic kidney disease 2 {IMAGE:455813}	455813 IMAGE Consortium	AA023786
5215 D	1 0218A1F0	IMAGE EST {IMAGE:455519}	455519 IMAGE Consortium	AA020602
5216 D	3 0218A1F0	ESTs {IMAGE:457767}	457767 IMAGE Consortium	AA021778
5217 D	5 0218A1F0	ESTs, Weakly similar to MEL-11 [C.elegans] {IMAGE:455598}	455598 IMAGE Consortium	AA020619
5218 D	7 0218A1F0	ESTs {IMAGE:457601}	457601 IMAGE Consortium	AA021806
5219 D	9 0218A1F0	EST {IMAGE:456294}	456294 IMAGE Consortium	AA023300

5220 D	11 0218A1F0	IMAGE EST {IMAGE:458555}	458555 IMAGE Consortium	AA023888
5221 F	1 0218A1F0	IMAGE EST {IMAGE:456831}	456831 IMAGE Consortium	AA023545
5222 F	3 0218A1F0	IMAGE EST {IMAGE:455432}	455432 IMAGE Consortium	AA020478
5223 F	5 0218A1F0	IMAGE EST {IMAGE:457455}	457455 IMAGE Consortium	AA021668
5224 F	7 0218A1F0	ESTs {IMAGE:455880}	455880 IMAGE Consortium	AA023058
5225 F	9 0218A1F0	IMAGE EST {IMAGE:455792}	455792 IMAGE Consortium	AA023119
5226 F	11 0218A1F0	ESTs {IMAGE:456344}	456344 IMAGE Consortium	AA023339
5227 H	1 0218A1F0	IMAGE EST {IMAGE:456234}	456234 IMAGE Consortium	AA023263
5228 H	3 0218A1F0	Mus musculus IER5 (Ier5) mRNA, complete cds {IMAGE:456536}	456536 IMAGE Consortium	AA023439
5229 H	5 0218A1F0	ESTs, Moderately similar to AMILORIDE-SENSITIVE AMINE OXIDASE [Rattus norvegicus] {IMAGE:456716}	456716 IMAGE Consortium	AA023491
5230 H	7 0218A1F0	IMAGE EST {IMAGE:459060}	459060 IMAGE Consortium	AA024260
5231 H	9 0218A1F0	IMAGE EST {IMAGE:455391}	455391 IMAGE Consortium	AA020437
5232 H	11 0218A1F0	ESTs {IMAGE:455371}	455371 IMAGE Consortium	AA020415
5233 B	1 021FA1F1	IMAGE EST {IMAGE:464405}	464405 IMAGE Consortium	AA030391
5234 B	3 021FA1F1	Mannosidase 2, alpha B1 {IMAGE:459599}	459599 IMAGE Consortium	AA027528
5235 B	5 021FA1F1	T-box 15 {IMAGE:463070}	463070 IMAGE Consortium	AA027675
5236 B	7 021FA1F1	IMAGE EST {IMAGE:463337}	463337 IMAGE Consortium	AA028697
5237 B	9 021FA1F1	IMAGE EST {IMAGE:464871}	464871 IMAGE Consortium	AA030527
5238 B	11 021FA1F1	ESTs, Moderately similar to bone-derived growth factor [H.sapiens] {IMAGE:458802}	458802 IMAGE Consortium	AA024091
5239 D	1 021FA1F1	ESTs {IMAGE:464728}	464728 IMAGE Consortium	AA030770
5240 D	3 021FA1F1	ESTs {IMAGE:465240}	465240 IMAGE Consortium	AA032564
5241 D	5 021FA1F1	Mus musculus CW17 gene {IMAGE:465005}	465005 IMAGE Consortium	AA032448
5242 D	7 021FA1F1	IMAGE EST {IMAGE:458874}	458874 IMAGE Consortium	AA024223
5243 D	9 021FA1F1	IMAGE EST {IMAGE:463860}	463860 IMAGE Consortium	AA028539
5244 D	11 021FA1F1	ESTs {IMAGE:463771}	463771 IMAGE Consortium	AA028368
5245 F	1 021FA1F1	IMAGE EST {IMAGE:465073}	465073 IMAGE Consortium	AA032406
5246 F	3 021FA1F1	IMAGE EST {IMAGE:463535}	463535 IMAGE Consortium	AA028635
5247 F	5 021FA1F1	ESTs, Highly similar to SINGLE-STRANDED DNA-BINDING PROTEIN, MITOCHONDRIAL PRECURSOR [Rattus norvegicus] {IMAGE:459094}	459094 IMAGE Consortium	AA024026
5248 F	7 021FA1F1	IMAGE EST {IMAGE:463819}	463819 IMAGE Consortium	AA028383
5249 F	9 021FA1F1	ESTs, Moderately similar to ZINC FINGER PROTEIN ZFP- 36 [Homo sapiens] {IMAGE:464215}	464215 IMAGE Consortium	AA028460
5250 F	11 021FA1F1	IMAGE EST {IMAGE:463186}	463186 IMAGE Consortium	AA030405
5251 H	1 021FA1F1	IMAGE EST {IMAGE:464152}	464152 IMAGE Consortium	AA028584
5252 H	3 021FA1F1	ESTs {IMAGE:463330}	463330 IMAGE Consortium	AA027604
5253 H	5 021FA1F1	Erythrocyte protein band 4.1 {IMAGE:463625}	463625 IMAGE Consortium	AA027522
5254 H	7 021FA1F1	IMAGE EST {IMAGE:459139}	459139 IMAGE Consortium	AA024084
5255 H	9 021FA1F1	ESTs {IMAGE:459562}	459562 IMAGE Consortium	AA027381
5256 H	11 021FA1F1	Mus musculus mRNA for vascular cadherin-2	459569 IMAGE Consortium	AA027396

		{IMAGE:459569}		
5257 B	1	021MA1F2 IMAGE EST {IMAGE:463313}	463313	IMAGE Consortium AA028673
5258 B	3	021MA1F2 IMAGE EST {IMAGE:464727}	464727	IMAGE Consortium AA030821
5259 B	5	021MA1F2 Alkaline phosphatase 2, liver {IMAGE:465052}	465052	IMAGE Consortium AA032457
5260 B	7	021MA1F2 ESTs, Weakly similar to RIBOKINASE [Bacillus subtilis] {IMAGE:465185}	465185	IMAGE Consortium AA032486
5261 B	9	021MA1F2 ESTs {IMAGE:467193}	467193	IMAGE Consortium AA034806
5262 B	11	021MA1F2 IMAGE EST {IMAGE:465744}	465744	IMAGE Consortium AA032916
5263 D	1	021MA1F2 IMAGE EST {IMAGE:466882}	466882	IMAGE Consortium AA031211
5264 D	3	021MA1F2 ESTs {IMAGE:466961}	466961	IMAGE Consortium AA034665
5265 D	5	021MA1F2 IMAGE EST {IMAGE:467550}	467550	IMAGE Consortium AA034611
5266 D	7	021MA1F2 Eyes absent 3 homolog (Drosophila) {IMAGE:466048}	466048	IMAGE Consortium AA033245
5267 D	9	021MA1F2 IMAGE EST {IMAGE:466371}	466371	IMAGE Consortium AA033504
5268 D	11	021MA1F2 IMAGE EST {IMAGE:466376}	466376	IMAGE Consortium AA032359
5269 F	1	021MA1F2 IMAGE EST {IMAGE:467535}	467535	IMAGE Consortium AA036459
5270 F	3	021MA1F2 ESTs {IMAGE:466401}	466401	IMAGE Consortium AA032380
5271 F	5	021MA1F2 Mus musculus cardiac calsequestrin mRNA, complete cds {IMAGE:466191}	466191	IMAGE Consortium AA033488
5272 F	7	021MA1F2 PEPTIDYL-PROLYL CIS-TRANS ISOMERASE B PRECURSOR {IMAGE:466632}	466632	IMAGE Consortium AA030995
5273 F	9	021MA1F2 IMAGE EST {IMAGE:466591}	466591	IMAGE Consortium AA030925
5274 F	11	021MA1F2 IMAGE EST {IMAGE:466623}	466623	IMAGE Consortium AA030949
5275 H	1	021MA1F2 IMAGE EST {IMAGE:468580}	468580	IMAGE Consortium AA014487
5276 H	3	021MA1F2 IMAGE EST {IMAGE:464996}	464996	IMAGE Consortium AA032655
5277 H	5	021MA1F2 Parathyroid hormone receptor {IMAGE:467172}	467172	IMAGE Consortium AA034782
5278 H	7	021MA1F2 IMAGE EST {IMAGE:466988}	466988	IMAGE Consortium AA034669
5279 H	9	021MA1F2 IMAGE EST {IMAGE:465680}	465680	IMAGE Consortium AA033034
5280 H	11	021MA1F2 IMAGE EST {IMAGE:465785}	465785	IMAGE Consortium AA032923
5281 B	1	021TA1F3 IMAGE EST {IMAGE:458903}	458903	IMAGE Consortium AA024250
5282 B	3	021TA1F3 IMAGE EST {IMAGE:459617}	459617	IMAGE Consortium AA027531
5283 B	5	021TA1F3 Cytochrome P450, 2e1, ethanol inducible {IMAGE:464224}	464224	IMAGE Consortium AA028481
5284 B	7	021TA1F3 IMAGE EST {IMAGE:463049}	463049	IMAGE Consortium AA030478
5285 B	9	021TA1F3 IMAGE EST {IMAGE:465170}	465170	IMAGE Consortium AA030746
5286 B	11	021TA1F3 ESTs {IMAGE:465223}	465223	IMAGE Consortium AA032563
5287 D	1	021TA1F3 IMAGE EST {IMAGE:463948}	463948	IMAGE Consortium AA028302
5288 D	3	021TA1F3 IMAGE EST {IMAGE:464121}	464121	IMAGE Consortium AA028312
5289 D	5	021TA1F3 IMAGE EST {IMAGE:464036}	464036	IMAGE Consortium AA028454
5290 D	7	021TA1F3 ESTs {IMAGE:464165}	464165	IMAGE Consortium AA028448
5291 D	9	021TA1F3 IMAGE EST {IMAGE:464245}	464245	IMAGE Consortium AA032909
5292 D	11	021TA1F3 IMAGE EST {IMAGE:463183}	463183	IMAGE Consortium AA030360
5293 F	1	021TA1F3 IMAGE EST {IMAGE:464317}	464317	IMAGE Consortium AA032853
5294 F	3	021TA1F3 ESTs, Highly similar to HYPOTHETICAL 15.5 KD PROTEIN IN BEM2-SPT2 INTERGENIC REGION [Saccharomyces	464416	IMAGE Consortium AA030665

		<i>cerevisiae</i> ] {IMAGE:464416}		
5295 F	5 021TA1F3	Peripherin {IMAGE:464493}	464493 IMAGE Consortium	AA030447
5296 F	7 021TA1F3	IMAGE EST {IMAGE:464717}	464717 IMAGE Consortium	AA030067
5297 F	9 021TA1F3	IMAGE EST {IMAGE:463006}	463006 IMAGE Consortium	AA030574
5298 F	11 021TA1F3	IMAGE EST {IMAGE:464827}	464827 IMAGE Consortium	AA032421
5299 H	1 021TA1F3	ESTs, Highly similar to HYPOTHETICAL 77.3 KD PROTEIN T05G5.8 IN CHROMOSOME III [Caenorhabditis elegans] {IMAGE:464279}	464279 IMAGE Consortium	AA033040
5300 H	3 021TA1F3	Transformation related protein 53 {IMAGE:464741}	464741 IMAGE Consortium	AA030090
5301 H	5 021TA1F3	IMAGE EST {IMAGE:464603}	464603 IMAGE Consortium	AA030153
5302 H	7 021TA1F3	IMAGE EST {IMAGE:464679}	464679 IMAGE Consortium	AA030461
5303 H	9 021TA1F3	IMAGE EST {IMAGE:464588}	464588 IMAGE Consortium	AA032966
5304 H	11 021TA1F3	Mus musculus E1B 19K/Bcl-2-binding protein homolog (Nip3) mRNA, nuclear gene encoding mitochondrial protein, complete cds {IMAGE:463388}	463388 IMAGE Consortium	AA030640
5305 B	1 0210A1F4	IMAGE EST {IMAGE:467204}	467204 IMAGE Consortium	AA034844
5306 B	3 0210A1F4	IMAGE EST {IMAGE:467377}	467377 IMAGE Consortium	AA036623
5307 B	5 0210A1F4	Cadherin 2 {IMAGE:467313}	467313 IMAGE Consortium	AA034594
5308 B	7 0210A1F4	ESTs {IMAGE:467873}	467873 IMAGE Consortium	AA036517
5309 B	9 0210A1F4	IMAGE EST {IMAGE:468171}	468171 IMAGE Consortium	AA015086
5310 B	11 0210A1F4	ESTs, Highly similar to 60S RIBOSOMAL PROTEIN L37A [Gallus gallus] {IMAGE:466384}	466384 IMAGE Consortium	AA034754
5311 D	1 0210A1F4	Mouse mRNA for the third largest RNA polymerase II subunit, complete cds {IMAGE:466648}	466648 IMAGE Consortium	AA031130
5312 D	3 0210A1F4	ESTs, Weakly similar to HYPOTHETICAL 54.9 KD PROTEIN C02F5.7 IN CHROMOSOME III [Caenorhabditis elegans] {IMAGE:466887}	466887 IMAGE Consortium	AA031219
5313 D	5 0210A1F4	IMAGE EST {IMAGE:467290}	467290 IMAGE Consortium	AA034624
5314 D	7 0210A1F4	IMAGE EST {IMAGE:467810}	467810 IMAGE Consortium	AA036499
5315 D	9 0210A1F4	Mus musculus 26S proteasome ATPase (CIP21) mRNA, complete cds {IMAGE:466254}	466254 IMAGE Consortium	AA033306
5316 D	11 0210A1F4	Poly (ADP-ribose) polymerase {IMAGE:466241}	466241 IMAGE Consortium	AA032357
5317 F	1 0210A1F4	ESTs {IMAGE:467811}	467811 IMAGE Consortium	AA036502
5318 F	3 0210A1F4	IMAGE EST {IMAGE:466093}	466093 IMAGE Consortium	AA036559
5319 F	5 0210A1F4	MyoD family inhibitor {IMAGE:466270}	466270 IMAGE Consortium	AA033404
5320 F	7 0210A1F4	IMAGE EST {IMAGE:465045}	465045 IMAGE Consortium	AA032622
5321 F	9 0210A1F4	ESTs {IMAGE:465555}	465555 IMAGE Consortium	AA032807
5322 F	11 0210A1F4	ESTs, Moderately similar to hypothetical human serine-threonine protein kinase R31240_1 [H.sapiens] {IMAGE:465490}	465490 IMAGE Consortium	AA032848
5323 H	1 0210A1F4	Myeloperoxidase {IMAGE:466693}	466693 IMAGE Consortium	AA031015
5324 H	3 0210A1F4	IMAGE EST {IMAGE:467048}	467048 IMAGE Consortium	AA034619
5325 H	5 0210A1F4	ESTs {IMAGE:465335}	465335 IMAGE Consortium	AA032437

5326 H	7 0210A1F4	Flap structure specific endonuclease 1 {IMAGE:467501}	467501 IMAGE Consortium	AA034561
5327 H	9 0210A1F4	ESTs {IMAGE:467316}	467316 IMAGE Consortium	AA034593
5328 H	11 0210A1F4	ESTs {IMAGE:467518}	467518 IMAGE Consortium	AA034718
5329 B	1 0217A1F5	IMAGE EST {IMAGE:465927}	465927 IMAGE Consortium	AA033268
5330 B	3 0217A1F5	IMAGE EST {IMAGE:466155}	466155 IMAGE Consortium	AA033351
5331 B	5 0217A1F5	IMAGE EST {IMAGE:466118}	466118 IMAGE Consortium	AA036564
5332 B	7 0217A1F5	IMAGE EST {IMAGE:466266}	466266 IMAGE Consortium	AA033402
5333 B	9 0217A1F5	ESTs {IMAGE:466637}	466637 IMAGE Consortium	AA031002
5334 B	11 0217A1F5	IMAGE EST {IMAGE:474317}	474317 IMAGE Consortium	AA038806
5335 D	1 0217A1F5	IMAGE EST {IMAGE:468900}	468900 IMAGE Consortium	AA035981
5336 D	3 0217A1F5	IMAGE EST {IMAGE:472299}	472299 IMAGE Consortium	AA036249
5337 D	5 0217A1F5	IMAGE EST {IMAGE:472801}	472801 IMAGE Consortium	AA037995
5338 D	7 0217A1F5	Voltage-dependent anion channel 3 {IMAGE:473770}	473770 IMAGE Consortium	AA038416
5339 D	9 0217A1F5	IMAGE EST {IMAGE:468498}	468498 IMAGE Consortium	AA014403
5340 D	11 0217A1F5	Polymerase (DNA directed), delta 2, regulatory subunit (50kDa) {IMAGE:468823}	468823 IMAGE Consortium	AA035902
5341 F	1 0217A1F5	IMAGE EST {IMAGE:474475}	474475 IMAGE Consortium	AA038984
5342 F	3 0217A1F5	IMAGE EST {IMAGE:472424}	472424 IMAGE Consortium	AA036387
5343 F	5 0217A1F5	ESTs {IMAGE:473419}	473419 IMAGE Consortium	AA038587
5344 F	7 0217A1F5	IMAGE EST {IMAGE:468500}	468500 IMAGE Consortium	AA014404
5345 F	9 0217A1F5	EST {IMAGE:474319}	474319 IMAGE Consortium	AA038807
5346 F	11 0217A1F5	IMAGE EST {IMAGE:469049}	469049 IMAGE Consortium	AA036060
5347 H	1 0217A1F5	IMAGE EST {IMAGE:473905}	473905 IMAGE Consortium	AA038470
5348 H	3 0217A1F5	IMAGE EST {IMAGE:468867}	468867 IMAGE Consortium	AA035956
5349 H	5 0217A1F5	ESTs {IMAGE:468838}	468838 IMAGE Consortium	AA035939
5350 H	7 0217A1F5	IMAGE EST {IMAGE:472831}	472831 IMAGE Consortium	AA038003
5351 H	9 0217A1F5	IMAGE EST {IMAGE:474025}	474025 IMAGE Consortium	AA038450
5352 H	11 0217A1F5	IMAGE EST {IMAGE:475419}	475419 IMAGE Consortium	AA041716
5353 B	1 021EA1F6	IMAGE EST {IMAGE:476865}	476865 IMAGE Consortium	AA050826
5354 B	3 021EA1F6	ESTs {IMAGE:477158}	477158 IMAGE Consortium	AA047929
5355 B	5 021EA1F6	ESTs {IMAGE:475903}	475903 IMAGE Consortium	AA044557
5356 B	7 021EA1F6	Mus musculus mRNA for GFRalpha-3, complete cds {IMAGE:475791}	475791 IMAGE Consortium	AA049894
5357 B	9 021EA1F6	Mus musculus branched-chain alpha-ketoacid dehydrogenase E1 beta-subunit mRNA sequence {IMAGE:476163}	476163 IMAGE Consortium	AA050586
5358 B	11 021EA1F6	ESTs {IMAGE:476369}	476369 IMAGE Consortium	AA050212
5359 D	1 021EA1F6	ESTs {IMAGE:476218}	476218 IMAGE Consortium	AA050204
5360 D	3 021EA1F6	IMAGE EST {IMAGE:477829}	477829 IMAGE Consortium	AA048626
5361 D	5 021EA1F6	IMAGE EST {IMAGE:478040}	478040 IMAGE Consortium	AA049480
5362 D	7 021EA1F6	IMAGE EST {IMAGE:478309}	478309 IMAGE Consortium	AA049436
5363 D	9 021EA1F6	IMAGE EST {IMAGE:475944}	475944 IMAGE Consortium	AA049712
5364 D	11 021EA1F6	IMAGE EST {IMAGE:476045}	476045 IMAGE Consortium	AA050277

5365 F	1 021EA1F6	IMAGE EST {IMAGE:477964}	477964 IMAGE Consortium	AA049417
5366 F	3 021EA1F6	Telomeric repeat binding factor 2 {IMAGE:478166}	478166 IMAGE Consortium	AA049607
5367 F	5 021EA1F6	IMAGE EST {IMAGE:475355}	475355 IMAGE Consortium	AA042063
5368 F	7 021EA1F6	Mus musculus kruppel-related zinc finger protein (Emzf1) mRNA, complete cds {IMAGE:477064}	477064 IMAGE Consortium	AA047965
5369 F	9 021EA1F6	ESTs {IMAGE:477140}	477140 IMAGE Consortium	AA048171
5370 F	11 021EA1F6	EST {IMAGE:475817}	475817 IMAGE Consortium	AA050145
5371 H	1 021EA1F6	IMAGE EST {IMAGE:477045}	477045 IMAGE Consortium	AA047964
5372 H	3 021EA1F6	Mus musculus mRNA for CDC6 protein {IMAGE:477516}	477516 IMAGE Consortium	AA048426
5373 H	5 021EA1F6	IMAGE EST {IMAGE:476296}	476296 IMAGE Consortium	AA050578
5374 H	7 021EA1F6	MyoD family inhibitor {IMAGE:477664}	477664 IMAGE Consortium	AA048578
5375 H	9 021EA1F6	ESTs, Weakly similar to K04G7.11 [C.elegans] {IMAGE:478012}	478012 IMAGE Consortium	AA049463
5376 H	11 021EA1F6	IMAGE EST {IMAGE:478310}	478310 IMAGE Consortium	AA049693
5377 B	1 021LA1F7	IMAGE EST {IMAGE:468911}	468911 IMAGE Consortium	AA036042
5378 B	3 021LA1F7	IMAGE EST {IMAGE:472808}	472808 IMAGE Consortium	AA038063
5379 B	5 021LA1F7	Snail homolog, (Drosophila) {IMAGE:469080}	469080 IMAGE Consortium	AA036380
5380 B	7 021LA1F7	IMAGE EST {IMAGE:474179}	474179 IMAGE Consortium	AA038685
5381 B	9 021LA1F7	IMAGE EST {IMAGE:472595}	472595 IMAGE Consortium	AA037955
5382 B	11 021LA1F7	IMAGE EST {IMAGE:475491}	475491 IMAGE Consortium	AA042014
5383 D	1 021LA1F7	ESTs, Highly similar to KERATIN, TYPE II CYTOSKELETAL 4 [Homo sapiens] {IMAGE:474388}	474388 IMAGE Consortium	AA038821
5384 D	3 021LA1F7	ESTs {IMAGE:468952}	468952 IMAGE Consortium	AA036085
5385 D	5 021LA1F7	IMAGE EST {IMAGE:472408}	472408 IMAGE Consortium	AA036134
5386 D	7 021LA1F7	IMAGE EST {IMAGE:473389}	473389 IMAGE Consortium	AA038548
5387 D	9 021LA1F7	ESTs {IMAGE:475296}	475296 IMAGE Consortium	AA041963
5388 D	11 021LA1F7	ESTs {IMAGE:475349}	475349 IMAGE Consortium	AA041969
5389 F	1 021LA1F7	Mus musculus betaglycan mRNA, complete cds {IMAGE:476264}	476264 IMAGE Consortium	AA050765
5390 F	3 021LA1F7	ESTs, Weakly similar to Marenostriin protein [H.sapiens] {IMAGE:476205}	476205 IMAGE Consortium	AA050695
5391 F	5 021LA1F7	IMAGE EST {IMAGE:476618}	476618 IMAGE Consortium	AA050479
5392 F	7 021LA1F7	ESTs {IMAGE:478157}	478157 IMAGE Consortium	AA049610
5393 F	9 021LA1F7	IMAGE EST {IMAGE:478044}	478044 IMAGE Consortium	AA049482
5394 F	11 021LA1F7	IMAGE EST {IMAGE:478336}	478336 IMAGE Consortium	AA049760
5395 H	1 021LA1F7	IMAGE EST {IMAGE:476609}	476609 IMAGE Consortium	AA050792
5396 H	3 021LA1F7	ESTs {IMAGE:478025}	478025 IMAGE Consortium	AA049360
5397 H	5 021LA1F7	Integrin alpha 5 (fibronectin receptor alpha) {IMAGE:476908}	476908 IMAGE Consortium	AA047917
5398 H	7 021LA1F7	IMAGE EST {IMAGE:475483}	475483 IMAGE Consortium	AA044468
5399 H	9 021LA1F7	Human putative holocytochrome c-type synthetase mRNA, complete cds {IMAGE:475656}	475656 IMAGE Consortium	AA049912
5400 H	11 021LA1F7	ESTs, Moderately similar to GAR1 PROTEIN [Saccharomyces cerevisiae] {IMAGE:475814}	475814 IMAGE Consortium	AA049897



5401 B	1	021SA1F8	ESTs {IMAGE:478008}	478008	IMAGE Consortium	AA049461
5402 B	3	021SA1F8	ESTs {IMAGE:478156}	478156	IMAGE Consortium	AA049602
5403 B	5	021SA1F8	Lamin B2 {IMAGE:478334}	478334	IMAGE Consortium	AA049647
5404 B	7	021SA1F8	Nuclear, factor, erythroid derived 2, like 2 {IMAGE:475505}	475505	IMAGE Consortium	AA044475
5405 B	9	021SA1F8	IMAGE EST {IMAGE:476019}	476019	IMAGE Consortium	AA049936
5406 B	11	021SA1F8	ESTs {IMAGE:475835}	475835	IMAGE Consortium	AA049709
5407 D	1	021SA1F8	Mus musculus skeletal muscle LIM protein (FLH1) mRNA, complete cds {IMAGE:477066}	477066	IMAGE Consortium	AA047966
5408 D	3	021SA1F8	IMAGE EST {IMAGE:475932}	475932	IMAGE Consortium	AA050339
5409 D	5	021SA1F8	IMAGE EST {IMAGE:477517}	477517	IMAGE Consortium	AA048433
5410 D	7	021SA1F8	IMAGE EST {IMAGE:476341}	476341	IMAGE Consortium	AA050453
5411 D	9	021SA1F8	ESTs, Weakly similar to F01F1.11 [C.elegans] {IMAGE:476061}	476061	IMAGE Consortium	AA049839
5412 D	11	021SA1F8	ESTs {IMAGE:478133}	478133	IMAGE Consortium	AA049537
5413 F	1	021SA1F8	ESTs {IMAGE:478878}	478878	IMAGE Consortium	AA051259
5414 F	3	021SA1F8	IMAGE EST {IMAGE:478788}	478788	IMAGE Consortium	AA051003
5415 F	5	021SA1F8	ESTs {IMAGE:480140}	480140	IMAGE Consortium	AA058013
5416 F	7	021SA1F8	EST {IMAGE:480600}	480600	IMAGE Consortium	AA064016
5417 F	9	021SA1F8	IMAGE EST {IMAGE:479480}	479480	IMAGE Consortium	AA049260
5418 F	11	021SA1F8	IMAGE EST {IMAGE:480569}	480569	IMAGE Consortium	AA059494
5419 H	1	021SA1F8	ESTs {IMAGE:478770}	478770	IMAGE Consortium	AA051207
5420 H	3	021SA1F8	ESTs {IMAGE:479305}	479305	IMAGE Consortium	AA048766
5421 H	5	021SA1F8	ESTs, Weakly similar to D52 [M.musculus] {IMAGE:479358}	479358	IMAGE Consortium	AA048792
5422 H	7	021SA1F8	Aquaporin 1 {IMAGE:480467}	480467	IMAGE Consortium	AA064011
5423 H	9	021SA1F8	ESTs, Highly similar to mevalonate pyrophosphate decarboxylase [R.norvegicus] {IMAGE:479201}	479201	IMAGE Consortium	AA049342
5424 H	11	021SA1F8	Protein kinase, cAMP dependent, catalytic, alpha {IMAGE:479691}	479691	IMAGE Consortium	AA051526
5425 B	1	021ZA1F9	ESTs, Highly similar to mevalonate pyrophosphate decarboxylase [R.norvegicus] {IMAGE:480621}	480621	IMAGE Consortium	AA059528
5426 B	3	021ZA1F9	ESTs {IMAGE:478477}	478477	IMAGE Consortium	AA050004
5427 B	5	021ZA1F9	IMAGE EST {IMAGE:478409}	478409	IMAGE Consortium	AA049955
5428 B	7	021ZA1F9	ESTs, Weakly similar to UGTrel1 [M.musculus] {IMAGE:478532}	478532	IMAGE Consortium	AA049844
5429 B	9	021ZA1F9	IMAGE EST {IMAGE:480149}	480149	IMAGE Consortium	AA051732
5430 B	11	021ZA1F9	ESTs {IMAGE:478975}	478975	IMAGE Consortium	AA048895
5431 D	1	021ZA1F9	IMAGE EST {IMAGE:478556}	478556	IMAGE Consortium	AA051330
5432 D	3	021ZA1F9	Cytochrome P450, 2e1, ethanol inducible {IMAGE:478752}	478752	IMAGE Consortium	AA051170
5433 D	5	021ZA1F9	ESTs {IMAGE:479076}	479076	IMAGE Consortium	AA048874
5434 D	7	021ZA1F9	IMAGE EST {IMAGE:479143}	479143	IMAGE Consortium	AA049116
5435 D	9	021ZA1F9	ESTs {IMAGE:480618}	480618	IMAGE Consortium	AA059523
5436 D	11	021ZA1F9	IMAGE EST {IMAGE:479375}	479375	IMAGE Consortium	AA049044
5437 F	1	021ZA1F9	IMAGE EST {IMAGE:480454}	480454	IMAGE Consortium	AA058140

5438 F	3 021ZA1F9	IMAGE EST {IMAGE:479471}	479471 IMAGE Consortium	AA049350
5439 F	5 021ZA1F9	ESTs, Highly similar to ESS1 PROTEIN [ <i>Saccharomyces cerevisiae</i> ] {IMAGE:478645}	478645 IMAGE Consortium	AA051390
5440 F	7 021ZA1F9	IMAGE EST {IMAGE:479709}	479709 IMAGE Consortium	AA051664
5441 F	9 021ZA1F9	IMAGE EST {IMAGE:479016}	479016 IMAGE Consortium	AA049268
5442 F	11 021ZA1F9	ESTs, Weakly similar to 36k hydrophobic nucleolar protein [ <i>M.musculus</i> ] {IMAGE:479053}	479053 IMAGE Consortium	AA048845
5443 H	1 021ZA1F9	IMAGE EST {IMAGE:480207}	480207 IMAGE Consortium	AA058195
5444 H	3 021ZA1F9	ESTs, Moderately similar to JAK3 [ <i>M.musculus</i> ] {IMAGE:480043}	480043 IMAGE Consortium	AA051649
5445 H	5 021ZA1F9	IMAGE EST {IMAGE:480534}	480534 IMAGE Consortium	AA058292
5446 H	7 021ZA1F9	Mus musculus sulfotransferase-related protein (SULT-X1) mRNA, complete cds {IMAGE:479196}	479196 IMAGE Consortium	AA049213
5447 H	9 021ZA1F9	Twist gene homolog, ( <i>Drosophila</i> ) {IMAGE:479367}	479367 IMAGE Consortium	AA048671
5448 H	11 021ZA1F9	ESTs, Highly similar to N-ACETYLGLUCOSAMINE-6-SULFATASE PRECURSOR [ <i>Homo sapiens</i> ] {IMAGE:480526}	480526 IMAGE Consortium	AA063941
5449 B	1 021JA1FA	IMAGE EST {IMAGE:481204}	481204 IMAGE Consortium	AA064326
5450 B	3 021JA1FA	Glutamate oxaloacetate transaminase 1, soluble {IMAGE:481381}	481381 IMAGE Consortium	AA060494
5451 B	5 021JA1FA	Mus musculus schwannoma-associated protein (SAM9) mRNA, complete cds {IMAGE:481051}	481051 IMAGE Consortium	AA060268
5452 B	7 021JA1FA	EST {IMAGE:481634}	481634 IMAGE Consortium	AA060150
5453 B	9 021JA1FA	ESTs {IMAGE:481784}	481784 IMAGE Consortium	AA060076
5454 B	11 021JA1FA	U1 small nuclear ribonucleoprotein 70 kDa polypeptide A {IMAGE:481692}	481692 IMAGE Consortium	AA060238
5455 D	1 021JA1FA	ESTs, Highly similar to 140 KD NUCLEOLAR PHOSPHOPROTEIN [ <i>Rattus norvegicus</i> ] {IMAGE:481372}	481372 IMAGE Consortium	AA060483
5456 D	3 021JA1FA	IMAGE EST {IMAGE:481400}	481400 IMAGE Consortium	AA060500
5457 D	5 021JA1FA	IMAGE EST {IMAGE:481405}	481405 IMAGE Consortium	AA060509
5458 D	7 021JA1FA	IMAGE EST {IMAGE:483033}	483033 IMAGE Consortium	AA060979
5459 D	9 021JA1FA	Acetyl coenzyme A dehydrogenase, medium chain {IMAGE:483333}	483333 IMAGE Consortium	AA061679
5460 D	11 021JA1FA	ESTs, Moderately similar to inositol 1,4,5-trisphosphate receptor type 2 [ <i>M.musculus</i> ] {IMAGE:481721}	481721 IMAGE Consortium	AA060171
5461 F	1 021JA1FA	ESTs {IMAGE:482652}	482652 IMAGE Consortium	AA061980
5462 F	3 021JA1FA	ESTs {IMAGE:482752}	482752 IMAGE Consortium	AA062075
5463 F	5 021JA1FA	ESTs {IMAGE:481316}	481316 IMAGE Consortium	AA060475
5464 F	7 021JA1FA	Max protein {IMAGE:481641}	481641 IMAGE Consortium	AA060226
5465 F	9 021JA1FA	IMAGE EST {IMAGE:481453}	481453 IMAGE Consortium	AA060396
5466 F	11 021JA1FA	ESTs, Moderately similar to ADP-RIBOSYLATION FACTOR 1 [ <i>Homo sapiens</i> ; <i>Bos taurus</i> ; <i>Rattus norvegicus</i> ] {IMAGE:481530}	481530 IMAGE Consortium	AA060086
5467 H	1 021JA1FA	ESTs {IMAGE:481373}	481373 IMAGE Consortium	AA060490

5468 H	3 021JA1FA	Vacuolar protein sorting-associated protein 45 {IMAGE:481192}	481192 IMAGE Consortium	AA060373
5469 H	5 021JA1FA	ESTs, Moderately similar to COLIPASE PRECURSOR [Rattus norvegicus] {IMAGE:481341}	481341 IMAGE Consortium	AA060526
5470 H	7 021JA1FA	IMAGE EST {IMAGE:482971}	482971 IMAGE Consortium	AA061042
5471 H	9 021JA1FA	ESTs {IMAGE:481507}	481507 IMAGE Consortium	AA060314
5472 H	11 021JA1FA	Mus musculus transcription factor PBX1b (PBX1b) mRNA, complete cds {IMAGE:483688}	483688 IMAGE Consortium	AA061468
5473 B	1 021QA1FB	IMAGE EST {IMAGE:480630}	480630 IMAGE Consortium	AA064149
5474 B	3 021QA1FB	ESTs, Weakly similar to unknown [M.musculus] {IMAGE:480562}	480562 IMAGE Consortium	AA060656
5475 B	5 021QA1FB	Laminin, alpha 5 {IMAGE:479439}	479439 IMAGE Consortium	AA049251
5476 B	7 021QA1FB	ESTs {IMAGE:479179}	479179 IMAGE Consortium	AA049336
5477 B	9 021QA1FB	IMAGE EST {IMAGE:478417}	478417 IMAGE Consortium	AA049820
5478 B	11 021QA1FB	ESTs, Highly similar to GLYCINE AMIDINOTRANSFERASE PRECURSOR [R.norvegicus] {IMAGE:478521}	478521 IMAGE Consortium	AA049981
5479 D	1 021QA1FB	IMAGE EST {IMAGE:480275}	480275 IMAGE Consortium	AA058029
5480 D	3 021QA1FB	ESTs {IMAGE:480568}	480568 IMAGE Consortium	AA059492
5481 D	5 021QA1FB	IMAGE EST {IMAGE:480452}	480452 IMAGE Consortium	AA058139
5482 D	7 021QA1FB	Mus musculus MO15-associated kinase (MO15) mRNA, complete cds {IMAGE:479857}	479857 IMAGE Consortium	AA051744
5483 D	9 021QA1FB	IMAGE EST {IMAGE:479819}	479819 IMAGE Consortium	AA051474
5484 D	11 021QA1FB	ESTs, Moderately similar to The ha3523 gene product is related to S.cerevisiae gene product located in chromosome III. [H.sapiens] {IMAGE:479042}	479042 IMAGE Consortium	AA049276
5485 F	1 021QA1FB	ESTs, Weakly similar to ZK1307.5 [C.elegans] {IMAGE:478665}	478665 IMAGE Consortium	AA051396
5486 F	3 021QA1FB	IMAGE EST {IMAGE:479748}	479748 IMAGE Consortium	AA051421
5487 F	5 021QA1FB	ESTs {IMAGE:479728}	479728 IMAGE Consortium	AA051666
5488 F	7 021QA1FB	ESTs {IMAGE:478701}	478701 IMAGE Consortium	AA051416
5489 F	9 021QA1FB	ESTs {IMAGE:478854}	478854 IMAGE Consortium	AA051252
5490 F	11 021QA1FB	ESTs {IMAGE:478740}	478740 IMAGE Consortium	AA051192
5491 H	1 021QA1FB	IMAGE EST {IMAGE:481108}	481108 IMAGE Consortium	AA064293
5492 H	3 021QA1FB	Inter-alpha trypsin inhibitor, heavy chain 3 {IMAGE:482895}	482895 IMAGE Consortium	AA062129
5493 H	5 021QA1FB	M.musculus mRNA for vacuolar adenosine triphosphatase subunit B {IMAGE:481410}	481410 IMAGE Consortium	AA060559
5494 H	7 021QA1FB	ESTs {IMAGE:483285}	483285 IMAGE Consortium	AA061485
5495 H	9 021QA1FB	IMAGE EST {IMAGE:483650}	483650 IMAGE Consortium	AA061359
5496 H	11 021QA1FB	ESTs {IMAGE:483633}	483633 IMAGE Consortium	AA061338
5497 B	1 021XA1FC	IMAGE EST {IMAGE:482986}	482986 IMAGE Consortium	AA060961
5498 B	3 021XA1FC	ESTs {IMAGE:481618}	481618 IMAGE Consortium	AA060217
5499 B	5 021XA1FC	IMAGE EST {IMAGE:483588}	483588 IMAGE Consortium	AA061886
5500 B	7 021XA1FC	CLATHRIN COAT ASSEMBLY PROTEIN AP47 {IMAGE:483377}	483377 IMAGE Consortium	AA061868

5501 B	9 021XA1FC	IMAGE EST {IMAGE:483647}	483647 IMAGE Consortium	AA061460
5502 B	11 021XA1FC	IMAGE EST {IMAGE:483756}	483756 IMAGE Consortium	AA061403
5503 D	1 021XA1FC	ESTs {IMAGE:483528}	483528 IMAGE Consortium	AA061801
5504 D	3 021XA1FC	ESTs {IMAGE:483636}	483636 IMAGE Consortium	AA061332
5505 D	5 021XA1FC	IMAGE EST {IMAGE:483689}	483689 IMAGE Consortium	AA061213
5506 D	7 021XA1FC	IMAGE EST {IMAGE:482052}	482052 IMAGE Consortium	AA059897
5507 D	9 021XA1FC	IMAGE EST {IMAGE:482062}	482062 IMAGE Consortium	AA059981
5508 D	11 021XA1FC	ESTs {IMAGE:482272}	482272 IMAGE Consortium	AA060793
5509 F	1 021XA1FC	Crystallin, gamma S {IMAGE:483498}	483498 IMAGE Consortium	AA060041
5510 F	3 021XA1FC	Laminin, gamma 1 {IMAGE:481837}	481837 IMAGE Consortium	AA059779
5511 F	5 021XA1FC	ESTs {IMAGE:482304}	482304 IMAGE Consortium	AA060891
5512 F	7 021XA1FC	IMAGE EST {IMAGE:480989}	480989 IMAGE Consortium	AA059561
5513 F	9 021XA1FC	IMAGE EST {IMAGE:480949}	480949 IMAGE Consortium	AA060003
5514 F	11 021XA1FC	ESTs {IMAGE:481202}	481202 IMAGE Consortium	AA064325
5515 H	1 021XA1FC	Retinol binding protein 2, cellular {IMAGE:483493}	483493 IMAGE Consortium	AA061550
5516 H	3 021XA1FC	EST {IMAGE:483643}	483643 IMAGE Consortium	AA061458
5517 H	5 021XA1FC	IMAGE EST {IMAGE:483724}	483724 IMAGE Consortium	AA061352
5518 H	7 021XA1FC	Thrombospondin 3 {IMAGE:483763}	483763 IMAGE Consortium	AA061474
5519 H	9 021XA1FC	IMAGE EST {IMAGE:484134}	484134 IMAGE Consortium	AA079969
5520 H	11 021XA1FC	Ubiquitin-conjugating enzyme 7 {IMAGE:482359}	482359 IMAGE Consortium	AA060906
5521 B	1 0214A1FD	IMAGE EST {IMAGE:483584}	483584 IMAGE Consortium	AA061884
5522 B	3 0214A1FD	ESTs, Highly similar to HYPOTHETICAL 70.2 KD PROTEIN IN GSH1-CHS6 INTERGENIC REGION [ <i>Saccharomyces cerevisiae</i> ] {IMAGE:483541}	483541 IMAGE Consortium	AA061763
5523 B	5 0214A1FD	IMAGE EST {IMAGE:482001}	482001 IMAGE Consortium	AA059886
5524 B	7 0214A1FD	ESTs, Highly similar to ELECTRON TRANSFER FLAVOPROTEIN ALPHA-SUBUNIT PRECURSOR [ <i>Homo sapiens</i> ] {IMAGE:481934}	481934 IMAGE Consortium	AA060723
5525 B	9 0214A1FD	ESTs, Highly similar to SH3 domain-containing protein SH3P17 [ <i>H.sapiens</i> ] {IMAGE:482370}	482370 IMAGE Consortium	AA063751
5526 B	11 0214A1FD	ESTs {IMAGE:482578}	482578 IMAGE Consortium	AA061899
5527 D	1 0214A1FD	Mouse mRNA for neuropsin, complete cds {IMAGE:483813}	483813 IMAGE Consortium	AA061507
5528 D	3 0214A1FD	IMAGE EST {IMAGE:482432}	482432 IMAGE Consortium	AA060844
5529 D	5 0214A1FD	ESTs {IMAGE:480854}	480854 IMAGE Consortium	AA064231
5530 D	7 0214A1FD	EST {IMAGE:481084}	481084 IMAGE Consortium	AA064265
5531 D	9 0214A1FD	ESTs {IMAGE:481375}	481375 IMAGE Consortium	AA060491
5532 D	11 0214A1FD	Mus musculus Swiss Webster demilune cell-specific protein mRNA, complete cds {IMAGE:482661}	482661 IMAGE Consortium	AA062054
5533 F	1 0214A1FD	ESTs {IMAGE:484327}	484327 IMAGE Consortium	AA080287
5534 F	3 0214A1FD	Male enhanced antigen 1 {IMAGE:484122}	484122 IMAGE Consortium	AA073695
5535 F	5 0214A1FD	Mus musculus cis-retinol androgen dehydrogenase 1 mRNA, complete cds {IMAGE:571559}	571559 IMAGE Consortium	AA108260
5536 F	7 0214A1FD	Bone morphogenetic protein 1 {IMAGE:492502}	492502 IMAGE Consortium	AA097223

5537 F	9 0214A1FD	Mus musculus G protein-coupled receptor mRNA, complete cds {IMAGE:493202}	493202 IMAGE Consortium	AA097490
5538 F	11 0214A1FD	IMAGE EST {IMAGE:572772}	572772 IMAGE Consortium	AA111754
5539 H	1 0214A1FD	Mus musculus nucleolar protein (MSP58) mRNA, complete cds {IMAGE:493227}	493227 IMAGE Consortium	AA098041
5540 H	3 0214A1FD	ESTs {IMAGE:572303}	572303 IMAGE Consortium	AA105161
5541 H	5 0214A1FD	IMAGE EST {IMAGE:493480}	493480 IMAGE Consortium	AA086931
5542 H	7 0214A1FD	ESTs, Highly similar to UBIQUINOL-CYTOCHROME C REDUCTASE COMPLEX 7.2 KD PROTEIN [Bos taurus] {IMAGE:493604}	493604 IMAGE Consortium	AA087137
5543 H	9 0214A1FD	Mus musculus mu-crystallin (Crym) mRNA, complete cds {IMAGE:570673}	570673 IMAGE Consortium	AA108639
5544 H	11 0214A1FD	IMAGE EST {IMAGE:573414}	573414 IMAGE Consortium	AA118751
5545 B	1 021BA1FE	ESTs {IMAGE:484125}	484125 IMAGE Consortium	AA073509
5546 B	3 021BA1FE	Mus musculus interleukin enhancer binding factor 3 (Ilf3) mRNA, complete cds {IMAGE:571805}	571805 IMAGE Consortium	AA108959
5547 B	5 021BA1FE	Apolipoprotein A-IV {IMAGE:492937}	492937 IMAGE Consortium	AA097421
5548 B	7 021BA1FE	Placental growth factor {IMAGE:493282}	493282 IMAGE Consortium	AA098166
5549 B	9 021BA1FE	ESTs {IMAGE:493295}	493295 IMAGE Consortium	AA098551
5550 B	11 021BA1FE	IMAGE EST {IMAGE:572840}	572840 IMAGE Consortium	AA110794
5551 D	1 021BA1FE	Mus musculus fatty acid transport protein 2 mRNA, complete cds {IMAGE:572379}	572379 IMAGE Consortium	AA108401
5552 D	3 021BA1FE	IMAGE EST {IMAGE:492953}	492953 IMAGE Consortium	AA097825
5553 D	5 021BA1FE	IMAGE EST {IMAGE:572513}	572513 IMAGE Consortium	AA109944
5554 D	7 021BA1FE	IMAGE EST {IMAGE:572904}	572904 IMAGE Consortium	AA116900
5555 D	9 021BA1FE	Ferritin heavy chain {IMAGE:572888}	572888 IMAGE Consortium	AA110807
5556 D	11 021BA1FE	ESTs {IMAGE:493702}	493702 IMAGE Consortium	AA098361
5557 F	1 021BA1FE	IMAGE EST {IMAGE:493555}	493555 IMAGE Consortium	AA087125
5558 F	3 021BA1FE	ESTs {IMAGE:493415}	493415 IMAGE Consortium	AA087054
5559 F	5 021BA1FE	IMAGE EST {IMAGE:570613}	570613 IMAGE Consortium	AA108603
5560 F	7 021BA1FE	IMAGE EST {IMAGE:571091}	571091 IMAGE Consortium	AA108648
5561 F	9 021BA1FE	ESTs {IMAGE:573520}	573520 IMAGE Consortium	AA110651
5562 F	11 021BA1FE	IMAGE EST {IMAGE:571451}	571451 IMAGE Consortium	AA109106
5563 H	1 021BA1FE	ESTs {IMAGE:573365}	573365 IMAGE Consortium	AA118680
5564 H	3 021BA1FE	ESTs {IMAGE:571369}	571369 IMAGE Consortium	AA108948
5565 H	5 021BA1FE	Mus musculus vacuolar adenosine triphosphatase subunit C mRNA, complete cds {IMAGE:484433}	484433 IMAGE Consortium	AA073887
5566 H	7 021BA1FE	IMAGE EST {IMAGE:571463}	571463 IMAGE Consortium	AA105370
5567 H	9 021BA1FE	Mus musculus fatty acid transport protein 4 mRNA, partial cds {IMAGE:493182}	493182 IMAGE Consortium	AA097486
5568 H	11 021BA1FE	IMAGE EST {IMAGE:493212}	493212 IMAGE Consortium	AA097795
5569 B	1 021IA1FF	IMAGE EST {IMAGE:572613}	572613 IMAGE Consortium	AA109917
5570 B	3 021IA1FF	ESTs, Highly similar to HYPOTHETICAL PROTEIN	493385 IMAGE Consortium	AA087036

		KIAA0069 [H.sapiens] {IMAGE:493385}		
5571 B	5 021IA1FF	ESTs {IMAGE:493652}	493652 IMAGE Consortium	AA098345
5572 B	7 021IA1FF	IMAGE EST {IMAGE:570788}	570788 IMAGE Consortium	AA108713
5573 B	9 021IA1FF	IMAGE EST {IMAGE:570978}	570978 IMAGE Consortium	AA109751
5574 B	11 021IA1FF	Mannosidase 2, alpha B1 {IMAGE:484259}	484259 IMAGE Consortium	AA080231
5575 D	1 021IA1FF	ESTs, Highly similar to HYPOTHETICAL MYELOID CELL LINE PROTEIN 6 [Homo sapiens] {IMAGE:573076}	573076 IMAGE Consortium	AA118706
5576 D	3 021IA1FF	IMAGE EST {IMAGE:572867}	572867 IMAGE Consortium	AA110804
5577 D	5 021IA1FF	Glycerolphosphate dehydrogenase 1, cytoplasmic adult {IMAGE:570675}	570675 IMAGE Consortium	AA108640
5578 D	7 021IA1FF	ESTs, Highly similar to ACETYL-COENZYME A SYNTHETASE [Bacillus subtilis] {IMAGE:570550}	570550 IMAGE Consortium	AA109675
5579 D	9 021IA1FF	IMAGE EST {IMAGE:484284}	484284 IMAGE Consortium	AA080236
5580 D	11 021IA1FF	ESTs, Moderately similar to eukaryotic translation initiation factor 3 subunit [H.sapiens] {IMAGE:571406}	571406 IMAGE Consortium	AA109090
5581 F	1 021IA1FF	ESTs {IMAGE:573410}	573410 IMAGE Consortium	AA118749
5582 F	3 021IA1FF	IMAGE EST {IMAGE:571124}	571124 IMAGE Consortium	AA109765
5583 F	5 021IA1FF	Keratin complex 1, acidic, gene 13 {IMAGE:484261}	484261 IMAGE Consortium	AA080232
5584 F	7 021IA1FF	IMAGE EST {IMAGE:484267}	484267 IMAGE Consortium	AA073827
5585 F	9 021IA1FF	ESTs, Moderately similar to LAMBDA-CRYSTALLIN [Oryctolagus cuniculus] {IMAGE:571803}	571803 IMAGE Consortium	AA108958
5586 F	11 021IA1FF	Mouse chromosome 6 BAC-284H12 (Research Genetics mouse BAC library) complete sequence {IMAGE:571954}	571954 IMAGE Consortium	AA108340
5587 H	1 021IA1FF	ESTs {IMAGE:573482}	573482 IMAGE Consortium	AA118868
5588 H	3 021IA1FF	Mus musculus mRNA for translation initiation factor eIF2 gamma Y {IMAGE:573320}	573320 IMAGE Consortium	AA118716
5589 H	5 021IA1FF	IMAGE EST {IMAGE:573398}	573398 IMAGE Consortium	AA118816
5590 H	7 021IA1FF	IMAGE EST {IMAGE:571447}	571447 IMAGE Consortium	AA109027
5591 H	9 021IA1FF	Mus musculus COP9 complex subunit 6 (COPS6) mRNA, complete cds {IMAGE:571385}	571385 IMAGE Consortium	AA105330
5592 H	11 021IA1FF	IMAGE EST {IMAGE:493104}	493104 IMAGE Consortium	AA097894
5593 B	1 021PA1FG	IMAGE EST {IMAGE:573771}	573771 IMAGE Consortium	AA119029
5594 B	3 021PA1FG	ESTs {IMAGE:575647}	575647 IMAGE Consortium	AA125243
5595 B	5 021PA1FG	IMAGE EST {IMAGE:574119}	574119 IMAGE Consortium	AA119086
5596 B	7 021PA1FG	IMAGE EST {IMAGE:575922}	575922 IMAGE Consortium	AA117606
5597 B	9 021PA1FG	ESTs {IMAGE:576315}	576315 IMAGE Consortium	AA120084
5598 B	11 021PA1FG	ESTs {IMAGE:574435}	574435 IMAGE Consortium	AA118290
5599 D	1 021PA1FG	IMAGE EST {IMAGE:575696}	575696 IMAGE Consortium	AA123006
5600 D	3 021PA1FG	ESTs {IMAGE:576172}	576172 IMAGE Consortium	AA119873
5601 D	5 021PA1FG	ESTs {IMAGE:576131}	576131 IMAGE Consortium	AA119797
5602 D	7 021PA1FG	IMAGE EST {IMAGE:574375}	574375 IMAGE Consortium	AA119571
5603 D	9 021PA1FG	ESTs {IMAGE:576344}	576344 IMAGE Consortium	AA120260
5604 D	11 021PA1FG	ESTs {IMAGE:576401}	576401 IMAGE Consortium	AA119676

5605 F	1 021PA1FG	ESTs, Highly similar to The KIAA0136 gene product is novel. [H.sapiens] {IMAGE:575554}	575554 IMAGE Consortium	AA125328
5606 F	3 021PA1FG	IMAGE EST {IMAGE:574287}	574287 IMAGE Consortium	AA117612
5607 F	5 021PA1FG	ESTs {IMAGE:574418}	574418 IMAGE Consortium	AA119581
5608 F	7 021PA1FG	ESTs {IMAGE:574447}	574447 IMAGE Consortium	AA117560
5609 F	9 021PA1FG	IMAGE EST {IMAGE:574676}	574676 IMAGE Consortium	AA120623
5610 F	11 021PA1FG	IMAGE EST {IMAGE:574995}	574995 IMAGE Consortium	AA124997
5611 H	1 021PA1FG	ESTs, Moderately similar to zinc finger protein [M.musculus] {IMAGE:574258}	574258 IMAGE Consortium	AA117644
5612 H	3 021PA1FG	IMAGE EST {IMAGE:576117}	576117 IMAGE Consortium	AA124563
5613 H	5 021PA1FG	Hepatic lipase {IMAGE:576579}	576579 IMAGE Consortium	AA124071
5614 H	7 021PA1FG	IMAGE EST {IMAGE:576440}	576440 IMAGE Consortium	AA119728
5615 H	9 021PA1FG	ESTs, Moderately similar to protein IEF SSP 9502 [H.sapiens] {IMAGE:576570}	576570 IMAGE Consortium	AA119826
5616 H	11 021PA1FG	ESTs, Highly similar to The ha3611 gene product is related to S.cerevisiae SNM1 protein. [H.sapiens] {IMAGE:576960}	576960 IMAGE Consortium	AA122530
5617 B	1 021WA1FH	IMAGE EST {IMAGE:574658}	574658 IMAGE Consortium	AA125255
5618 B	3 021WA1FH	IMAGE EST {IMAGE:574519}	574519 IMAGE Consortium	AA117642
5619 B	5 021WA1FH	IMAGE EST {IMAGE:574693}	574693 IMAGE Consortium	AA125230
5620 B	7 021WA1FH	ESTs, Highly similar to URACIL PHOSPHORIBOSYLTRANSFERASE [Saccharomyces cerevisiae] {IMAGE:576642}	576642 IMAGE Consortium	AA119830
5621 B	9 021WA1FH	ESTs {IMAGE:576745}	576745 IMAGE Consortium	AA123827
5622 B	11 021WA1FH	ESTs {IMAGE:576918}	576918 IMAGE Consortium	AA124161
5623 D	1 021WA1FH	IMAGE EST {IMAGE:574240}	574240 IMAGE Consortium	AA111228
5624 D	3 021WA1FH	ESTs {IMAGE:576157}	576157 IMAGE Consortium	AA119804
5625 D	5 021WA1FH	Mouse putative guanylate binding protein mRNA, complete cds {IMAGE:576022}	576022 IMAGE Consortium	AA119111
5626 D	7 021WA1FH	IMAGE EST {IMAGE:576269}	576269 IMAGE Consortium	AA120051
5627 D	9 021WA1FH	Lymphoid enhancer binding factor 1 {IMAGE:576355}	576355 IMAGE Consortium	AA119479
5628 D	11 021WA1FH	IMAGE EST {IMAGE:576674}	576674 IMAGE Consortium	AA123769
5629 F	1 021WA1FH	ESTs {IMAGE:574045}	574045 IMAGE Consortium	AA119003
5630 F	3 021WA1FH	Mus musculus calpain Lp82 mRNA, complete cds {IMAGE:574491}	574491 IMAGE Consortium	AA117626
5631 F	5 021WA1FH	ESTs {IMAGE:574352}	574352 IMAGE Consortium	AA119641
5632 F	7 021WA1FH	IMAGE EST {IMAGE:574679}	574679 IMAGE Consortium	AA120627
5633 F	9 021WA1FH	IMAGE EST {IMAGE:576962}	576962 IMAGE Consortium	AA122531
5634 F	11 021WA1FH	Mus musculus selenocysteine tRNA gene transcription activating factor (m-Staf) mRNA, complete cds {IMAGE:574939}	574939 IMAGE Consortium	AA118392
5635 H	1 021WA1FH	ESTs, Highly similar to DYNEIN BETA CHAIN, CILIARY [Anthocidaris crassispina] {IMAGE:574464}	574464 IMAGE Consortium	AA117561
5636 H	3 021WA1FH	ESTs {IMAGE:574520}	574520 IMAGE Consortium	AA125159

5637 H	5 021WA1FH	ESTs {IMAGE:574824}	574824 IMAGE Consortium	AA125186
5638 H	7 021WA1FH	IMAGE EST {IMAGE:574974}	574974 IMAGE Consortium	AA124989
5639 H	9 021WA1FH	ESTs {IMAGE:574887}	574887 IMAGE Consortium	AA124900
5640 H	11 021WA1FH	IMAGE EST {IMAGE:577023}	577023 IMAGE Consortium	AA124375
5641 B	1 0213A1FI	IMAGE EST {IMAGE:580613}	580613 IMAGE Consortium	AA139544
5642 B	3 0213A1FI	IMAGE EST {IMAGE:577567}	577567 IMAGE Consortium	AA139179
5643 B	5 0213A1FI	IMAGE EST {IMAGE:581104}	581104 IMAGE Consortium	AA138660
5644 B	7 0213A1FI	ESTs {IMAGE:577974}	577974 IMAGE Consortium	AA122881
5645 B	9 0213A1FI	Mus musculus ERG-associated protein ESET mRNA, complete cds {IMAGE:577866}	577866 IMAGE Consortium	AA116277
5646 B	11 0213A1FI	ESTs {IMAGE:582532}	582532 IMAGE Consortium	AA153093
5647 D	1 0213A1FI	IMAGE EST {IMAGE:582199}	582199 IMAGE Consortium	AA153915
5648 D	3 0213A1FI	IMAGE EST {IMAGE:582541}	582541 IMAGE Consortium	AA153956
5649 D	5 0213A1FI	ESTs {IMAGE:582575}	582575 IMAGE Consortium	AA144169
5650 D	7 0213A1FI	Syntaxin binding protein 2 {IMAGE:579970}	579970 IMAGE Consortium	AA137517
5651 D	9 0213A1FI	D-amino acid oxidase {IMAGE:579983}	579983 IMAGE Consortium	AA122470
5652 D	11 0213A1FI	ESTs {IMAGE:580168}	580168 IMAGE Consortium	AA124387
5653 F	1 0213A1FI	IMAGE EST {IMAGE:577071}	577071 IMAGE Consortium	AA123802
5654 F	3 0213A1FI	Tripeptidyl peptidase II {IMAGE:577425}	577425 IMAGE Consortium	AA139288
5655 F	5 0213A1FI	IMAGE EST {IMAGE:581226}	581226 IMAGE Consortium	AA138750
5656 F	7 0213A1FI	ESTs, Moderately similar to hypothetical protein [M.musculus] {IMAGE:581932}	581932 IMAGE Consortium	AA152980
5657 F	9 0213A1FI	ESTs {IMAGE:581964}	581964 IMAGE Consortium	AA154129
5658 F	11 0213A1FI	ESTs {IMAGE:582726}	582726 IMAGE Consortium	AA152867
5659 H	1 0213A1FI	ESTs, Highly similar to CORONIN [Dictyostelium discoideum] {IMAGE:577627}	577627 IMAGE Consortium	AA140511
5660 H	3 0213A1FI	ESTs {IMAGE:577856}	577856 IMAGE Consortium	AA122848
5661 H	5 0213A1FI	IMAGE EST {IMAGE:582711}	582711 IMAGE Consortium	AA154152
5662 H	7 0213A1FI	ESTs {IMAGE:577181}	577181 IMAGE Consortium	AA124535
5663 H	9 0213A1FI	IMAGE EST {IMAGE:580623}	580623 IMAGE Consortium	AA139711
5664 H	11 0213A1FI	G protein gamma 3 linked gene {IMAGE:581125}	581125 IMAGE Consortium	AA140079
5665 B	1 021AA1FJ	IMAGE EST {IMAGE:576295}	576295 IMAGE Consortium	AA120078
5666 B	3 021AA1FJ	ESTs {IMAGE:574896}	574896 IMAGE Consortium	AA125197
5667 B	5 021AA1FJ	IMAGE EST {IMAGE:575094}	575094 IMAGE Consortium	AA125373
5668 B	7 021AA1FJ	ESTs, Moderately similar to microsomal glutathione S-transferase 3 [H.sapiens] {IMAGE:574792}	574792 IMAGE Consortium	AA120639
5669 B	9 021AA1FJ	IMAGE EST {IMAGE:574748}	574748 IMAGE Consortium	AA118335
5670 B	11 021AA1FJ	ESTs, Weakly similar to HYPOTHETICAL 51.6 KD PROTEIN ZK353.8 IN CHROMOSOME III [Caenorhabditis elegans] {IMAGE:575451}	575451 IMAGE Consortium	AA125221
5671 D	1 021AA1FJ	IMAGE EST {IMAGE:574529}	574529 IMAGE Consortium	AA120574
5672 D	3 021AA1FJ	ESTs, Highly similar to ARGINYL-TRNA SYNTHETASE, MITOCHONDRIAL PRECURSOR [Saccharomyces	576572 IMAGE Consortium	AA119827



		cerevisiae] {IMAGE:576572}		
5673 D	5 021AA1FJ	IMAGE EST {IMAGE:574888}	574888 IMAGE Consortium	AA120755
5674 D	7 021AA1FJ	ESTs, Weakly similar to cDNA EST EMBL:D70402 comes from this gene [C.elegans] {IMAGE:576761}	576761 IMAGE Consortium	AA120013
5675 D	9 021AA1FJ	IMAGE EST {IMAGE:575484}	575484 IMAGE Consortium	AA124670
5676 D	11 021AA1FJ	ESTs, Highly similar to DEAD-box protein p72 [H.sapiens] {IMAGE:573857}	573857 IMAGE Consortium	AA111063
5677 F	1 021AA1FJ	ESTs {IMAGE:575285}	575285 IMAGE Consortium	AA125312
5678 F	3 021AA1FJ	IMAGE EST {IMAGE:575486}	575486 IMAGE Consortium	AA124671
5679 F	5 021AA1FJ	IMAGE EST {IMAGE:575274}	575274 IMAGE Consortium	AA119236
5680 F	7 021AA1FJ	IMAGE EST {IMAGE:575592}	575592 IMAGE Consortium	AA125404
5681 F	9 021AA1FJ	IMAGE EST {IMAGE:575512}	575512 IMAGE Consortium	AA119529
5682 F	11 021AA1FJ	IMAGE EST {IMAGE:576154}	576154 IMAGE Consortium	AA119801
5683 H	1 021AA1FJ	IMAGE EST {IMAGE:574990}	574990 IMAGE Consortium	AA118268
5684 H	3 021AA1FJ	ESTs {IMAGE:575333}	575333 IMAGE Consortium	AA123966
5685 H	5 021AA1FJ	IMAGE EST {IMAGE:580381}	580381 IMAGE Consortium	AA137688
5686 H	7 021AA1FJ	Mus musculus thioredoxin-related protein mRNA, complete cds {IMAGE:577221}	577221 IMAGE Consortium	AA116519
5687 H	9 021AA1FJ	IMAGE EST {IMAGE:581199}	581199 IMAGE Consortium	AA139630
5688 H	11 021AA1FJ	ESTs {IMAGE:581835}	581835 IMAGE Consortium	AA154427
5689 B	1 021HA1FK	ESTs {IMAGE:579608}	579608 IMAGE Consortium	AA116661
5690 B	3 021HA1FK	ESTs, Highly similar to MVP1 PROTEIN [Saccharomyces cerevisiae] {IMAGE:579756}	579756 IMAGE Consortium	AA116950
5691 B	5 021HA1FK	Mus musculus cyclin-dependent kinase-2 alpha (Cdk2-alpha) mRNA, complete cds {IMAGE:577063}	577063 IMAGE Consortium	AA124465
5692 B	7 021HA1FK	ESTs, Highly similar to CALCINEURIN B SUBUNIT ISOFORM 1 [Homo sapiens; Bos taurus; Rattus norvegicus] {IMAGE:580505}	580505 IMAGE Consortium	AA137748
5693 B	9 021HA1FK	Solute carrier family 22, member 1 {IMAGE:581126}	581126 IMAGE Consortium	AA140078
5694 B	11 021HA1FK	ESTs {IMAGE:577485}	577485 IMAGE Consortium	AA137916
5695 D	1 021HA1FK	ESTs {IMAGE:581198}	581198 IMAGE Consortium	AA139627
5696 D	3 021HA1FK	IMAGE EST {IMAGE:582001}	582001 IMAGE Consortium	AA153004
5697 D	5 021HA1FK	IMAGE EST {IMAGE:581766}	581766 IMAGE Consortium	AA153851
5698 D	7 021HA1FK	IMAGE EST {IMAGE:581678}	581678 IMAGE Consortium	AA152806
5699 D	9 021HA1FK	ESTs {IMAGE:578525}	578525 IMAGE Consortium	AA162920
5700 D	11 021HA1FK	ESTs {IMAGE:578367}	578367 IMAGE Consortium	AA124043
5701 F	1 021HA1FK	IMAGE EST {IMAGE:577790}	577790 IMAGE Consortium	AA139233
5702 F	3 021HA1FK	IMAGE EST {IMAGE:577970}	577970 IMAGE Consortium	AA122879
5703 F	5 021HA1FK	IMAGE EST {IMAGE:578146}	578146 IMAGE Consortium	AA125521
5704 F	7 021HA1FK	Ral-interacting protein 1 {IMAGE:582585}	582585 IMAGE Consortium	AA153988
5705 F	9 021HA1FK	ESTs, Moderately similar to calpastatin [R.norvegicus] {IMAGE:577089}	577089 IMAGE Consortium	AA123853
5706 F	11 021HA1FK	ESTs, Moderately similar to RIBOSOMAL PROTEIN S6	577196 IMAGE Consortium	AA116335

5707 H	1 021HA1FK	KINASE [Homo sapiens] {IMAGE:577196} Mus musculus thioredoxin peroxidase (Tpx) mRNA, complete cds {IMAGE:579867}	579867 IMAGE Consortium	AA117023
5708 H	3 021HA1FK	IMAGE EST {IMAGE:577137}	577137 IMAGE Consortium	AA124142
5709 H	5 021HA1FK	IMAGE EST {IMAGE:577590}	577590 IMAGE Consortium	AA139183
5710 H	7 021HA1FK	Mus musculus Mlark mRNA, complete cds {IMAGE:577527}	577527 IMAGE Consortium	AA139442
5711 H	9 021HA1FK	ESTs {IMAGE:577837}	577837 IMAGE Consortium	AA122846
5712 H	11 021HA1FK	IMAGE EST {IMAGE:582746}	582746 IMAGE Consortium	AA152873
5713 B	1 021OA1FL	ESTs {IMAGE:597797}	597797 IMAGE Consortium	AA163875
5714 B	3 021OA1FL	Oncostatin receptor {IMAGE:598146}	598146 IMAGE Consortium	AA265259
5715 B	5 021OA1FL	IMAGE EST {IMAGE:598299}	598299 IMAGE Consortium	AA162662
5716 B	7 021OA1FL	IMAGE EST {IMAGE:596231}	596231 IMAGE Consortium	AA122443
5717 B	9 021OA1FL	Core binding factor beta {IMAGE:596552}	596552 IMAGE Consortium	AA146442
5718 B	11 021OA1FL	ESTs {IMAGE:596539}	596539 IMAGE Consortium	AA138311
5719 D	1 021OA1FL	ESTs {IMAGE:583932}	583932 IMAGE Consortium	AA154057
5720 D	3 021OA1FL	M.musculus GEG-154 mRNA {IMAGE:583867}	583867 IMAGE Consortium	AA145819
5721 D	5 021OA1FL	IMMUNOGLOBULIN J CHAIN PRECURSOR {IMAGE:595848}	595848 IMAGE Consortium	AA137942
5722 D	7 021OA1FL	IMAGE EST {IMAGE:595950}	595950 IMAGE Consortium	AA140540
5723 D	9 021OA1FL	ESTs {IMAGE:596200}	596200 IMAGE Consortium	AA122759
5724 D	11 021OA1FL	ESTs {IMAGE:596400}	596400 IMAGE Consortium	AA138544
5725 F	1 021OA1FL	IMAGE EST {IMAGE:583712}	583712 IMAGE Consortium	AA153422
5726 F	3 021OA1FL	IMAGE EST {IMAGE:583669}	583669 IMAGE Consortium	AA153416
5727 F	5 021OA1FL	ESTs, Moderately similar to The ha1539 protein is related to cyclophilin. [H.sapiens] {IMAGE:583580}	583580 IMAGE Consortium	AA146002
5728 F	7 021OA1FL	IMAGE EST {IMAGE:595944}	595944 IMAGE Consortium	AA140539
5729 F	9 021OA1FL	ESTs {IMAGE:596022}	596022 IMAGE Consortium	AA137291
5730 F	11 021OA1FL	IMAGE EST {IMAGE:598188}	598188 IMAGE Consortium	AA162537
5731 H	1 021OA1FL	ESTs {IMAGE:597384}	597384 IMAGE Consortium	AA153153
5732 H	3 021OA1FL	ESTs, Highly similar to T-CELL SURFACE PROTEIN TACTILE PRECURSOR [Homo sapiens] {IMAGE:597729}	597729 IMAGE Consortium	AA154092
5733 H	5 021OA1FL	IMAGE EST {IMAGE:583765}	583765 IMAGE Consortium	AA153078
5734 H	7 021OA1FL	IMAGE EST {IMAGE:597920}	597920 IMAGE Consortium	AA162154
5735 H	9 021OA1FL	IMAGE EST {IMAGE:595951}	595951 IMAGE Consortium	AA140542
5736 H	11 021OA1FL	IMAGE EST {IMAGE:598452}	598452 IMAGE Consortium	AA161770
5737 B	1 021VA1FM	EST {IMAGE:583534}	583534 IMAGE Consortium	AA146384
5738 B	3 021VA1FM	ESTs {IMAGE:597661}	597661 IMAGE Consortium	AA152976
5739 B	5 021VA1FM	Chemokine (C-C) receptor 5 {IMAGE:597691}	597691 IMAGE Consortium	AA144482
5740 B	7 021VA1FM	IMAGE EST {IMAGE:597843}	597843 IMAGE Consortium	AA164069
5741 B	9 021VA1FM	ESTs, Weakly similar to p76 [H.sapiens] {IMAGE:598117}	598117 IMAGE Consortium	AA265145
5742 B	11 021VA1FM	IMAGE EST {IMAGE:595978}	595978 IMAGE Consortium	AA137467
5743 D	1 021VA1FM	ESTs, Weakly similar to R10D12.12 [C.elegans] {IMAGE:583535}	583535 IMAGE Consortium	AA146388

5744 D	3 021VA1FM	Mus musculus amino levulinate synthase (ALAS-H) mRNA, 3' end {IMAGE:583638}	583638 IMAGE Consortium	AA144013
5745 D	5 021VA1FM	IMAGE EST {IMAGE:583833}	583833 IMAGE Consortium	AA152947
5746 D	7 021VA1FM	IMAGE EST {IMAGE:583769}	583769 IMAGE Consortium	AA146401
5747 D	9 021VA1FM	ESTs {IMAGE:597854}	597854 IMAGE Consortium	AA163827
5748 D	11 021VA1FM	IMAGE EST {IMAGE:598202}	598202 IMAGE Consortium	AA161815
5749 F	1 021VA1FM	Tripeptidyl peptidase II {IMAGE:583926}	583926 IMAGE Consortium	AA154542
5750 F	3 021VA1FM	IMAGE EST {IMAGE:583929}	583929 IMAGE Consortium	AA154059
5751 F	5 021VA1FM	IMAGE EST {IMAGE:583698}	583698 IMAGE Consortium	AA146278
5752 F	7 021VA1FM	ESTs, Highly similar to CYTOCHROME P450 IIC13, MALE-SPECIFIC [Rattus norvegicus] {IMAGE:583799}	583799 IMAGE Consortium	AA146215
5753 F	9 021VA1FM	IMAGE EST {IMAGE:595973}	595973 IMAGE Consortium	AA137298
5754 F	11 021VA1FM	IMAGE EST {IMAGE:595933}	595933 IMAGE Consortium	AA137953
5755 H	1 021VA1FM	Proprotein convertase subtilisin/kexin type 3 {IMAGE:583759}	583759 IMAGE Consortium	AA152814
5756 H	3 021VA1FM	IMAGE EST {IMAGE:583930}	583930 IMAGE Consortium	AA154056
5757 H	5 021VA1FM	ESTs {IMAGE:598060}	598060 IMAGE Consortium	AA265239
5758 H	7 021VA1FM	ESTs {IMAGE:597905}	597905 IMAGE Consortium	AA163652
5759 H	9 021VA1FM	IMAGE EST {IMAGE:598418}	598418 IMAGE Consortium	AA163849
5760 H	11 021VA1FM	IMAGE EST {IMAGE:598448}	598448 IMAGE Consortium	AA162119
5761 B	1 0212A1FN	IMAGE EST {IMAGE:597278}	597278 IMAGE Consortium	AA145344
5762 B	3 0212A1FN	IMAGE EST {IMAGE:597600}	597600 IMAGE Consortium	AA152725
5763 B	5 0212A1FN	IMAGE EST {IMAGE:597634}	597634 IMAGE Consortium	AA152969
5764 B	7 0212A1FN	ESTs {IMAGE:583719}	583719 IMAGE Consortium	AA153429
5765 B	9 0212A1FN	IMAGE EST {IMAGE:583892}	583892 IMAGE Consortium	AA145824
5766 B	11 0212A1FN	IMAGE EST {IMAGE:598078}	598078 IMAGE Consortium	AA265212
5767 D	1 0212A1FN	IMAGE EST {IMAGE:583520}	583520 IMAGE Consortium	AA142684
5768 D	3 0212A1FN	ESTs, Highly similar to HYPOTHETICAL 66.5 KD PROTEIN IN ADE12-RAP1 INTERGENIC REGION [Saccharomyces cerevisiae] {IMAGE:583479}	583479 IMAGE Consortium	AA142678
5769 D	5 0212A1FN	ESTs {IMAGE:583684}	583684 IMAGE Consortium	AA144098
5770 D	7 0212A1FN	ESTs, Moderately similar to CH-TOG PROTEIN [H.sapiens] {IMAGE:583569}	583569 IMAGE Consortium	AA144046
5771 D	9 0212A1FN	Mus musculus PS-2 mRNA, complete cds {IMAGE:597527}	597527 IMAGE Consortium	AA144382
5772 D	11 0212A1FN	IMAGE EST {IMAGE:597849}	597849 IMAGE Consortium	AA163828
5773 F	1 0212A1FN	Cytochrome P450, 2e1, ethanol inducible {IMAGE:596870}	596870 IMAGE Consortium	AA145212
5774 F	3 0212A1FN	ESTs {IMAGE:596879}	596879 IMAGE Consortium	AA138580
5775 F	5 0212A1FN	ESTs, Weakly similar to nuclear factor YC [M.musculus] {IMAGE:583433}	583433 IMAGE Consortium	AA146265
5776 F	7 0212A1FN	IMAGE EST {IMAGE:583781}	583781 IMAGE Consortium	AA152816
5777 F	9 0212A1FN	IMAGE EST {IMAGE:597499}	597499 IMAGE Consortium	AA144372
5778 F	11 0212A1FN	IMAGE EST {IMAGE:598128}	598128 IMAGE Consortium	AA265439
5779 H	1 0212A1FN	IMAGE EST {IMAGE:599075}	599075 IMAGE Consortium	AA172852
5780 H	3 0212A1FN	ESTs {IMAGE:597223}	597223 IMAGE Consortium	AA145252

5781 H	5 0212A1FN	Thymopoietin {IMAGE:597352}	597352 IMAGE Consortium	AA138415
5782 H	7 0212A1FN	ESTs {IMAGE:597579}	597579 IMAGE Consortium	AA152722
5783 H	9 0212A1FN	IMAGE EST {IMAGE:597493}	597493 IMAGE Consortium	AA154419
5784 H	11 0212A1FN	ESTs {IMAGE:597547}	597547 IMAGE Consortium	AA144388
5785 B	1 0219A1FO	ESTs {IMAGE:616648}	616648 IMAGE Consortium	AA170464
5786 B	3 0219A1FO	ESTs {IMAGE:619079}	619079 IMAGE Consortium	AA170629
5787 B	5 0219A1FO	IMAGE EST {IMAGE:617424}	617424 IMAGE Consortium	AA174440
5788 B	7 0219A1FO	ESTs {IMAGE:619687}	619687 IMAGE Consortium	AA175976
5789 B	9 0219A1FO	IMAGE EST {IMAGE:619861}	619861 IMAGE Consortium	AA177984
5790 B	11 0219A1FO	IMAGE EST {IMAGE:618199}	618199 IMAGE Consortium	AA175243
5791 D	1 0219A1FO	ESTs {IMAGE:617657}	617657 IMAGE Consortium	AA170654
5792 D	3 0219A1FO	ESTs {IMAGE:619723}	619723 IMAGE Consortium	AA172624
5793 D	5 0219A1FO	Mus musculus protease-activated receptor 3 (PAR3) mRNA, complete cds {IMAGE:620379}	620379 IMAGE Consortium	AA177828
5794 D	7 0219A1FO	ESTs, Weakly similar to D2085.4 [C.elegans] {IMAGE:618327}	618327 IMAGE Consortium	AA175034
5795 D	9 0219A1FO	IMAGE EST {IMAGE:620596}	620596 IMAGE Consortium	AA178292
5796 D	11 0219A1FO	ESTs {IMAGE:620751}	620751 IMAGE Consortium	AA178265
5797 F	1 0219A1FO	IMAGE EST {IMAGE:617860}	617860 IMAGE Consortium	AA175259
5798 F	3 0219A1FO	Polycystic kidney disease 2 {IMAGE:620097}	620097 IMAGE Consortium	AA176035
5799 F	5 0219A1FO	ESTs {IMAGE:620080}	620080 IMAGE Consortium	AA172490
5800 F	7 0219A1FO	Glycoprotein galactosyltransferase alpha 1, 3 {IMAGE:618535}	618535 IMAGE Consortium	AA175441
5801 F	9 0219A1FO	IMAGE EST {IMAGE:618667}	618667 IMAGE Consortium	AA170344
5802 F	11 0219A1FO	IMAGE EST {IMAGE:619013}	619013 IMAGE Consortium	AA175929
5803 H	1 0219A1FO	IMAGE EST {IMAGE:618386}	618386 IMAGE Consortium	AA175082
5804 H	3 0219A1FO	IMAGE EST {IMAGE:618160}	618160 IMAGE Consortium	AA170488
5805 H	5 0219A1FO	Mus musculus SH2/SH3 adaptor protein (mNck-alpha) mRNA, complete cds {IMAGE:618165}	618165 IMAGE Consortium	AA170493
5806 H	7 0219A1FO	ESTs {IMAGE:618584}	618584 IMAGE Consortium	AA175911
5807 H	9 0219A1FO	IMAGE EST {IMAGE:618691}	618691 IMAGE Consortium	AA170350
5808 H	11 0219A1FO	IMAGE EST {IMAGE:619253}	619253 IMAGE Consortium	AA175709
5809 B	1 021GA1FP	ESTs, Moderately similar to BILIVERDIN REDUCTASE A PRECURSOR [Rattus norvegicus] {IMAGE:620320}	620320 IMAGE Consortium	AA172748
5810 B	3 021GA1FP	ESTs, Weakly similar to CMRF35 ANTIGEN PRECURSOR [H.sapiens] {IMAGE:618824}	618824 IMAGE Consortium	AA175290
5811 B	5 021GA1FP	ESTs, Highly similar to UBIQUITIN-ACTIVATING ENZYME E1 HOMOLOG [Homo sapiens] {IMAGE:618859}	618859 IMAGE Consortium	AA170444
5812 B	7 021GA1FP	IMAGE EST {IMAGE:619141}	619141 IMAGE Consortium	AA175618
5813 B	9 021GA1FP	ESTs, Moderately similar to T20D3.3 [C.elegans] {IMAGE:617288}	617288 IMAGE Consortium	AA174312
5814 B	11 021GA1FP	Myristoylated alanine rich protein kinase C substrate {IMAGE:617200}	617200 IMAGE Consortium	AA170009

5815 D	1	021GA1FP	Murine Glvr-1 mRNA, complete cds {IMAGE:620195}	620195	IMAGE Consortium	AA177949
5816 D	3	021GA1FP	Mouse thymidylate kinase homologue mRNA, complete cds {IMAGE:620057}	620057	IMAGE Consortium	AA172346
5817 D	5	021GA1FP	Ubiquitin specific protease 9, X chromosome {IMAGE:620646}	620646	IMAGE Consortium	AA178383
5818 D	7	021GA1FP	ESTs {IMAGE:620469}	620469	IMAGE Consortium	AA174364
5819 D	9	021GA1FP	ESTs {IMAGE:621175}	621175	IMAGE Consortium	AA178703
5820 D	11	021GA1FP	ESTs {IMAGE:618926}	618926	IMAGE Consortium	AA175466
5821 F	1	021GA1FP	ESTs {IMAGE:618440}	618440	IMAGE Consortium	AA175322
5822 F	3	021GA1FP	ESTs, Highly similar to gene CC1 protein [M.musculus] {IMAGE:618505}	618505	IMAGE Consortium	AA175386
5823 F	5	021GA1FP	ESTs {IMAGE:618609}	618609	IMAGE Consortium	AA174940
5824 F	7	021GA1FP	ESTs {IMAGE:618644}	618644	IMAGE Consortium	AA170288
5825 F	9	021GA1FP	IMAGE EST {IMAGE:619207}	619207	IMAGE Consortium	AA175695
5826 F	11	021GA1FP	IMAGE EST {IMAGE:616968}	616968	IMAGE Consortium	AA166315
5827 H	1	021GA1FP	M.musculus mRNA for cell surface glycoprotein CD53 {IMAGE:620268}	620268	IMAGE Consortium	AA178076
5828 H	3	021GA1FP	ESTs {IMAGE:620131}	620131	IMAGE Consortium	AA172550
5829 H	5	021GA1FP	Myristoylated alanine rich protein kinase C substrate {IMAGE:618681}	618681	IMAGE Consortium	AA175094
5830 H	7	021GA1FP	IMAGE EST {IMAGE:620819}	620819	IMAGE Consortium	AA178322
5831 H	9	021GA1FP	ESTs {IMAGE:621186}	621186	IMAGE Consortium	AA177368
5832 H	11	021GA1FP	IMAGE EST {IMAGE:617016}	617016	IMAGE Consortium	AA166344
5833 B	1	021NA1FQ	ESTs {IMAGE:634992}	634992	IMAGE Consortium	AA154838
5834 B	3	021NA1FQ	ESTs, Highly similar to CYTOCHROME C OXIDASE ASSEMBLY PROTEIN COX11 [Saccharomyces cerevisiae] {IMAGE:635096}	635096	IMAGE Consortium	AA178821
5835 B	5	021NA1FQ	Mus musculus SH3-containing protein SH3P2 mRNA, partial cds {IMAGE:635746}	635746	IMAGE Consortium	AA166372
5836 B	7	021NA1FQ	ESTs, Moderately similar to JAK3 [M.musculus] {IMAGE:635916}	635916	IMAGE Consortium	AA183273
5837 B	9	021NA1FQ	ESTs {IMAGE:636372}	636372	IMAGE Consortium	AA183226
5838 B	11	021NA1FQ	IMAGE EST {IMAGE:634133}	634133	IMAGE Consortium	AA184625
5839 D	1	021NA1FQ	Protein tyrosine phosphatase, receptor type, A {IMAGE:622319}	622319	IMAGE Consortium	AA177749
5840 D	3	021NA1FQ	ESTs {IMAGE:635296}	635296	IMAGE Consortium	AA189776
5841 D	5	021NA1FQ	ESTs {IMAGE:635907}	635907	IMAGE Consortium	AA182195
5842 D	7	021NA1FQ	IMAGE EST {IMAGE:622790}	622790	IMAGE Consortium	AA184267
5843 D	9	021NA1FQ	IMAGE EST {IMAGE:622914}	622914	IMAGE Consortium	AA177703
5844 D	11	021NA1FQ	ESTs {IMAGE:621316}	621316	IMAGE Consortium	AA178549
5845 F	1	021NA1FQ	Immunoglobulin heavy chain 8 (heavy chain of IgG3) {IMAGE:635556}	635556	IMAGE Consortium	AA166427
5846 F	3	021NA1FQ	IMAGE EST {IMAGE:623030}	623030	IMAGE Consortium	AA185885
5847 F	5	021NA1FQ	ESTs {IMAGE:621412}	621412	IMAGE Consortium	AA178467

5848 F	7 021NA1FQ	Mus musculus major histocompatibility locus class II region; Fas-binding protein Daxx (DAXX) gene, partial cds; Bing1 (BING1), tapasin (tapasin), RalGDS-like factor (RLF), KE2 (KE2), BING4 (BING4), beta1, 3-galactosyl transferase {IMAGE:634344}	634344 IMAGE Consortium	AA185515
5849 F	9 021NA1FQ	IMAGE EST {IMAGE:621738}	621738 IMAGE Consortium	AA177582
5850 F	11 021NA1FQ	ESTs {IMAGE:621910}	621910 IMAGE Consortium	AA178392
5851 H	1 021NA1FQ	IMAGE EST {IMAGE:621786}	621786 IMAGE Consortium	AA177340
5852 H	3 021NA1FQ	IMAGE EST {IMAGE:634849}	634849 IMAGE Consortium	AA163500
5853 H	5 021NA1FQ	IMAGE EST {IMAGE:621969}	621969 IMAGE Consortium	AA183914
5854 H	7 021NA1FQ	IMAGE EST {IMAGE:635020}	635020 IMAGE Consortium	AA163237
5855 H	9 021NA1FQ	ESTs {IMAGE:622383}	622383 IMAGE Consortium	AA184207
5856 H	11 021NA1FQ	ESTs {IMAGE:635654}	635654 IMAGE Consortium	AA166545
5857 B	1 021UA1FR	ESTs, Moderately similar to GEPHYRIN [Rattus norvegicus] {IMAGE:618257}	618257 IMAGE Consortium	AA170590
5858 B	3 021UA1FR	ESTs {IMAGE:618602}	618602 IMAGE Consortium	AA174970
5859 B	5 021UA1FR	ESTs {IMAGE:620905}	620905 IMAGE Consortium	AA178356
5860 B	7 021UA1FR	IMAGE EST {IMAGE:620868}	620868 IMAGE Consortium	AA178360
5861 B	9 021UA1FR	ESTs, Weakly similar to ADP-RIBOSYLATION FACTOR 2 [Saccharomyces cerevisiae] {IMAGE:619214}	619214 IMAGE Consortium	AA175745
5862 B	11 021UA1FR	IMAGE EST {IMAGE:617316}	617316 IMAGE Consortium	AA174320
5863 D	1 021UA1FR	IMAGE EST {IMAGE:618626}	618626 IMAGE Consortium	AA174978
5864 D	3 021UA1FR	IMAGE EST {IMAGE:621097}	621097 IMAGE Consortium	AA178680
5865 D	5 021UA1FR	Nitric oxide synthase 3, endothelial cell {IMAGE:620940}	620940 IMAGE Consortium	AA177240
5866 D	7 021UA1FR	ESTs {IMAGE:618927}	618927 IMAGE Consortium	AA175470
5867 D	9 021UA1FR	DIK1-like homolog (Drosophila) {IMAGE:619006}	619006 IMAGE Consortium	AA170607
5868 D	11 021UA1FR	IMAGE EST {IMAGE:619398}	619398 IMAGE Consortium	AA175541
5869 F	1 021UA1FR	ESTs {IMAGE:618675}	618675 IMAGE Consortium	AA174996
5870 F	3 021UA1FR	IMAGE EST {IMAGE:618986}	618986 IMAGE Consortium	AA175920
5871 F	5 021UA1FR	Alpha-galactosidase {IMAGE:617018}	617018 IMAGE Consortium	AA166345
5872 F	7 021UA1FR	ESTs {IMAGE:617455}	617455 IMAGE Consortium	AA174582
5873 F	9 021UA1FR	IMAGE EST {IMAGE:617583}	617583 IMAGE Consortium	AA174881
5874 F	11 021UA1FR	CD1d1 antigen {IMAGE:619932}	619932 IMAGE Consortium	AA178002
5875 H	1 021UA1FR	ESTs {IMAGE:621629}	621629 IMAGE Consortium	AA183687
5876 H	3 021UA1FR	IMAGE EST {IMAGE:634338}	634338 IMAGE Consortium	AA182369
5877 H	5 021UA1FR	ESTs {IMAGE:634728}	634728 IMAGE Consortium	AA163094
5878 H	7 021UA1FR	IMAGE EST {IMAGE:634733}	634733 IMAGE Consortium	AA163279
5879 H	9 021UA1FR	ESTs {IMAGE:622334}	622334 IMAGE Consortium	AA183628
5880 H	11 021UA1FR	ESTs, Highly similar to RAS-RELATED PROTEIN RAP-1A [Homo sapiens; Rattus norvegicus; Bos taurus] {IMAGE:621983}	621983 IMAGE Consortium	AA177450
5881 B	1 0211A1FS	ESTs, Highly similar to INTERFERON-ACTIVATABLE PROTEIN 204 [Mus musculus] {IMAGE:622037}	622037 IMAGE Consortium	AA183835

5882 B	3 0211A1FS	ESTs {IMAGE:634747}	634747 IMAGE Consortium	AA154647
5883 B	5 0211A1FS	IMAGE EST {IMAGE:635218}	635218 IMAGE Consortium	AA189710
5884 B	7 0211A1FS	ESTs {IMAGE:635774}	635774 IMAGE Consortium	AA166378
5885 B	9 0211A1FS	IMAGE EST {IMAGE:636318}	636318 IMAGE Consortium	AA189229
5886 B	11 0211A1FS	Interferon concensus sequence binding protein {IMAGE:621246}	621246 IMAGE Consortium	AA178779
5887 D	1 0211A1FS	ESTs {IMAGE:622542}	622542 IMAGE Consortium	AA183587
5888 D	3 0211A1FS	IMAGE EST {IMAGE:622624}	622624 IMAGE Consortium	AA177481
5889 D	5 0211A1FS	IMAGE EST {IMAGE:622605}	622605 IMAGE Consortium	AA177479
5890 D	7 0211A1FS	ESTs {IMAGE:636007}	636007 IMAGE Consortium	AA183350
5891 D	9 0211A1FS	MutS homolog 3 (E. coli) {IMAGE:636392}	636392 IMAGE Consortium	AA183199
5892 D	11 0211A1FS	IMAGE EST {IMAGE:621542}	621542 IMAGE Consortium	AA178786
5893 F	1 0211A1FS	GA repeat binding protein, beta 1 {IMAGE:622424}	622424 IMAGE Consortium	AA184214
5894 F	3 0211A1FS	IMAGE EST {IMAGE:622653}	622653 IMAGE Consortium	AA177538
5895 F	5 0211A1FS	IMAGE EST {IMAGE:636080}	636080 IMAGE Consortium	AA189449
5896 F	7 0211A1FS	ESTs {IMAGE:636165}	636165 IMAGE Consortium	AA183114
5897 F	9 0211A1FS	Mouse mRNA for RNA polymerase II subuunit RPB14, complete cds {IMAGE:636513}	636513 IMAGE Consortium	AA182240
5898 F	11 0211A1FS	M.musculus mRNA for casein kinase I-alpha {IMAGE:636472}	636472 IMAGE Consortium	AA185259
5899 H	1 0211A1FS	ESTs, Weakly similar to hypothetical protein [M.musculus] {IMAGE:635141}	635141 IMAGE Consortium	AA162504
5900 H	3 0211A1FS	IMAGE EST {IMAGE:635621}	635621 IMAGE Consortium	AA165777
5901 H	5 0211A1FS	IMAGE EST {IMAGE:635638}	635638 IMAGE Consortium	AA163134
5902 H	7 0211A1FS	IMAGE EST {IMAGE:623003}	623003 IMAGE Consortium	AA184418
5903 H	9 0211A1FS	ESTs {IMAGE:621464}	621464 IMAGE Consortium	AA178585
5904 H	11 0211A1FS	IMAGE EST {IMAGE:621303}	621303 IMAGE Consortium	AA178604
5905 B	1 0218A1FT	ESTs, Highly similar to INTERFERON-ACTIVATABLE PROTEIN 204 [Mus musculus] {IMAGE:622643}	622643 IMAGE Consortium	AA183597
5906 B	3 0218A1FT	Interleukin 4 receptor, alpha {IMAGE:635530}	635530 IMAGE Consortium	AA166418
5907 B	5 0218A1FT	ESTs {IMAGE:636105}	636105 IMAGE Consortium	AA189456
5908 B	7 0218A1FT	IMAGE EST {IMAGE:636193}	636193 IMAGE Consortium	AA183187
5909 B	9 0218A1FT	ESTs {IMAGE:636323}	636323 IMAGE Consortium	AA190115
5910 B	11 0218A1FT	IMAGE EST {IMAGE:621297}	621297 IMAGE Consortium	AA178601
5911 D	1 0218A1FT	IMAGE EST {IMAGE:622548}	622548 IMAGE Consortium	AA184337
5912 D	3 0218A1FT	ESTs {IMAGE:622722}	622722 IMAGE Consortium	AA177554
5913 D	5 0218A1FT	ESTs {IMAGE:622920}	622920 IMAGE Consortium	AA184365
5914 D	7 0218A1FT	ESTs, Weakly similar to EGF repeat transmembrane protein [M.musculus] {IMAGE:636129}	636129 IMAGE Consortium	AA189462
5915 D	9 0218A1FT	ESTs {IMAGE:638900}	638900 IMAGE Consortium	AA200955
5916 D	11 0218A1FT	IMAGE EST {IMAGE:636829}	636829 IMAGE Consortium	AA183303
5917 F	1 0218A1FT	IMAGE EST {IMAGE:640674}	640674 IMAGE Consortium	AA204302
5918 F	3 0218A1FT	IMAGE EST {IMAGE:640987}	640987 IMAGE Consortium	AA204389

5919 F	5 0218A1FT	IMAGE EST {IMAGE:639008}	639008 IMAGE Consortium	AA200172
5920 F	7 0218A1FT	IMAGE EST {IMAGE:636584}	636584 IMAGE Consortium	AA183038
5921 F	9 0218A1FT	ESTs, Highly similar to major vault protein [R.norvegicus] {IMAGE:639502}	639502 IMAGE Consortium	AA200827
5922 F	11 0218A1FT	ESTs, Weakly similar to predicted using Genefinder [C.elegans] {IMAGE:636955}	636955 IMAGE Consortium	AA184998
5923 H	1 0218A1FT	Immunoglobulin heavy chain 1 (serum IgG2a) {IMAGE:636695}	636695 IMAGE Consortium	AA182334
5924 H	3 0218A1FT	IMAGE EST {IMAGE:636944}	636944 IMAGE Consortium	AA185118
5925 H	5 0218A1FT	ESTs, Moderately similar to HYPOTHETICAL 22.7 KD PROTEIN C28H8.9 IN CHROMOSOME III [Caenorhabditis elegans] {IMAGE:637469}	637469 IMAGE Consortium	AA184903
5926 H	7 0218A1FT	ESTs {IMAGE:640013}	640013 IMAGE Consortium	AA198524
5927 H	9 0218A1FT	IMAGE EST {IMAGE:640735}	640735 IMAGE Consortium	AA203882
5928 H	11 0218A1FT	ESTs {IMAGE:638350}	638350 IMAGE Consortium	AA200703
5929 B	1 021FA1FU	IMAGE EST {IMAGE:641081}	641081 IMAGE Consortium	AA204463
5930 B	3 021FA1FU	Zinc finger protein 36 {IMAGE:641523}	641523 IMAGE Consortium	AA200282
5931 B	5 021FA1FU	Neutrophil cytosolic factor 4 {IMAGE:641512}	641512 IMAGE Consortium	AA200535
5932 B	7 021FA1FU	ESTs, Moderately similar to SART-1 [H.sapiens] {IMAGE:639601}	639601 IMAGE Consortium	AA198906
5933 B	9 021FA1FU	IMAGE EST {IMAGE:636714}	636714 IMAGE Consortium	AA182335
5934 B	11 021FA1FU	IMAGE EST {IMAGE:639867}	639867 IMAGE Consortium	AA197644
5935 D	1 021FA1FU	Dlk1-like homolog (Drosophila) {IMAGE:636875}	636875 IMAGE Consortium	AA189431
5936 D	3 021FA1FU	IMAGE EST {IMAGE:637254}	637254 IMAGE Consortium	AA189791
5937 D	5 021FA1FU	CD6 antigen {IMAGE:637369}	637369 IMAGE Consortium	AA185065
5938 D	7 021FA1FU	IMAGE EST {IMAGE:637523}	637523 IMAGE Consortium	AA184738
5939 D	9 021FA1FU	IMAGE EST {IMAGE:640091}	640091 IMAGE Consortium	AA197362
5940 D	11 021FA1FU	ESTs {IMAGE:638452}	638452 IMAGE Consortium	AA216986
5941 F	1 021FA1FU	ESTs {IMAGE:637237}	637237 IMAGE Consortium	AA189902
5942 F	3 021FA1FU	IMAGE EST {IMAGE:637394}	637394 IMAGE Consortium	AA190264
5943 F	5 021FA1FU	Mus musculus mRNA for G-protein coupled receptor kinase 6-B {IMAGE:640178}	640178 IMAGE Consortium	AA197538
5944 F	7 021FA1FU	IMAGE EST {IMAGE:640331}	640331 IMAGE Consortium	AA199476
5945 F	9 021FA1FU	ESTs {IMAGE:640026}	640026 IMAGE Consortium	AA197412
5946 F	11 021FA1FU	IMAGE EST {IMAGE:638267}	638267 IMAGE Consortium	AA200197
5947 H	1 021FA1FU	ESTs {IMAGE:637888}	637888 IMAGE Consortium	AA189578
5948 H	3 021FA1FU	ESTs {IMAGE:638334}	638334 IMAGE Consortium	AA200779
5949 H	5 021FA1FU	IMAGE EST {IMAGE:638244}	638244 IMAGE Consortium	AA200336
5950 H	7 021FA1FU	IMAGE EST {IMAGE:640971}	640971 IMAGE Consortium	AA204223
5951 H	9 021FA1FU	ESTs {IMAGE:641130}	641130 IMAGE Consortium	AA204476
5952 H	11 021FA1FU	ESTs {IMAGE:641530}	641530 IMAGE Consortium	AA200475
5953 B	1 021MA1FV	ESTs, Weakly similar to HYPOTHETICAL PROTEIN F44G4.1 IN CHROMOSOME II [C.elegans] {IMAGE:637493}	637493 IMAGE Consortium	AA190292



5954 B	3 021MA1FV	Mus musculus interferon regulatory factor 5 (mirf5) mRNA, complete cds {IMAGE:637338}	637338 IMAGE Consortium	AA185025
5955 B	5 021MA1FV	Apoptosis inhibitor 1 {IMAGE:640046}	640046 IMAGE Consortium	AA197349
5956 B	7 021MA1FV	IMAGE EST {IMAGE:640125}	640125 IMAGE Consortium	AA197697
5957 B	9 021MA1FV	ESTs {IMAGE:640496}	640496 IMAGE Consortium	AA204424
5958 B	11 021MA1FV	IMAGE EST {IMAGE:638496}	638496 IMAGE Consortium	AA200077
5959 D	1 021MA1FV	ESTs {IMAGE:637173}	637173 IMAGE Consortium	AA189628
5960 D	3 021MA1FV	IMAGE EST {IMAGE:640062}	640062 IMAGE Consortium	AA198531
5961 D	5 021MA1FV	ESTs {IMAGE:638401}	638401 IMAGE Consortium	AA217009
5962 D	7 021MA1FV	ESTs {IMAGE:638264}	638264 IMAGE Consortium	AA200342
5963 D	9 021MA1FV	ESTs {IMAGE:638593}	638593 IMAGE Consortium	AA200890
5964 D	11 021MA1FV	Protein tyrosine phosphatase, receptor type, C {IMAGE:639168}	639168 IMAGE Consortium	AA201097
5965 F	1 021MA1FV	Mouse mRNA for Bach protein 2, complete cds {IMAGE:640200}	640200 IMAGE Consortium	AA198553
5966 F	3 021MA1FV	ESTs {IMAGE:640023}	640023 IMAGE Consortium	AA198550
5967 F	5 021MA1FV	ESTs {IMAGE:640682}	640682 IMAGE Consortium	AA203817
5968 F	7 021MA1FV	ESTs, Highly similar to The KIAA0136 gene product is novel. [H.sapiens] {IMAGE:638129}	638129 IMAGE Consortium	AA200976
5969 F	9 021MA1FV	IMAGE EST {IMAGE:640910}	640910 IMAGE Consortium	AA210519
5970 F	11 021MA1FV	IMAGE EST {IMAGE:641223}	641223 IMAGE Consortium	AA200448
5971 H	1 021MA1FV	ESTs, Moderately similar to CYTOCHROME B5, OUTER MITOCHONDRIAL MEMBRANE [Rattus norvegicus] {IMAGE:640762}	640762 IMAGE Consortium	AA203975
5972 H	3 021MA1FV	IMAGE EST {IMAGE:638739}	638739 IMAGE Consortium	AA200942
5973 H	5 021MA1FV	ESTs, Moderately similar to p116Rip [M.musculus] {IMAGE:638631}	638631 IMAGE Consortium	AA200000
5974 H	7 021MA1FV	IMAGE EST {IMAGE:639145}	639145 IMAGE Consortium	AA201091
5975 H	9 021MA1FV	IMAGE EST {IMAGE:639062}	639062 IMAGE Consortium	AA199299
5976 H	11 021MA1FV	ESTs {IMAGE:636578}	636578 IMAGE Consortium	AA183098
5977 B	1 021TA1FW	IMAGE EST {IMAGE:642330}	642330 IMAGE Consortium	AA185249
5978 B	3 021TA1FW	ESTs {IMAGE:642795}	642795 IMAGE Consortium	AA185624
5979 B	5 021TA1FW	ESTs, Highly similar to putative ATP/GTP-binding protein [H.sapiens] {IMAGE:642977}	642977 IMAGE Consortium	AA184876
5980 B	7 021TA1FW	IMAGE EST {IMAGE:643106}	643106 IMAGE Consortium	AA189867
5981 B	9 021TA1FW	ESTs {IMAGE:657773}	657773 IMAGE Consortium	AA199196
5982 B	11 021TA1FW	ESTs {IMAGE:658233}	658233 IMAGE Consortium	AA198258
5983 D	1 021TA1FW	ESTs {IMAGE:653427}	653427 IMAGE Consortium	AA240351
5984 D	3 021TA1FW	ESTs {IMAGE:642820}	642820 IMAGE Consortium	AA185638
5985 D	5 021TA1FW	ESTs {IMAGE:643274}	643274 IMAGE Consortium	AA189587
5986 D	7 021TA1FW	ESTs {IMAGE:643437}	643437 IMAGE Consortium	AA185995
5987 D	9 021TA1FW	Mus musculus protein phosphatase 5 (PP5) mRNA, complete cds {IMAGE:658090}	658090 IMAGE Consortium	AA198158

5988 D	11 021TA1FW	IMAGE EST {IMAGE:658172}	658172 IMAGE Consortium	AA198291
5989 F	1 021TA1FW	ESTs, Moderately similar to BETA GALACTOSIDASE-RELATED PROTEIN PRECURSOR [Homo sapiens] {IMAGE:642988}	642988 IMAGE Consortium	AA185584
5990 F	3 021TA1FW	IMAGE EST {IMAGE:657528}	657528 IMAGE Consortium	AA199416
5991 F	5 021TA1FW	Retinitis pigmentosa 3 homolog {IMAGE:643295}	643295 IMAGE Consortium	AA189891
5992 F	7 021TA1FW	IMAGE EST {IMAGE:658378}	658378 IMAGE Consortium	AA198312
5993 F	9 021TA1FW	Hormone receptor {IMAGE:641865}	641865 IMAGE Consortium	AA209882
5994 F	11 021TA1FW	ESTs, Moderately similar to NOF1 [H.sapiens] {IMAGE:641826}	641826 IMAGE Consortium	AA210176
5995 H	1 021TA1FW	IMAGE EST {IMAGE:643309}	643309 IMAGE Consortium	AA189880
5996 H	3 021TA1FW	Mus musculus SIK similar protein mRNA, complete cds {IMAGE:643463}	643463 IMAGE Consortium	AA186002
5997 H	5 021TA1FW	IMAGE EST {IMAGE:643837}	643837 IMAGE Consortium	AA204151
5998 H	7 021TA1FW	ESTs, Moderately similar to Similarity to S.pombe hypothetical protein C2F7.02C [C.elegans] {IMAGE:641889}	641889 IMAGE Consortium	AA209897
5999 H	9 021TA1FW	IMAGE EST {IMAGE:641755}	641755 IMAGE Consortium	AA210038
6000 H	11 021TA1FW	ESTs, Weakly similar to weak similarity to ribosome releasing factors [C.elegans] {IMAGE:652555}	652555 IMAGE Consortium	AA217489
6001 B	1 0210A1FX	ESTs {IMAGE:644928}	644928 IMAGE Consortium	AA210309
6002 B	3 0210A1FX	ESTs {IMAGE:644916}	644916 IMAGE Consortium	AA210321
6003 B	5 0210A1FX	IMAGE EST {IMAGE:652561}	652561 IMAGE Consortium	AA217230
6004 B	7 0210A1FX	IMAGE EST {IMAGE:642129}	642129 IMAGE Consortium	AA212159
6005 B	9 0210A1FX	IMAGE EST {IMAGE:642313}	642313 IMAGE Consortium	AA185106
6006 B	11 0210A1FX	IMAGE EST {IMAGE:653339}	653339 IMAGE Consortium	AA241237
6007 D	1 0210A1FX	IMAGE EST {IMAGE:641828}	641828 IMAGE Consortium	AA210177
6008 D	3 0210A1FX	IMAGE EST {IMAGE:642177}	642177 IMAGE Consortium	AA212233
6009 D	5 0210A1FX	IMAGE EST {IMAGE:652906}	652906 IMAGE Consortium	AA217082
6010 D	7 0210A1FX	Mus musculus mRNA for platelet glycoprotein IX, complete cds {IMAGE:642372}	642372 IMAGE Consortium	AA184800
6011 D	9 0210A1FX	IMAGE EST {IMAGE:642382}	642382 IMAGE Consortium	AA185200
6012 D	11 0210A1FX	ESTs, Moderately similar to protein kinase [H.sapiens] {IMAGE:653513}	653513 IMAGE Consortium	AA217475
6013 F	1 0210A1FX	ESTs, Highly similar to axonemal dynein heavy chain [H.sapiens] {IMAGE:642201}	642201 IMAGE Consortium	AA212241
6014 F	3 0210A1FX	ESTs {IMAGE:642409}	642409 IMAGE Consortium	AA184882
6015 F	5 0210A1FX	ESTs {IMAGE:653444}	653444 IMAGE Consortium	AA240530
6016 F	7 0210A1FX	IMAGE EST {IMAGE:653561}	653561 IMAGE Consortium	AA217780
6017 F	9 0210A1FX	ESTs {IMAGE:657889}	657889 IMAGE Consortium	AA199266
6018 F	11 0210A1FX	ESTs, Weakly similar to Similarity to Yeast cobalt uptake protein cot1 [C.elegans] {IMAGE:643732}	643732 IMAGE Consortium	AA204011
6019 H	1 0210A1FX	ESTs, Highly similar to PTB-ASSOCIATED SPLICING FACTOR [Homo sapiens] {IMAGE:643398}	643398 IMAGE Consortium	AA190087
6020 H	3 0210A1FX	Mus musculus IER5 (Ier5) mRNA, complete cds	643745 IMAGE Consortium	AA203798

		{IMAGE:643745}		
6021 H	5 0210A1FX	IMAGE EST {IMAGE:641868}	641868 IMAGE Consortium	AA209880
6022 H	7 0210A1FX	ESTs, Moderately similar to cofactor E [H.sapiens] {IMAGE:644967}	644967 IMAGE Consortium	AA210383
6023 H	9 0210A1FX	ESTs {IMAGE:652539}	652539 IMAGE Consortium	AA217223
6024 H	11 0210A1FX	IMAGE EST {IMAGE:652528}	652528 IMAGE Consortium	AA217458
6025 B	1 0217A1FY	IMAGE EST {IMAGE:851140}	851140 IMAGE Consortium	AA462751
6026 B	3 0217A1FY	ESTs {IMAGE:851421}	851421 IMAGE Consortium	AA461767
6027 B	5 0217A1FY	Orosomucoid 1 {IMAGE:851877}	851877 IMAGE Consortium	AA522210
6028 B	7 0217A1FY	Mus musculus Ras-like GTP-binding protein Rad mRNA, complete cds {IMAGE:850142}	850142 IMAGE Consortium	AA450695
6029 B	9 0217A1FY	ESTs {IMAGE:851770}	851770 IMAGE Consortium	AA501113
6030 B	11 0217A1FY	ESTs {IMAGE:850390}	850390 IMAGE Consortium	AA451233
6031 D	1 0217A1FY	ESTs {IMAGE:851282}	851282 IMAGE Consortium	AA463077
6032 D	3 0217A1FY	ESTs, Highly similar to THERMOSENSITIVE GLUCONOKINASE [Escherichia coli] {IMAGE:850210}	850210 IMAGE Consortium	AA451431
6033 D	5 0217A1FY	IMAGE EST {IMAGE:850988}	850988 IMAGE Consortium	AA462393
6034 D	7 0217A1FY	ESTs {IMAGE:851593}	851593 IMAGE Consortium	AA500704
6035 D	9 0217A1FY	IMAGE EST {IMAGE:850156}	850156 IMAGE Consortium	AA451381
6036 D	11 0217A1FY	IMAGE EST {IMAGE:851585}	851585 IMAGE Consortium	AA461917
6037 F	1 0217A1FY	IMAGE EST {IMAGE:860324}	860324 IMAGE Consortium	AA500990
6038 F	3 0217A1FY	ESTs, Moderately similar to ACTIN-LIKE PROTEIN 14D [Drosophila melanogaster] {IMAGE:850164}	850164 IMAGE Consortium	AA451385
6039 F	5 0217A1FY	ESTs, Highly similar to HYPOTHETICAL 64.5 KD PROTEIN ZK652.9 IN CHROMOSOME III [Caenorhabditis elegans] {IMAGE:851115}	851115 IMAGE Consortium	AA462739
6040 F	7 0217A1FY	ESTs {IMAGE:851708}	851708 IMAGE Consortium	AA522156
6041 F	9 0217A1FY	IMAGE EST {IMAGE:873864}	873864 IMAGE Consortium	AA472027
6042 F	11 0217A1FY	IMAGE EST {IMAGE:874192}	874192 IMAGE Consortium	AA472299
6043 H	1 0217A1FY	IMAGE EST {IMAGE:873587}	873587 IMAGE Consortium	AA472773
6044 H	3 0217A1FY	ESTs {IMAGE:873973}	873973 IMAGE Consortium	AA472164
6045 H	5 0217A1FY	ESTs {IMAGE:874356}	874356 IMAGE Consortium	AA474934
6046 H	7 0217A1FY	ESTs, Highly similar to PROTEIN PHOSPHATASE PP2A, 55 KD REGULATORY SUBUNIT, ALPHA ISOFORM [Rattus norvegicus] {IMAGE:871532}	871532 IMAGE Consortium	AA462108
6047 H	9 0217A1FY	Mus musculus cytochrome P450 monooxygenase (CYP2J5) mRNA, complete cds {IMAGE:872125}	872125 IMAGE Consortium	AA473629
6048 H	11 0217A1FY	Splicing factor, arginine/serine-rich 3 (SRp20) {IMAGE:873314}	873314 IMAGE Consortium	AA473794
6049 B	1 021EA1FZ	IMAGE EST {IMAGE:657952}	657952 IMAGE Consortium	AA198145
6050 B	3 021EA1FZ	HGF-regulated tyrosine kinase substrate {IMAGE:658311}	658311 IMAGE Consortium	AA198392
6051 B	5 021EA1FZ	ESTs {IMAGE:658638}	658638 IMAGE Consortium	AA230822
6052 B	7 021EA1FZ	IMAGE EST {IMAGE:644883}	644883 IMAGE Consortium	AA210604

6053 B	9 021EA1FZ	ESTs {IMAGE:644825}	644825 IMAGE Consortium	AA210267
6054 B	11 021EA1FZ	ESTs {IMAGE:652565}	652565 IMAGE Consortium	AA217231
6055 D	1 021EA1FZ	ESTs {IMAGE:644382}	644382 IMAGE Consortium	AA265295
6056 D	3 021EA1FZ	IMAGE EST {IMAGE:641605}	641605 IMAGE Consortium	AA209869
6057 D	5 021EA1FZ	ESTs {IMAGE:641585}	641585 IMAGE Consortium	AA221282
6058 D	7 021EA1FZ	ESTs, Moderately similar to HYPOTHETICAL 29.5 KD PROTEIN C05B5.7 IN CHROMOSOME III [Caenorhabditis elegans] {IMAGE:652543}	652543 IMAGE Consortium	AA217225
6059 D	9 021EA1FZ	IMAGE EST {IMAGE:652802}	652802 IMAGE Consortium	AA217509
6060 D	11 021EA1FZ	Calcium modulating ligand {IMAGE:652907}	652907 IMAGE Consortium	AA217093
6061 F	1 021EA1FZ	ESTs, Highly similar to HYPOTHETICAL 58.6 KD PROTEIN ZK632.3 IN CHROMOSOME III [Caenorhabditis elegans] {IMAGE:831764}	831764 IMAGE Consortium	AA444950
6062 F	3 021EA1FZ	M.musculus mRNA for WDNM1 protein {IMAGE:832158}	832158 IMAGE Consortium	AA423584
6063 F	5 021EA1FZ	ESTs {IMAGE:832048}	832048 IMAGE Consortium	AA444721
6064 F	7 021EA1FZ	Leucocyte specific transcript 1 {IMAGE:846366}	846366 IMAGE Consortium	AA435060
6065 F	9 021EA1FZ	ESTs {IMAGE:831741}	831741 IMAGE Consortium	AA444941
6066 F	11 021EA1FZ	IMAGE EST {IMAGE:846536}	846536 IMAGE Consortium	AA433553
6067 H	1 021EA1FZ	ESTs, Highly similar to PHOSPHOLIPASE A-2-ACTIVATING PROTEIN [M.musculus] {IMAGE:831959}	831959 IMAGE Consortium	AA444488
6068 H	3 021EA1FZ	ESTs, Weakly similar to PYRROLINE-5-CARBOXYLATE REDUCTASE [Glycine max] {IMAGE:846965}	846965 IMAGE Consortium	AA433727
6069 H	5 021EA1FZ	ESTs {IMAGE:848629}	848629 IMAGE Consortium	AA445398
6070 H	7 021EA1FZ	IMAGE EST {IMAGE:849758}	849758 IMAGE Consortium	AA458194
6071 H	9 021EA1FZ	ESTs {IMAGE:847082}	847082 IMAGE Consortium	AA419788
6072 H	11 021EA1FZ	ESTs {IMAGE:832008}	832008 IMAGE Consortium	AA444689
6073 B	1 021EA1G0	IMAGE EST {IMAGE:874190}	874190 IMAGE Consortium	AA472298
6074 B	3 021EA1G0	ESTs {IMAGE:874540}	874540 IMAGE Consortium	AA472314
6075 B	5 021EA1G0	IMAGE EST {IMAGE:871509}	871509 IMAGE Consortium	AA462096
6076 B	7 021EA1G0	Mus musculus mRNA for pyruvate kinase, complete cds {IMAGE:873690}	873690 IMAGE Consortium	AA475121
6077 B	9 021EA1G0	IMAGE EST {IMAGE:873956}	873956 IMAGE Consortium	AA475183
6078 B	11 021EA1G0	Lactotransferrin {IMAGE:874383}	874383 IMAGE Consortium	AA474964
6079 D	1 021EA1G0	Mus musculus HS1-associating protein (mHAX-1s) mRNA, complete cds {IMAGE:874591}	874591 IMAGE Consortium	AA472437
6080 D	3 021EA1G0	IMAGE EST {IMAGE:871753}	871753 IMAGE Consortium	AA462598
6081 D	5 021EA1G0	ESTs {IMAGE:864361}	864361 IMAGE Consortium	AA458136
6082 D	7 021EA1G0	ESTs {IMAGE:864608}	864608 IMAGE Consortium	AA458317
6083 D	9 021EA1G0	IMAGE EST {IMAGE:874458}	874458 IMAGE Consortium	AA475099
6084 D	11 021EA1G0	ESTs {IMAGE:864273}	864273 IMAGE Consortium	AA458080
6085 F	1 021EA1G0	IMAGE EST {IMAGE:871700}	871700 IMAGE Consortium	AA462371
6086 F	3 021EA1G0	IMAGE EST {IMAGE:875285}	875285 IMAGE Consortium	AA475864
6087 F	5 021EA1G0	Mouse mRNA for Dhm1 protein, complete cds	875395 IMAGE Consortium	AA475568

6088 F	7 021EA1G0	{IMAGE:875395} Mus Musculus mRNA for hypothetical protein {IMAGE:873849}	873849 IMAGE Consortium	AA471960
6089 F	9 021EA1G0	Pre-mRNA protein kinase {IMAGE:876060}	876060 IMAGE Consortium	AA475423
6090 F	11 021EA1G0	ESTs {IMAGE:876395}	876395 IMAGE Consortium	AA475823
6091 H	1 021EA1G0	ESTs {IMAGE:876715}	876715 IMAGE Consortium	AA497410
6092 H	3 021EA1G0	ESTs {IMAGE:888864}	888864 IMAGE Consortium	AA466288
6093 H	5 021EA1G0	IMAGE EST {IMAGE:889311}	889311 IMAGE Consortium	AA498300
6094 H	7 021EA1G0	FRUCTOSE-BISPHOSPHATE ALDOLASE A {IMAGE:903419}	903419 IMAGE Consortium	AA518639
6095 H	9 021EA1G0	ESTs {IMAGE:876063}	876063 IMAGE Consortium	AA475435
6096 H	11 021EA1G0	ESTs {IMAGE:891193}	891193 IMAGE Consortium	AA510283
6097 B	1 021LA1G1	ESTs {IMAGE:876190}	876190 IMAGE Consortium	AA475616
6098 B	3 021LA1G1	ESTs {IMAGE:876596}	876596 IMAGE Consortium	AA476030
6099 B	5 021LA1G1	Mus musculus aquaporin-8 mRNA, complete cds {IMAGE:888820}	888820 IMAGE Consortium	AA498131
6100 B	7 021LA1G1	ESTs, Highly similar to HYPOTHETICAL 13.6 KD PROTEIN IN NUP170-ILS1 INTERGENIC REGION [ <i>Saccharomyces cerevisiae</i> ] {IMAGE:889720}	889720 IMAGE Consortium	AA498356
6101 B	9 021LA1G1	ESTs, Weakly similar to The KIAA0151 gene product is classified into serine/threonine kinase. [ <i>H.sapiens</i> ] {IMAGE:875470}	875470 IMAGE Consortium	AA475369
6102 B	11 021LA1G1	IMAGE EST {IMAGE:876446}	876446 IMAGE Consortium	AA475902
6103 D	1 021LA1G1	IMAGE EST {IMAGE:891082}	891082 IMAGE Consortium	AA510126
6104 D	3 021LA1G1	ESTs {IMAGE:902076}	902076 IMAGE Consortium	AA518600
6105 D	5 021LA1G1	ESTs {IMAGE:888952}	888952 IMAGE Consortium	AA466323
6106 D	7 021LA1G1	ESTs, Weakly similar to secretory carrier membrane protein 37 [ <i>R.norvegicus</i> ] {IMAGE:889697}	889697 IMAGE Consortium	AA498328
6107 D	9 021LA1G1	ESTs, Moderately similar to GALECTIN-2 [ <i>Homo sapiens</i> ] {IMAGE:903822}	903822 IMAGE Consortium	AA521545
6108 D	11 021LA1G1	Cytochrome P450, 2e1, ethanol inducible {IMAGE:891122}	891122 IMAGE Consortium	AA510149
6109 F	1 021LA1G1	ESTs {IMAGE:902086}	902086 IMAGE Consortium	AA518605
6110 F	3 021LA1G1	IMAGE EST {IMAGE:903004}	903004 IMAGE Consortium	AA518428
6111 F	5 021LA1G1	Erythrocyte protein band 4.1 {IMAGE:889824}	889824 IMAGE Consortium	AA498495
6112 F	7 021LA1G1	ESTs {IMAGE:889894}	889894 IMAGE Consortium	AA498586
6113 F	9 021LA1G1	ESTs, Weakly similar to retrovirus restriction polypeptide [ <i>M.musculus</i> ] {IMAGE:904137}	904137 IMAGE Consortium	AA518000
6114 F	11 021LA1G1	Calcium channel, voltage-dependent, L type, alpha 2 delta subunit {IMAGE:890932}	890932 IMAGE Consortium	AA511037
6115 H	1 021LA1G1	ESTs {IMAGE:891158}	891158 IMAGE Consortium	AA510252
6116 H	3 021LA1G1	Adaptor-related protein complex AP-1, beta 1 subunit {IMAGE:876929}	876929 IMAGE Consortium	AA497573
6117 H	5 021LA1G1	ESTs, Moderately similar to DARPP-32 {IMAGE:888501}	888501 IMAGE Consortium	AA497615
6118 H	7 021LA1G1	IMAGE EST {IMAGE:890588}	890588 IMAGE Consortium	AA509739

6119 H	9 021LA1G1	IMAGE EST {IMAGE:876341}	876341 IMAGE Consortium	AA476197
6120 H	11 021LA1G1	Mus musculus methyl-CpG binding protein MBD2 (Mbd2) mRNA, complete cds {IMAGE:891284}	891284 IMAGE Consortium	AA510881
6121 B	1 021SA1G2	ESTs {IMAGE:922216}	922216 IMAGE Consortium	AA512638
6122 B	3 021SA1G2	IMAGE EST {IMAGE:922479}	922479 IMAGE Consortium	AA511327
6123 B	5 021SA1G2	Mus musculus G protein signaling regulator RGS5 (rgs5) mRNA, complete cds {IMAGE:935524}	935524 IMAGE Consortium	AA543469
6124 B	7 021SA1G2	Mus musculus protein-tyrosine phosphatase mRNA, complete cds {IMAGE:935557}	935557 IMAGE Consortium	AA543569
6125 B	9 021SA1G2	IMAGE EST {IMAGE:904796}	904796 IMAGE Consortium	AA521940
6126 B	11 021SA1G2	IMAGE EST {IMAGE:948539}	948539 IMAGE Consortium	AA544627
6127 D	1 021SA1G2	ESTs {IMAGE:920701}	920701 IMAGE Consortium	AA538356
6128 D	3 021SA1G2	ESTs, Weakly similar to cytochrome P450 Cyp4a [M.musculus] {IMAGE:922126}	922126 IMAGE Consortium	AA512402
6129 D	5 021SA1G2	ESTs {IMAGE:935343}	935343 IMAGE Consortium	AA542183
6130 D	7 021SA1G2	ESTs {IMAGE:533790}	533790 IMAGE Consortium	AA071762
6131 D	9 021SA1G2	IMAGE EST {IMAGE:920524}	920524 IMAGE Consortium	AA541888
6132 D	11 021SA1G2	IMAGE EST {IMAGE:922971}	922971 IMAGE Consortium	AA511420
6133 F	1 021SA1G2	Mus musculus mRNA for ribosomal protein L35a {IMAGE:659928}	659928 IMAGE Consortium	AA221110
6134 F	3 021SA1G2	Mesenchyme homeobox 2 {IMAGE:672437}	672437 IMAGE Consortium	AA231293
6135 F	5 021SA1G2	M.musculus mRNA for N-glycan alpha 2,8-sialyltransferase {IMAGE:673010}	673010 IMAGE Consortium	AA231471
6136 F	7 021SA1G2	Histocompatibility 2, complement component factor B {IMAGE:676164}	676164 IMAGE Consortium	AA209640
6137 F	9 021SA1G2	ESTs, Moderately similar to PLASMA KALLIKREIN PRECURSOR [M.musculus] {IMAGE:676536}	676536 IMAGE Consortium	AA208793
6138 F	11 021SA1G2	ESTs, Highly similar to plakophilin 2a [H.sapiens] {IMAGE:676792}	676792 IMAGE Consortium	AA212868
6139 H	1 021SA1G2	IMAGE EST {IMAGE:672656}	672656 IMAGE Consortium	AA231457
6140 H	3 021SA1G2	IMAGE EST {IMAGE:660952}	660952 IMAGE Consortium	AA230681
6141 H	5 021SA1G2	ESTs {IMAGE:669838}	669838 IMAGE Consortium	AA230456
6142 H	7 021SA1G2	IMAGE EST {IMAGE:670164}	670164 IMAGE Consortium	AA221472
6143 H	9 021SA1G2	IMAGE EST {IMAGE:670393}	670393 IMAGE Consortium	AA221250
6144 H	11 021SA1G2	ESTs {IMAGE:676967}	676967 IMAGE Consortium	AA213299
6145 B	1 021ZA1G3	ESTs {IMAGE:949044}	949044 IMAGE Consortium	AA543898
6146 B	3 021ZA1G3	IMAGE EST {IMAGE:922298}	922298 IMAGE Consortium	AA512750
6147 B	5 021ZA1G3	Mus musculus PS-2 mRNA, complete cds {IMAGE:949564}	949564 IMAGE Consortium	AA537185
6148 B	7 021ZA1G3	IMAGE EST {IMAGE:904835}	904835 IMAGE Consortium	AA522127
6149 B	9 021ZA1G3	ESTs, Weakly similar to F43E2.7 [C.elegans] {IMAGE:920215}	920215 IMAGE Consortium	AA500298
6150 B	11 021ZA1G3	IMAGE EST {IMAGE:920496}	920496 IMAGE Consortium	AA541835
6151 D	1 021ZA1G3	ESTs {IMAGE:904836}	904836 IMAGE Consortium	AA522117

6152 D	3 021ZA1G3	Fibrous sheath component 1 {IMAGE:516191}	516191 IMAGE Consortium	AA106031
6153 D	5 021ZA1G3	House mouse; Musculus domesticus female mammary carcinoma mRNA for mCDC47, complete cds {IMAGE:948894}	948894 IMAGE Consortium	AA544948
6154 D	7 021ZA1G3	IMAGE EST {IMAGE:949400}	949400 IMAGE Consortium	AA542206
6155 D	9 021ZA1G3	IMAGE EST {IMAGE:949589}	949589 IMAGE Consortium	AA537219
6156 D	11 021ZA1G3	IMAGE EST {IMAGE:935626}	935626 IMAGE Consortium	AA543680
6157 F	1 021ZA1G3	ESTs {IMAGE:904777}	904777 IMAGE Consortium	AA521942
6158 F	3 021ZA1G3	IMAGE EST {IMAGE:552603}	552603 IMAGE Consortium	AA096614
6159 F	5 021ZA1G3	Mouse chromatin nonhistone high mobility group protein (HGM-I(Y), complete cds {IMAGE:920268}	920268 IMAGE Consortium	AA538243
6160 F	7 021ZA1G3	ESTs, Weakly similar to cDNA EST EMBL:M89154 comes from this gene [C.elegans] {IMAGE:948841}	948841 IMAGE Consortium	AA544857
6161 F	9 021ZA1G3	IMAGE EST {IMAGE:920587}	920587 IMAGE Consortium	AA541969
6162 F	11 021ZA1G3	ESTs {IMAGE:922330}	922330 IMAGE Consortium	AA512307
6163 H	1 021ZA1G3	ESTs, Weakly similar to HuEMAP [H.sapiens] {IMAGE:949710}	949710 IMAGE Consortium	AA537426
6164 H	3 021ZA1G3	ESTs, Highly similar to Similar to S.cerevisiae hypothetical protein N0330 [H.sapiens] {IMAGE:904685}	904685 IMAGE Consortium	AA521593
6165 H	5 021ZA1G3	ESTs {IMAGE:904650}	904650 IMAGE Consortium	AA518135
6166 H	7 021ZA1G3	IMAGE EST {IMAGE:920211}	920211 IMAGE Consortium	AA500296
6167 H	9 021ZA1G3	ESTs {IMAGE:948842}	948842 IMAGE Consortium	AA544848
6168 H	11 021ZA1G3	E26 avian leukemia oncogene 2, 3' domain {IMAGE:949055}	949055 IMAGE Consortium	AA543913
6169 B	1 0216A1G4	ESTs {IMAGE:660896}	660896 IMAGE Consortium	AA221135
6170 B	3 0216A1G4	ESTs, Highly similar to SERINE HYDROXYMETHYLTRANSFERASE, MITOCHONDRIAL [Oryctolagus cuniculus] {IMAGE:676311}	676311 IMAGE Consortium	AA208877
6171 B	5 0216A1G4	Apolipoprotein E {IMAGE:676579}	676579 IMAGE Consortium	AA209061
6172 B	7 0216A1G4	ESTs, Weakly similar to cytoplasmic protein Ndr1 [M.musculus] {IMAGE:677163}	677163 IMAGE Consortium	AA213094
6173 B	9 0216A1G4	ESTs, Moderately similar to golgi peripheral membrane protein p65 [R.norvegicus] {IMAGE:677054}	677054 IMAGE Consortium	AA213185
6174 B	11 0216A1G4	Mus musculus flavin-containing monooxygenase 3 mRNA, complete cds {IMAGE:677605}	677605 IMAGE Consortium	AA213017
6175 D	1 0216A1G4	ESTs {IMAGE:676911}	676911 IMAGE Consortium	AA213062
6176 D	3 0216A1G4	ESTs, Highly similar to CALCINEURIN B SUBUNIT ISOFORM 1 [Homo sapiens; Bos taurus; Rattus norvegicus] {IMAGE:677488}	677488 IMAGE Consortium	AA213263
6177 D	5 0216A1G4	IMAGE EST {IMAGE:659533}	659533 IMAGE Consortium	AA218234
6178 D	7 0216A1G4	IMAGE EST {IMAGE:659768}	659768 IMAGE Consortium	AA220047
6179 D	9 0216A1G4	Mus musculus amyloid beta-peptide binding protein (ERAB) mRNA, complete cds {IMAGE:660171}	660171 IMAGE Consortium	AA220432
6180 D	11 0216A1G4	ESTs {IMAGE:660601}	660601 IMAGE Consortium	AA222865
6181 F	1 0216A1G4	IMAGE EST {IMAGE:677402}	677402 IMAGE Consortium	AA212861

6182 F	3 0216A1G4	ESTs, Moderately similar to MAK16 PROTEIN [Saccharomyces cerevisiae] {IMAGE:670893}	670893 IMAGE Consortium	AA250407
6183 F	5 0216A1G4	Small nuclear ribonucleoprotein B {IMAGE:671171}	671171 IMAGE Consortium	AA222348
6184 F	7 0216A1G4	ESTs, Highly similar to TNF-alpha stimulated ABC protein [H.sapiens] {IMAGE:659557}	659557 IMAGE Consortium	AA218241
6185 F	9 0216A1G4	IMAGE EST {IMAGE:659745}	659745 IMAGE Consortium	AA220044
6186 F	11 0216A1G4	GATA-binding protein 4 {IMAGE:672424}	672424 IMAGE Consortium	AA230989
6187 H	1 0216A1G4	Tubulin, beta 4 {IMAGE:671340}	671340 IMAGE Consortium	AA222216
6188 H	3 0216A1G4	ESTs {IMAGE:672024}	672024 IMAGE Consortium	AA245119
6189 H	5 0216A1G4	IMAGE EST {IMAGE:672189}	672189 IMAGE Consortium	AA231161
6190 H	7 0216A1G4	ESTs {IMAGE:660716}	660716 IMAGE Consortium	AA230553
6191 H	9 0216A1G4	ESTs, Weakly similar to similar to gamma-butyrobetaine,2-oxoglutarate dioxygenase [C.elegans] {IMAGE:676349}	676349 IMAGE Consortium	AA208820
6192 H	11 0216A1G4	ESTs {IMAGE:676123}	676123 IMAGE Consortium	AA209095
6193 B	1 021DA1G5	ESTs {IMAGE:672859}	672859 IMAGE Consortium	AA231422
6194 B	3 021DA1G5	ESTs, Highly similar to CYSTEINE-RICH PROTEIN 2 [Rattus norvegicus] {IMAGE:676134}	676134 IMAGE Consortium	AA208934
6195 B	5 021DA1G5	ESTs {IMAGE:670111}	670111 IMAGE Consortium	AA221321
6196 B	7 021DA1G5	Mus musculus DNA cytosine-5 methyltransferase 3A (Dnmt3a) mRNA, complete cds {IMAGE:670948}	670948 IMAGE Consortium	AA240794
6197 B	9 021DA1G5	ESTs, Highly similar to HEAVY METAL TOLERANCE PROTEIN PRECURSOR [Schizosaccharomyces pombe] {IMAGE:677418}	677418 IMAGE Consortium	AA212806
6198 B	11 021DA1G5	IMAGE EST {IMAGE:658905}	658905 IMAGE Consortium	AA217908
6199 D	1 021DA1G5	Growth factor receptor bound protein 2 {IMAGE:670112}	670112 IMAGE Consortium	AA231029
6200 D	3 021DA1G5	ESTs {IMAGE:670245}	670245 IMAGE Consortium	AA221231
6201 D	5 021DA1G5	IMAGE EST {IMAGE:677012}	677012 IMAGE Consortium	AA213280
6202 D	7 021DA1G5	M.musculus rhoC mRNA {IMAGE:670886}	670886 IMAGE Consortium	AA221997
6203 D	9 021DA1G5	ESTs {IMAGE:671087}	671087 IMAGE Consortium	AA240009
6204 D	11 021DA1G5	Delta-aminolevulinate dehydratase {IMAGE:671205}	671205 IMAGE Consortium	AA222320
6205 F	1 021DA1G5	TRANSPLANTATION ANTIGEN P35B {IMAGE:677601}	677601 IMAGE Consortium	AA213015
6206 F	3 021DA1G5	IMAGE EST {IMAGE:671280}	671280 IMAGE Consortium	AA222481
6207 F	5 021DA1G5	IMAGE EST {IMAGE:659374}	659374 IMAGE Consortium	AA218405
6208 F	7 021DA1G5	ESTs {IMAGE:672198}	672198 IMAGE Consortium	AA230984
6209 F	9 021DA1G5	ESTs {IMAGE:676152}	676152 IMAGE Consortium	AA208998
6210 F	11 021DA1G5	ESTs {IMAGE:669765}	669765 IMAGE Consortium	AA230444
6211 H	1 021DA1G5	ESTs {IMAGE:672291}	672291 IMAGE Consortium	AA231535
6212 H	3 021DA1G5	ESTs, Moderately similar to COMPLEMENT C1S COMPONENT PRECURSOR [H.sapiens] {IMAGE:676176}	676176 IMAGE Consortium	AA209006
6213 H	5 021DA1G5	Huntington disease gene homolog {IMAGE:669939}	669939 IMAGE Consortium	AA250402
6214 H	7 021DA1G5	ESTs {IMAGE:676898}	676898 IMAGE Consortium	AA212963
6215 H	9 021DA1G5	Protease (prosome, macropain) 28 subunit, alpha {IMAGE:678768}	678768 IMAGE Consortium	AA239485



6216 H	11 021DA1G5	ESTs, Highly similar to PS-PLA1 [R.norvegicus] {IMAGE:679080}	679080 IMAGE Consortium	AA244721
6217 B	1 021KA1G6	Matrix metalloproteinase 15 {IMAGE:680964}	680964 IMAGE Consortium	AA242457
6218 B	3 021KA1G6	ESTs {IMAGE:678608}	678608 IMAGE Consortium	AA238106
6219 B	5 021KA1G6	ESTs {IMAGE:679086}	679086 IMAGE Consortium	AA244724
6220 B	7 021KA1G6	Apolipoprotein CI {IMAGE:681599}	681599 IMAGE Consortium	AA237977
6221 B	9 021KA1G6	ESTs {IMAGE:692552}	692552 IMAGE Consortium	AA250720
6222 B	11 021KA1G6	Mus musculus cytochrome P450 monooxygenase (CYP2J5) mRNA, complete cds {IMAGE:692850}	692850 IMAGE Consortium	AA238492
6223 D	1 021KA1G6	ESTs, Moderately similar to MALONYL-COA DECARBOXYLASE PRECURSOR [Anser anser anser] {IMAGE:692290}	692290 IMAGE Consortium	AA238355
6224 D	3 021KA1G6	ESTs {IMAGE:679706}	679706 IMAGE Consortium	AA245131
6225 D	5 021KA1G6	IMAGE EST {IMAGE:693148}	693148 IMAGE Consortium	AA253678
6226 D	7 021KA1G6	M.musculus mRNA for e1 protein {IMAGE:693257}	693257 IMAGE Consortium	AA397020
6227 D	9 021KA1G6	Mus musculus Ah receptor-interacting protein (AIP) mRNA, complete cds {IMAGE:693415}	693415 IMAGE Consortium	AA245324
6228 D	11 021KA1G6	ESTs, Weakly similar to PUTATIVE ACID PHOSPHATASE F26C11.1 [Caenorhabditis elegans] {IMAGE:693719}	693719 IMAGE Consortium	AA265095
6229 F	1 021KA1G6	ESTs {IMAGE:679563}	679563 IMAGE Consortium	AA242669
6230 F	3 021KA1G6	ESTs {IMAGE:692827}	692827 IMAGE Consortium	AA238488
6231 F	5 021KA1G6	ESTs, Highly similar to a-glucosidase I [H.sapiens] {IMAGE:680150}	680150 IMAGE Consortium	AA237872
6232 F	7 021KA1G6	ESTs {IMAGE:680063}	680063 IMAGE Consortium	AA244796
6233 F	9 021KA1G6	ESTs {IMAGE:680407}	680407 IMAGE Consortium	AA250595
6234 F	11 021KA1G6	M.musculus mRNA for interleukin 1 receptor accessory protein {IMAGE:694135}	694135 IMAGE Consortium	AA239010
6235 H	1 021KA1G6	ESTs {IMAGE:693560}	693560 IMAGE Consortium	AA239254
6236 H	3 021KA1G6	ESTs {IMAGE:693938}	693938 IMAGE Consortium	AA238831
6237 H	5 021KA1G6	ESTs, Highly similar to GLUCOSE 1-DEHYDROGENASE [Bacillus subtilis] {IMAGE:680935}	680935 IMAGE Consortium	AA241896
6238 H	7 021KA1G6	IMAGE EST {IMAGE:681143}	681143 IMAGE Consortium	AA242117
6239 H	9 021KA1G6	Glutamyl aminopeptidase {IMAGE:681802}	681802 IMAGE Consortium	AA244590
6240 H	11 021KA1G6	Prolactin receptor {IMAGE:679305}	679305 IMAGE Consortium	AA244758
6241 B	1 021RA1G7	IMAGE EST {IMAGE:693542}	693542 IMAGE Consortium	AA239252
6242 B	3 021RA1G7	ESTs {IMAGE:694035}	694035 IMAGE Consortium	AA238521
6243 B	5 021RA1G7	Mus musculus betaine homocysteine methyl transferase (BHMT) mRNA, complete cds {IMAGE:680854}	680854 IMAGE Consortium	AA241859
6244 B	7 021RA1G7	ESTs {IMAGE:678748}	678748 IMAGE Consortium	AA239479
6245 B	9 021RA1G7	Mus musculus pituitary tumor transforming gene protein (PTTG) mRNA, complete cds {IMAGE:678793}	678793 IMAGE Consortium	AA250500
6246 B	11 021RA1G7	Growth factor receptor bound protein 7 {IMAGE:679007}	679007 IMAGE Consortium	AA244622
6247 D	1 021RA1G7	ESTs {IMAGE:679464}	679464 IMAGE Consortium	AA244557
6248 D	3 021RA1G7	IMAGE EST {IMAGE:679594}	679594 IMAGE Consortium	AA242617

6249 D	5 021RA1G7	Mus musculus cell death activator CIDE-B (Cide-b) mRNA, complete cds {IMAGE:679895}	679895 IMAGE Consortium	AA238193
6250 D	7 021RA1G7	Mouse gamma-aminobutyric acid transporter protein (GABA transporter) mRNA {IMAGE:693366}	693366 IMAGE Consortium	AA397130
6251 D	9 021RA1G7	ESTs {IMAGE:680450}	680450 IMAGE Consortium	AA237174
6252 D	11 021RA1G7	Aldehyde dehydrogenase 4, liver microsomal (class 3) {IMAGE:694108}	694108 IMAGE Consortium	AA238875
6253 F	1 021RA1G7	Mus musculus Masp-2 mRNA for mannose-binding lectin associated serine protease-2, complete cds {IMAGE:680232}	680232 IMAGE Consortium	AA238260
6254 F	3 021RA1G7	CCAAT/enhancer binding protein alpha (C/EBP), related sequence 1 {IMAGE:680123}	680123 IMAGE Consortium	AA244944
6255 F	5 021RA1G7	ESTs {IMAGE:680225}	680225 IMAGE Consortium	AA237611
6256 F	7 021RA1G7	D-E-A-D (aspartate-glutamate-alanine-aspartate) box polypeptide 5 {IMAGE:680475}	680475 IMAGE Consortium	AA237184
6257 F	9 021RA1G7	ESTs {IMAGE:678415}	678415 IMAGE Consortium	AA237424
6258 F	11 021RA1G7	ESTs {IMAGE:678854}	678854 IMAGE Consortium	AA250462
6259 H	1 021RA1G7	ESTs, Weakly similar to 36k hydrophobic nucleolar protein [M.musculus] {IMAGE:680512}	680512 IMAGE Consortium	AA237136
6260 H	3 021RA1G7	ESTs {IMAGE:680761}	680761 IMAGE Consortium	AA242411
6261 H	5 021RA1G7	IMAGE EST {IMAGE:680916}	680916 IMAGE Consortium	AA242445
6262 H	7 021RA1G7	Histocompatibility 2, D region locus 1 {IMAGE:694651}	694651 IMAGE Consortium	AA221044
6263 H	9 021RA1G7	IMAGE EST {IMAGE:681424}	681424 IMAGE Consortium	AA237757
6264 H	11 021RA1G7	IMAGE EST {IMAGE:681542}	681542 IMAGE Consortium	AA237540
6265 B	1 021YA1G8	ESTs, Highly similar to MELANOMA-ASSOCIATED ANTIGEN XP [Homo sapiens] {IMAGE:681587}	681587 IMAGE Consortium	AA237820
6266 B	3 021YA1G8	IMAGE EST {IMAGE:679148}	679148 IMAGE Consortium	AA245738
6267 B	5 021YA1G8	ESTs, Highly similar to CYSTATHIONINE GAMMA-LYASE [Homo sapiens] {IMAGE:679235}	679235 IMAGE Consortium	AA245993
6268 B	7 021YA1G8	IMAGE EST {IMAGE:679341}	679341 IMAGE Consortium	AA245672
6269 B	9 021YA1G8	ESTs, Moderately similar to Similar to a C.elegans protein in cosmid C14H10 [H.sapiens] {IMAGE:679641}	679641 IMAGE Consortium	AA242192
6270 B	11 021YA1G8	ESTs {IMAGE:679938}	679938 IMAGE Consortium	AA238201
6271 D	1 021YA1G8	IMAGE EST {IMAGE:717798}	717798 IMAGE Consortium	AA250469
6272 D	3 021YA1G8	Nidogen {IMAGE:697887}	697887 IMAGE Consortium	AA238652
6273 D	5 021YA1G8	Mus musculus mRNA for mouse rabaptin-5, complete cds {IMAGE:718521}	718521 IMAGE Consortium	AA266332
6274 D	7 021YA1G8	IMAGE EST {IMAGE:695269}	695269 IMAGE Consortium	AA220487
6275 D	9 021YA1G8	ESTs {IMAGE:695589}	695589 IMAGE Consortium	AA220551
6276 D	11 021YA1G8	ESTs {IMAGE:716725}	716725 IMAGE Consortium	AA266146
6277 F	1 021YA1G8	IMAGE EST {IMAGE:718665}	718665 IMAGE Consortium	AA265636
6278 F	3 021YA1G8	Integrin alpha 7 {IMAGE:695845}	695845 IMAGE Consortium	AA222315
6279 F	5 021YA1G8	Cytochrome P450, 2e1, ethanol inducible {IMAGE:696374}	696374 IMAGE Consortium	AA239467
6280 F	7 021YA1G8	Heart and neural crest derivatives expressed transcript 1 {IMAGE:696795}	696795 IMAGE Consortium	AA240305

6281 F	9 021YA1G8	ESTs {IMAGE:717191}	717191 IMAGE Consortium	AA266478
6282 F	11 021YA1G8	PCTAIRE-motif protein kinase 1 {IMAGE:697169}	697169 IMAGE Consortium	AA230417
6283 H	1 021YA1G8	IMAGE EST {IMAGE:695447}	695447 IMAGE Consortium	AA220498
6284 H	3 021YA1G8	IMAGE EST {IMAGE:716750}	716750 IMAGE Consortium	AA266151
6285 H	5 021YA1G8	Mus musculus FGF-binding protein (FGF-BP) mRNA, complete cds {IMAGE:717457}	717457 IMAGE Consortium	AA403432
6286 H	7 021YA1G8	ESTs {IMAGE:717299}	717299 IMAGE Consortium	AA403861
6287 H	9 021YA1G8	ESTs {IMAGE:717816}	717816 IMAGE Consortium	AA250473
6288 H	11 021YA1G8	ESTs {IMAGE:717948}	717948 IMAGE Consortium	AA249917
6289 B	1 0215A1G9	ESTs {IMAGE:696728}	696728 IMAGE Consortium	AA240276
6290 B	3 0215A1G9	ESTs {IMAGE:717709}	717709 IMAGE Consortium	AA250450
6291 B	5 0215A1G9	IMAGE EST {IMAGE:697511}	697511 IMAGE Consortium	AA253924
6292 B	7 0215A1G9	ESTs, Weakly similar to cDNA EST yk486b9.3 comes from this gene [C.elegans] {IMAGE:697835}	697835 IMAGE Consortium	AA238637
6293 B	9 0215A1G9	IMAGE EST {IMAGE:698242}	698242 IMAGE Consortium	AA239129
6294 B	11 0215A1G9	ESTs {IMAGE:699462}	699462 IMAGE Consortium	AA245526
6295 D	1 0215A1G9	ESTs {IMAGE:717910}	717910 IMAGE Consortium	AA250561
6296 D	3 0215A1G9	ESTs {IMAGE:718193}	718193 IMAGE Consortium	AA266355
6297 D	5 0215A1G9	IMAGE EST {IMAGE:718495}	718495 IMAGE Consortium	AA266294
6298 D	7 0215A1G9	IMAGE EST {IMAGE:718666}	718666 IMAGE Consortium	AA265560
6299 D	9 0215A1G9	IMAGE EST {IMAGE:695231}	695231 IMAGE Consortium	AA220507
6300 D	11 0215A1G9	Mus musculus mRNA for sarco/endoplasmic reticulum Ca <sup>2+</sup> - ATPase (SERCA2) {IMAGE:695695}	695695 IMAGE Consortium	AA222567
6301 F	1 0215A1G9	IMAGE EST {IMAGE:698221}	698221 IMAGE Consortium	AA239126
6302 F	3 0215A1G9	ESTs {IMAGE:699290}	699290 IMAGE Consortium	AA245395
6303 F	5 0215A1G9	ESTs, Moderately similar to predicted using Genefinder [C.elegans] {IMAGE:699416}	699416 IMAGE Consortium	AA245448
6304 F	7 0215A1G9	ESTs {IMAGE:699377}	699377 IMAGE Consortium	AA245492
6305 F	9 0215A1G9	Rad51 homolog (S. cerevisiae) {IMAGE:717102}	717102 IMAGE Consortium	AA266114
6306 F	11 0215A1G9	Mus musculus eIF4E-like protein 4E-LP mRNA, complete cds {IMAGE:717088}	717088 IMAGE Consortium	AA266082
6307 H	1 0215A1G9	Cytochrome P450, 2f2 {IMAGE:695667}	695667 IMAGE Consortium	AA220582
6308 H	3 0215A1G9	ESTs, Highly similar to DNA-binding protein [H.sapiens] {IMAGE:716588}	716588 IMAGE Consortium	AA265312
6309 H	5 0215A1G9	ESTs {IMAGE:716815}	716815 IMAGE Consortium	AA266387
6310 H	7 0215A1G9	Carbon catabolite repression 4 homolog (S. cerevisiae) {IMAGE:717197}	717197 IMAGE Consortium	AA404014
6311 H	9 0215A1G9	Lectin, galactose binding, soluble 6 {IMAGE:717208}	717208 IMAGE Consortium	AA404174
6312 H	11 0215A1G9	ESTs {IMAGE:717819}	717819 IMAGE Consortium	AA250522
6313 B	1 021PA1GA	Myeloblastosis oncogene {IMAGE:721803}	721803 IMAGE Consortium	AA267899
6314 B	3 021PA1GA	IMAGE EST {IMAGE:721936}	721936 IMAGE Consortium	AA267969
6315 B	5 021PA1GA	ESTs, Highly similar to The ha0919 gene product is novel. [H.sapiens] {IMAGE:719597}	719597 IMAGE Consortium	AA255088

6316 B	7	021PA1GA	IMAGE EST {IMAGE:722318}	722318	IMAGE Consortium	AA254095
6317 B	9	021PA1GA	IMAGE EST {IMAGE:722761}	722761	IMAGE Consortium	AA254547
6318 B	11	021PA1GA	ESTs {IMAGE:720349}	720349	IMAGE Consortium	AA270888
6319 D	1	021PA1GA	ESTs {IMAGE:721889}	721889	IMAGE Consortium	AA268522
6320 D	3	021PA1GA	ESTs {IMAGE:722402}	722402	IMAGE Consortium	AA254182
6321 D	5	021PA1GA	Zinc finger protein 28 {IMAGE:722409}	722409	IMAGE Consortium	AA254285
6322 D	7	021PA1GA	IMAGE EST {IMAGE:722910}	722910	IMAGE Consortium	AA254649
6323 D	9	021PA1GA	ESTs {IMAGE:720937}	720937	IMAGE Consortium	AA267578
6324 D	11	021PA1GA	ESTs {IMAGE:720707}	720707	IMAGE Consortium	AA267512
6325 F	1	021PA1GA	ESTs {IMAGE:719264}	719264	IMAGE Consortium	AA254744
6326 F	3	021PA1GA	NEDD1 PROTEIN {IMAGE:722426}	722426	IMAGE Consortium	AA254225
6327 F	5	021PA1GA	ESTs {IMAGE:722433}	722433	IMAGE Consortium	AA254293
6328 F	7	021PA1GA	M.musculus mRNA for mTGIF protein {IMAGE:722623}	722623	IMAGE Consortium	AA260654
6329 F	9	021PA1GA	Pyruvate decarboxylase {IMAGE:722754}	722754	IMAGE Consortium	AA260717
6330 F	11	021PA1GA	IMAGE EST {IMAGE:723156}	723156	IMAGE Consortium	AA254480
6331 H	1	021PA1GA	Mus musculus COP9 complex subunit 7b (COPS7b) mRNA, complete cds {IMAGE:719496}	719496	IMAGE Consortium	AA254769
6332 H	3	021PA1GA	ESTs {IMAGE:722297}	722297	IMAGE Consortium	AA254212
6333 H	5	021PA1GA	M.musculus mRNA for alpha-2 antiplasmin {IMAGE:720340}	720340	IMAGE Consortium	AA270852
6334 H	7	021PA1GA	ESTs, Highly similar to APOLIPOPROTEIN B-100 PRECURSOR [Homo sapiens] {IMAGE:723038}	723038	IMAGE Consortium	AA254389
6335 H	9	021PA1GA	Mus musculus 76 kDa tyrosine phosphoprotein SLP-76 mRNA, complete cds {IMAGE:723285}	723285	IMAGE Consortium	AA267811
6336 H	11	021PA1GA	ESTs, Moderately similar to KINESIN LIGHT CHAIN [Rattus norvegicus] {IMAGE:733346}	733346	IMAGE Consortium	AA261327
6337 B	1	021WA1GB	IMAGE EST {IMAGE:717342}	717342	IMAGE Consortium	AA403748
6338 B	3	021WA1GB	IMAGE EST {IMAGE:717628}	717628	IMAGE Consortium	AA250329
6339 B	5	021WA1GB	ESTs, Weakly similar to KELL BLOOD GROUP GLYCOPROTEIN [H.sapiens] {IMAGE:718248}	718248	IMAGE Consortium	AA261590
6340 B	7	021WA1GB	ESTs, Weakly similar to zinc finger protein [M.musculus] {IMAGE:718195}	718195	IMAGE Consortium	AA266356
6341 B	9	021WA1GB	Mus musculus TDAG51 (TDAG51) mRNA, complete cds {IMAGE:698297}	698297	IMAGE Consortium	AA239282
6342 B	11	021WA1GB	IMAGE EST {IMAGE:699411}	699411	IMAGE Consortium	AA245350
6343 D	1	021WA1GB	IMAGE EST {IMAGE:695723}	695723	IMAGE Consortium	AA222181
6344 D	3	021WA1GB	B-cell leukemia/lymphoma 3 {IMAGE:716976}	716976	IMAGE Consortium	AA266002
6345 D	5	021WA1GB	ESTs {IMAGE:716877}	716877	IMAGE Consortium	AA266195
6346 D	7	021WA1GB	ESTs, Moderately similar to Phosphatidic acid phosphatase [M.musculus] {IMAGE:697092}	697092	IMAGE Consortium	AA220316
6347 D	9	021WA1GB	ESTs {IMAGE:697506}	697506	IMAGE Consortium	AA253913
6348 D	11	021WA1GB	ESTs, Moderately similar to ZINC FINGER PROTEIN ZFP-36 [Homo sapiens] {IMAGE:718565}	718565	IMAGE Consortium	AA266614
6349 F	1	021WA1GB	ESTs {IMAGE:722406}	722406	IMAGE Consortium	AA254184

6350 F	3	021WA1GB	IMAGE EST {IMAGE:719965}	719965	IMAGE Consortium	AA255150
6351 F	5	021WA1GB	IMAGE EST {IMAGE:720408}	720408	IMAGE Consortium	AA261436
6352 F	7	021WA1GB	House mouse; Musculus domesticus male brain mRNA for ARF1, complete cds {IMAGE:720444}	720444	IMAGE Consortium	AA261520
6353 F	9	021WA1GB	ESTs, Moderately similar to [Human chromosome 3p21.1 gene sequence, complete cds.], gene product [H.sapiens] {IMAGE:720573}	720573	IMAGE Consortium	AA261422
6354 F	11	021WA1GB	IMAGE EST {IMAGE:723600}	723600	IMAGE Consortium	AA260793
6355 H	1	021WA1GB	ESTs {IMAGE:719167}	719167	IMAGE Consortium	AA267315
6356 H	3	021WA1GB	ESTs {IMAGE:722039}	722039	IMAGE Consortium	AA268483
6357 H	5	021WA1GB	IMAGE EST {IMAGE:722336}	722336	IMAGE Consortium	AA254100
6358 H	7	021WA1GB	ESTs {IMAGE:719894}	719894	IMAGE Consortium	AA255141
6359 H	9	021WA1GB	IMAGE EST {IMAGE:720457}	720457	IMAGE Consortium	AA261076
6360 H	11	021WA1GB	IMAGE EST {IMAGE:720454}	720454	IMAGE Consortium	AA261662
6361 B	1	0213A1GC	ESTs {IMAGE:722154}	722154	IMAGE Consortium	AA268414
6362 B	3	0213A1GC	IMAGE EST {IMAGE:722451}	722451	IMAGE Consortium	AA254245
6363 B	5	0213A1GC	Adrenomedullin receptor {IMAGE:720232}	720232	IMAGE Consortium	AA255046
6364 B	7	0213A1GC	ESTs, Weakly similar to Ylr323cp [S.cerevisiae] {IMAGE:723182}	723182	IMAGE Consortium	AA254511
6365 B	9	0213A1GC	ESTs {IMAGE:723118}	723118	IMAGE Consortium	AA254455
6366 B	11	0213A1GC	NEURONAL PROTEIN 3.1 {IMAGE:733420}	733420	IMAGE Consortium	AA259807
6367 D	1	0213A1GC	ESTs {IMAGE:722506}	722506	IMAGE Consortium	AA254055
6368 D	3	0213A1GC	IMAGE EST {IMAGE:720114}	720114	IMAGE Consortium	AA255317
6369 D	5	0213A1GC	ESTs, Moderately similar to MEA6 [H.sapiens] {IMAGE:720342}	720342	IMAGE Consortium	AA270853
6370 D	7	0213A1GC	Fibroblast growth factor 1 {IMAGE:720400}	720400	IMAGE Consortium	AA261582
6371 D	9	0213A1GC	IMAGE EST {IMAGE:720973}	720973	IMAGE Consortium	AA266744
6372 D	11	0213A1GC	ESTs, Highly similar to RIBOSOMAL PROTEIN S6 KINASE [Homo sapiens] {IMAGE:720977}	720977	IMAGE Consortium	AA266868
6373 F	1	0213A1GC	ESTs {IMAGE:722435}	722435	IMAGE Consortium	AA254294
6374 F	3	0213A1GC	ESTs {IMAGE:722418}	722418	IMAGE Consortium	AA253969
6375 F	5	0213A1GC	ESTs {IMAGE:720558}	720558	IMAGE Consortium	AA261697
6376 F	7	0213A1GC	ESTs {IMAGE:720640}	720640	IMAGE Consortium	AA261454
6377 F	9	0213A1GC	IMAGE EST {IMAGE:723166}	723166	IMAGE Consortium	AA254483
6378 F	11	0213A1GC	Cytochrome P450, 2e1, ethanol inducible {IMAGE:721382}	721382	IMAGE Consortium	AA267254
6379 H	1	0213A1GC	ESTs, Highly similar to PROTEASOME COMPONENT C2 [Homo sapiens] {IMAGE:723267}	723267	IMAGE Consortium	AA267785
6380 H	3	0213A1GC	Huntingtin-associated protein 1 {IMAGE:723088}	723088	IMAGE Consortium	AA254430
6381 H	5	0213A1GC	IMAGE EST {IMAGE:723190}	723190	IMAGE Consortium	AA254513
6382 H	7	0213A1GC	ESTs {IMAGE:721167}	721167	IMAGE Consortium	AA267233
6383 H	9	0213A1GC	IMAGE EST {IMAGE:733846}	733846	IMAGE Consortium	AA268263
6384 H	11	0213A1GC	IMAGE EST {IMAGE:721461}	721461	IMAGE Consortium	AA266935
6385 B	1	021AA1GD	Mus musculus gamma-glutamyl hydrolase precursor, mRNA,	733456	IMAGE Consortium	AA259661

		nuclear gene encoding lysosomal protein, complete cds {IMAGE:733456}		
6386 B	3 021AA1GD	IMAGE EST {IMAGE:733899}	733899 IMAGE Consortium	AA269328
6387 B	5 021AA1GD	ESTs, Highly similar to LZTR-1 [H.sapiens] {IMAGE:733862}	733862 IMAGE Consortium	AA268165
6388 B	7 021AA1GD	Mus musculus nimA-related kinase 2 (Nek2) mRNA, complete cds {IMAGE:733966}	733966 IMAGE Consortium	AA268349
6389 B	9 021AA1GD	ESTs {IMAGE:718768}	718768 IMAGE Consortium	AA265700
6390 B	11 021AA1GD	ESTs {IMAGE:721846}	721846 IMAGE Consortium	AA267717
6391 D	1 021AA1GD	ESTs {IMAGE:721241}	721241 IMAGE Consortium	AA267202
6392 D	3 021AA1GD	IMAGE EST {IMAGE:721547}	721547 IMAGE Consortium	AA267442
6393 D	5 021AA1GD	Mouse mRNA for squalene epoxidase, complete cds {IMAGE:734170}	734170 IMAGE Consortium	AA268608
6394 D	7 021AA1GD	ESTs {IMAGE:737688}	737688 IMAGE Consortium	AA277652
6395 D	9 021AA1GD	ESTs {IMAGE:737605}	737605 IMAGE Consortium	AA277142
6396 D	11 021AA1GD	ESTs, Weakly similar to QUINONE OXIDOREDUCTASE [M.musculus] {IMAGE:735186}	735186 IMAGE Consortium	AA259674
6397 F	1 021AA1GD	ESTs, Highly similar to PERIPHERAL-TYPE BENZODIAZEPINE RECEPTOR [Rattus norvegicus] {IMAGE:736628}	736628 IMAGE Consortium	AA270584
6398 F	3 021AA1GD	Mus musculus retinoid X receptor interacting protein (RIP14- 1No.6) mRNA, complete cds {IMAGE:747875}	747875 IMAGE Consortium	AA261313
6399 F	5 021AA1GD	ESTs, Weakly similar to Trio [H.sapiens] {IMAGE:734314}	734314 IMAGE Consortium	AA268669
6400 F	7 021AA1GD	ESTs {IMAGE:734808}	734808 IMAGE Consortium	AA259365
6401 F	9 021AA1GD	ESTs {IMAGE:737629}	737629 IMAGE Consortium	AA277149
6402 F	11 021AA1GD	IMAGE EST {IMAGE:737782}	737782 IMAGE Consortium	AA277831
6403 H	1 021AA1GD	ESTs, Weakly similar to ORF YOR281c [S.cerevisiae] {IMAGE:747612}	747612 IMAGE Consortium	AA268866
6404 H	3 021AA1GD	ESTs {IMAGE:737442}	737442 IMAGE Consortium	AA277843
6405 H	5 021AA1GD	ESTs, Highly similar to Shb protein [H.sapiens] {IMAGE:737572}	737572 IMAGE Consortium	AA277474
6406 H	7 021AA1GD	ESTs, Highly similar to HYPOTHETICAL 272.0 KD PROTEIN C50C3.6 IN CHROMOSOME III [Caenorhabditis elegans] {IMAGE:737750}	737750 IMAGE Consortium	AA277696
6407 H	9 021AA1GD	IMAGE EST {IMAGE:737954}	737954 IMAGE Consortium	AA271006
6408 H	11 021AA1GD	Mus musculus pantophysin gene, complete cds {IMAGE:737944}	737944 IMAGE Consortium	AA271505
6409 B	1 021HA1GE	ESTs, Highly similar to NADH-UBIQUINONE OXIDOREDUCTASE PDSW SUBUNIT [Bos taurus] {IMAGE:746481}	746481 IMAGE Consortium	AA260194
6410 B	3 021HA1GE	ESTs {IMAGE:746908}	746908 IMAGE Consortium	AA274470
6411 B	5 021HA1GE	IMAGE EST {IMAGE:746798}	746798 IMAGE Consortium	AA260428
6412 B	7 021HA1GE	ESTs, Weakly similar to Schizosaccharomyces pombe 4-nitrophenylphosphatase [C.elegans] {IMAGE:747384}	747384 IMAGE Consortium	AA268225

6413 B	9	021HA1GE	ESTs {IMAGE:747350}	747350 IMAGE Consortium	AA268133
6414 B	11	021HA1GE	Histocompatibility 2, class II antigen A, alpha {IMAGE:747378}	747378 IMAGE Consortium	AA272807
6415 D	1	021HA1GE	ESTs {IMAGE:735445}	735445 IMAGE Consortium	AA269921
6416 D	3	021HA1GE	IMAGE EST {IMAGE:735524}	735524 IMAGE Consortium	AA269841
6417 D	5	021HA1GE	ESTs {IMAGE:735850}	735850 IMAGE Consortium	AA269594
6418 D	7	021HA1GE	Mus musculus mRNA for F1F0-ATP synthase, g subunit {IMAGE:735887}	735887 IMAGE Consortium	AA269701
6419 D	9	021HA1GE	ESTs, Highly similar to GLYCOGEN SYNTHASE KINASE-3 BETA [R.norvegicus] {IMAGE:736232}	736232 IMAGE Consortium	AA270371
6420 D	11	021HA1GE	IMAGE EST {IMAGE:736193}	736193 IMAGE Consortium	AA269950
6421 F	1	021HA1GE	IMAGE EST {IMAGE:735089}	735089 IMAGE Consortium	AA259720
6422 F	3	021HA1GE	IMAGE EST {IMAGE:738140}	738140 IMAGE Consortium	AA271043
6423 F	5	021HA1GE	ESTs, Moderately similar to PL6 protein [H.sapiens] {IMAGE:746547}	746547 IMAGE Consortium	AA260135
6424 F	7	021HA1GE	ESTs {IMAGE:735901}	735901 IMAGE Consortium	AA269613
6425 F	9	021HA1GE	IMAGE EST {IMAGE:736200}	736200 IMAGE Consortium	AA270366
6426 F	11	021HA1GE	ESTs, Highly similar to DEOXYHYPUSINE SYNTHASE [H.sapiens] {IMAGE:736328}	736328 IMAGE Consortium	AA270493
6427 H	1	021HA1GE	IMAGE EST {IMAGE:746643}	746643 IMAGE Consortium	AA268085
6428 H	3	021HA1GE	IMAGE EST {IMAGE:735662}	735662 IMAGE Consortium	AA269798
6429 H	5	021HA1GE	ESTs, Highly similar to HYPOTHETICAL 29.2 KD PROTEIN IN PHD1-PTM1 INTERGENIC REGION [Saccharomyces cerevisiae] {IMAGE:736297}	736297 IMAGE Consortium	AA270607
6430 H	7	021HA1GE	Mus musculus MPS1 gene and mRNA, 3'end {IMAGE:747364}	747364 IMAGE Consortium	AA268219
6431 H	9	021HA1GE	Laminin, alpha 4 {IMAGE:736616}	736616 IMAGE Consortium	AA270669
6432 H	11	021HA1GE	ESTs {IMAGE:747624}	747624 IMAGE Consortium	AA268758
6433 B	1	021OA1GF	ESTs {IMAGE:734810}	734810 IMAGE Consortium	AA259366
6434 B	3	021OA1GF	ESTs, Weakly similar to PROTHROMBINASE PRECURSOR [M.musculus] {IMAGE:734759}	734759 IMAGE Consortium	AA261769
6435 B	5	021OA1GF	IMAGE EST {IMAGE:737931}	737931 IMAGE Consortium	AA270993
6436 B	7	021OA1GF	Mus musculus coagulation factor VII (fVII) mRNA, complete cds {IMAGE:738136}	738136 IMAGE Consortium	AA271041
6437 B	9	021OA1GF	Thymoma viral proto-oncogene 2 {IMAGE:735648}	735648 IMAGE Consortium	AA269678
6438 B	11	021OA1GF	Gap junction membrane channel protein alpha 4 {IMAGE:746939}	746939 IMAGE Consortium	AA274472
6439 D	1	021OA1GF	IMAGE EST {IMAGE:734718}	734718 IMAGE Consortium	AA259304
6440 D	3	021OA1GF	IMAGE EST {IMAGE:737872}	737872 IMAGE Consortium	AA277240
6441 D	5	021OA1GF	ESTs {IMAGE:735141}	735141 IMAGE Consortium	AA259775
6442 D	7	021OA1GF	ESTs {IMAGE:738190}	738190 IMAGE Consortium	AA271058
6443 D	9	021OA1GF	ESTs {IMAGE:746599}	746599 IMAGE Consortium	AA268055
6444 D	11	021OA1GF	IMAGE EST {IMAGE:746445}	746445 IMAGE Consortium	AA259797
6445 F	1	021OA1GF	ESTs {IMAGE:734856}	734856 IMAGE Consortium	AA259388

6446 F	3	021OA1GF	IMAGE EST {IMAGE:737934}	737934	IMAGE Consortium	AA270991
6447 F	5	021OA1GF	WEE1-LIKE PROTEIN KINASE {IMAGE:746432}	746432	IMAGE Consortium	AA260001
6448 F	7	021OA1GF	ESTs, Highly similar to EUKARYOTIC TRANSLATION INITIATION FACTOR 1A [M.musculus] {IMAGE:746931}	746931	IMAGE Consortium	AA274423
6449 F	9	021OA1GF	Tenascin C {IMAGE:736372}	736372	IMAGE Consortium	AA270625
6450 F	11	021OA1GF	IMAGE EST {IMAGE:747443}	747443	IMAGE Consortium	AA272787
6451 H	1	021OA1GF	ESTs {IMAGE:738151}	738151	IMAGE Consortium	AA270943
6452 H	3	021OA1GF	ESTs {IMAGE:738162}	738162	IMAGE Consortium	AA271049
6453 H	5	021OA1GF	IMAGE EST {IMAGE:746955}	746955	IMAGE Consortium	AA275074
6454 H	7	021OA1GF	IMAGE EST {IMAGE:736334}	736334	IMAGE Consortium	AA270496
6455 H	9	021OA1GF	ESTs, Weakly similar to predicted using Genefinder [C.elegans] {IMAGE:736584}	736584	IMAGE Consortium	AA270652
6456 H	11	021OA1GF	ESTs, Weakly similar to NEDD-4 PROTEIN [M.musculus] {IMAGE:747655}	747655	IMAGE Consortium	AA268771
6457 B	1	021VA1GG	Lumican {IMAGE:746644}	746644	IMAGE Consortium	AA268082
6458 B	3	021VA1GG	ESTs {IMAGE:747208}	747208	IMAGE Consortium	AA260490
6459 B	5	021VA1GG	IMAGE EST {IMAGE:736556}	736556	IMAGE Consortium	AA270745
6460 B	7	021VA1GG	ESTs {IMAGE:737427}	737427	IMAGE Consortium	AA277184
6461 B	9	021VA1GG	ESTs, Weakly similar to Similarity to Yeast D-lactate dehydrogenase [C.elegans] {IMAGE:748228}	748228	IMAGE Consortium	AA274974
6462 B	11	021VA1GG	IMAGE EST {IMAGE:748263}	748263	IMAGE Consortium	AA274549
6463 D	1	021VA1GG	IMAGE EST {IMAGE:749482}	749482	IMAGE Consortium	AA289666
6464 D	3	021VA1GG	M.musculus T10 mRNA {IMAGE:750147}	750147	IMAGE Consortium	AA396437
6465 D	5	021VA1GG	ESTs, Weakly similar to PROBABLE LEUCYL-TRNA SYNTHETASE [C.elegans] {IMAGE:748061}	748061	IMAGE Consortium	AA275028
6466 D	7	021VA1GG	Suppressor of Ty 4 homolog (S. cerevisiae) {IMAGE:748241}	748241	IMAGE Consortium	AA274811
6467 D	9	021VA1GG	ESTs {IMAGE:750382}	750382	IMAGE Consortium	AA396764
6468 D	11	021VA1GG	ESTs {IMAGE:750782}	750782	IMAGE Consortium	AA290313
6469 F	1	021VA1GG	Mus musculus methyl-CpG binding protein MBD1 (Mbd1) mRNA, complete cds {IMAGE:750171}	750171	IMAGE Consortium	AA396451
6470 F	3	021VA1GG	ESTs, Moderately similar to DD96 [H.sapiens] {IMAGE:762437}	762437	IMAGE Consortium	AA274146
6471 F	5	021VA1GG	ESTs {IMAGE:750579}	750579	IMAGE Consortium	AA472952
6472 F	7	021VA1GG	IMAGE EST {IMAGE:748563}	748563	IMAGE Consortium	AA269488
6473 F	9	021VA1GG	ESTs {IMAGE:748330}	748330	IMAGE Consortium	AA267991
6474 F	11	021VA1GG	ESTs {IMAGE:748363}	748363	IMAGE Consortium	AA267936
6475 H	1	021VA1GG	Growth hormone releasing hormone {IMAGE:749837}	749837	IMAGE Consortium	AA388461
6476 H	3	021VA1GG	ESTs {IMAGE:762318}	762318	IMAGE Consortium	AA273761
6477 H	5	021VA1GG	Zinc finger protein 30 {IMAGE:747957}	747957	IMAGE Consortium	AA260956
6478 H	7	021VA1GG	H-2 CLASS II HISTOCOMPATIBILITY ANTIGEN, I-A BETA CHAIN PRECURSOR {IMAGE:748587}	748587	IMAGE Consortium	AA269724
6479 H	9	021VA1GG	IMAGE EST {IMAGE:750711}	750711	IMAGE Consortium	AA290235
6480 H	11	021VA1GG	ESTs, Highly similar to KIAA0436 [H.sapiens]	748947	IMAGE Consortium	AA288420



		{IMAGE:748947}		
6481 B	1	0212A1GH	Small inducible cytokine subfamily D, 1 {IMAGE:761708}	761708 IMAGE Consortium AA272232
6482 B	3	0212A1GH	Mus musculus transmembrane protein (Wfs1) mRNA, complete cds {IMAGE:762223}	762223 IMAGE Consortium AA277777
6483 B	5	0212A1GH	IMAGE EST {IMAGE:750521}	750521 IMAGE Consortium AA472962
6484 B	7	0212A1GH	ESTs {IMAGE:750857}	750857 IMAGE Consortium AA290125
6485 B	9	0212A1GH	ESTs {IMAGE:751160}	751160 IMAGE Consortium AA290491
6486 B	11	0212A1GH	ESTs {IMAGE:751473}	751473 IMAGE Consortium AA395994
6487 D	1	0212A1GH	Fibroblast growth factor inducible 15 {IMAGE:750018}	750018 IMAGE Consortium AA388985
6488 D	3	0212A1GH	Uncoupling protein 2, mitochondrial {IMAGE:748112}	748112 IMAGE Consortium AA260521
6489 D	5	0212A1GH	Mus musculus amine N-sulfotransferase (SULT-N) mRNA, complete cds {IMAGE:748125}	748125 IMAGE Consortium AA275042
6490 D	7	0212A1GH	IMAGE EST {IMAGE:750722}	750722 IMAGE Consortium AA289193
6491 D	9	0212A1GH	ESTs {IMAGE:751113}	751113 IMAGE Consortium AA289466
6492 D	11	0212A1GH	IMAGE EST {IMAGE:751619}	751619 IMAGE Consortium AA396068
6493 F	1	0212A1GH	IMAGE EST {IMAGE:748619}	748619 IMAGE Consortium AA270185
6494 F	3	0212A1GH	TATA box binding protein {IMAGE:751730}	751730 IMAGE Consortium AA396081
6495 F	5	0212A1GH	ESTs {IMAGE:751500}	751500 IMAGE Consortium AA396004
6496 F	7	0212A1GH	ESTs, Moderately similar to Similar to a C.elegans protein in cosmid C14H10 [H.sapiens] {IMAGE:749282}	749282 IMAGE Consortium AA290448
6497 F	9	0212A1GH	ESTs {IMAGE:749347}	749347 IMAGE Consortium AA289695
6498 F	11	0212A1GH	IMAGE EST {IMAGE:752348}	752348 IMAGE Consortium AA290064
6499 H	1	0212A1GH	ESTs {IMAGE:751572}	751572 IMAGE Consortium AA395943
6500 H	3	0212A1GH	IMAGE EST {IMAGE:751481}	751481 IMAGE Consortium AA389916
6501 H	5	0212A1GH	IMAGE EST {IMAGE:749337}	749337 IMAGE Consortium AA288562
6502 H	7	0212A1GH	Mus musculus mRNA for type II phosphatidylinositolphosphate kinase-alpha, complete cds {IMAGE:752194}	752194 IMAGE Consortium AA289027
6503 H	9	0212A1GH	ESTs {IMAGE:749510}	749510 IMAGE Consortium AA289544
6504 H	11	0212A1GH	IMAGE EST {IMAGE:749889}	749889 IMAGE Consortium AA389158
6505 B	1	0219A1GI	Integrin, alpha E, epithelial-associated {IMAGE:777194}	777194 IMAGE Consortium AA276764
6506 B	3	0219A1GI	IMAGE EST {IMAGE:762209}	762209 IMAGE Consortium AA277335
6507 B	5	0219A1GI	ESTs {IMAGE:762624}	762624 IMAGE Consortium AA271206
6508 B	7	0219A1GI	IMAGE EST {IMAGE:762542}	762542 IMAGE Consortium AA271510
6509 B	9	0219A1GI	ESTs {IMAGE:763085}	763085 IMAGE Consortium AA286605
6510 B	11	0219A1GI	IMAGE EST {IMAGE:775803}	775803 IMAGE Consortium AA275962
6511 D	1	0219A1GI	ESTs {IMAGE:762353}	762353 IMAGE Consortium AA273353
6512 D	3	0219A1GI	ESTs {IMAGE:762505}	762505 IMAGE Consortium AA271421
6513 D	5	0219A1GI	ESTs {IMAGE:762782}	762782 IMAGE Consortium AA271705
6514 D	7	0219A1GI	ESTs, Weakly similar to MEN chimeric transcription factor {IMAGE:762895}	762895 IMAGE Consortium AA286583
6515 D	9	0219A1GI	ESTs {IMAGE:775875}	775875 IMAGE Consortium AA276079
6516 D	11	0219A1GI	IMAGE EST {IMAGE:763365}	763365 IMAGE Consortium AA285678

6517 F	1 0219A1GI	Mouse mRNA for PSD-95/SAP90A, complete cds {IMAGE:765270}	765270 IMAGE Consortium	AA273310
6518 F	3 0219A1GI	ESTs {IMAGE:765377}	765377 IMAGE Consortium	AA275124
6519 F	5 0219A1GI	IMAGE EST {IMAGE:762377}	762377 IMAGE Consortium	AA273360
6520 F	7 0219A1GI	ESTs {IMAGE:762555}	762555 IMAGE Consortium	AA271495
6521 F	9 0219A1GI	ESTs {IMAGE:762854}	762854 IMAGE Consortium	AA271175
6522 F	11 0219A1GI	ESTs {IMAGE:763563}	763563 IMAGE Consortium	AA285875
6523 H	1 0219A1GI	Chromodomain helicase DNA binding protein 1 {IMAGE:765223}	765223 IMAGE Consortium	AA273291
6524 H	3 0219A1GI	IMAGE EST {IMAGE:765306}	765306 IMAGE Consortium	AA275237
6525 H	5 0219A1GI	Mus musculus Ets-family transcription factor Elf-1 mRNA, complete cds {IMAGE:765699}	765699 IMAGE Consortium	AA277070
6526 H	7 0219A1GI	IMAGE EST {IMAGE:766065}	766065 IMAGE Consortium	AA278005
6527 H	9 0219A1GI	ESTs {IMAGE:766685}	766685 IMAGE Consortium	AA275562
6528 H	11 0219A1GI	Mus musculus retinal short-chain dehydrogenase/reductase retSDR1 mRNA, complete cds {IMAGE:766482}	766482 IMAGE Consortium	AA275737
6529 B	1 021GA1GJ	IMAGE EST {IMAGE:751577}	751577 IMAGE Consortium	AA389933
6530 B	3 021GA1GJ	ESTs, Highly similar to PTB-ASSOCIATED SPLICING FACTOR [Homo sapiens] {IMAGE:751758}	751758 IMAGE Consortium	AA396118
6531 B	5 021GA1GJ	ESTs, Weakly similar to cDNA EST EMBL:D71941 comes from this gene [C.elegans] {IMAGE:752290}	752290 IMAGE Consortium	AA288972
6532 B	7 021GA1GJ	ESTs, Weakly similar to RANBP8 [H.sapiens] {IMAGE:749558}	749558 IMAGE Consortium	AA288567
6533 B	9 021GA1GJ	ESTs {IMAGE:761605}	761605 IMAGE Consortium	AA272367
6534 B	11 021GA1GJ	Mus musculus cGMP phosphodiesterase (PDE9A*1) mRNA, complete cds {IMAGE:762313}	762313 IMAGE Consortium	AA273765
6535 D	1 021GA1GJ	ESTs {IMAGE:749427}	749427 IMAGE Consortium	AA289578
6536 D	3 021GA1GJ	ESTs {IMAGE:749350}	749350 IMAGE Consortium	AA289693
6537 D	5 021GA1GJ	IMAGE EST {IMAGE:752302}	752302 IMAGE Consortium	AA290049
6538 D	7 021GA1GJ	IMAGE EST {IMAGE:761677}	761677 IMAGE Consortium	AA272146
6539 D	9 021GA1GJ	AMP deaminase 3 {IMAGE:750480}	750480 IMAGE Consortium	AA389897
6540 D	11 021GA1GJ	ESTs {IMAGE:750420}	750420 IMAGE Consortium	AA390005
6541 F	1 021GA1GJ	ESTs {IMAGE:752326}	752326 IMAGE Consortium	AA290057
6542 F	3 021GA1GJ	ESTs {IMAGE:749954}	749954 IMAGE Consortium	AA388790
6543 F	5 021GA1GJ	ESTs {IMAGE:749843}	749843 IMAGE Consortium	AA389146
6544 F	7 021GA1GJ	IMAGE EST {IMAGE:750313}	750313 IMAGE Consortium	AA396698
6545 F	9 021GA1GJ	ESTs, Moderately similar to GLUTAMATE DECARBOXYLASE, 67 KD ISOFORM [Rattus norvegicus] {IMAGE:748262}	748262 IMAGE Consortium	AA274541
6546 F	11 021GA1GJ	IMAGE EST {IMAGE:748471}	748471 IMAGE Consortium	AA268104
6547 H	1 021GA1GJ	Signal transducer and activator of transcription 1 {IMAGE:749544}	749544 IMAGE Consortium	AA388678
6548 H	3 021GA1GJ	IMAGE EST {IMAGE:749480}	749480 IMAGE Consortium	AA289665
6549 H	5 021GA1GJ	IMAGE EST {IMAGE:749752}	749752 IMAGE Consortium	AA288606

6550 H	7 021GA1GJ	IMAGE EST {IMAGE:749907}	749907 IMAGE Consortium	AA388656
6551 H	9 021GA1GJ	IMAGE EST {IMAGE:750577}	750577 IMAGE Consortium	AA472951
6552 H	11 021GA1GJ	IMAGE EST {IMAGE:765841}	765841 IMAGE Consortium	AA275367
6553 B	1 021NA1GK	ESTs {IMAGE:777225}	777225 IMAGE Consortium	AA276789
6554 B	3 021NA1GK	IMAGE EST {IMAGE:762210}	762210 IMAGE Consortium	AA277332
6555 B	5 021NA1GK	EST {IMAGE:762785}	762785 IMAGE Consortium	AA271590
6556 B	7 021NA1GK	ESTs {IMAGE:763042}	763042 IMAGE Consortium	AA271583
6557 B	9 021NA1GK	IMAGE EST {IMAGE:775616}	775616 IMAGE Consortium	AA275926
6558 B	11 021NA1GK	ESTs, Moderately similar to HYPOTHETICAL 13.6 KD PROTEIN IN NUP170-ILS1 INTERGENIC REGION [Saccharomyces cerevisiae] {IMAGE:775912}	775912 IMAGE Consortium	AA276338
6559 D	1 021NA1GK	IMAGE EST {IMAGE:777306}	777306 IMAGE Consortium	AA276959
6560 D	3 021NA1GK	ESTs {IMAGE:765724}	765724 IMAGE Consortium	AA277077
6561 D	5 021NA1GK	Male enhanced antigen 1 {IMAGE:762546}	762546 IMAGE Consortium	AA271081
6562 D	7 021NA1GK	IMAGE EST {IMAGE:766676}	766676 IMAGE Consortium	AA275644
6563 D	9 021NA1GK	ESTs, Moderately similar to SAS [H.sapiens] {IMAGE:763311}	763311 IMAGE Consortium	AA272449
6564 D	11 021NA1GK	Cytochrome P450, 2e1, ethanol inducible {IMAGE:775823}	775823 IMAGE Consortium	AA276059
6565 F	1 021NA1GK	ESTs {IMAGE:762558}	762558 IMAGE Consortium	AA271494
6566 F	3 021NA1GK	ESTs, Highly similar to PROTEASOME COMPONENT C9 [Homo sapiens] {IMAGE:766686}	766686 IMAGE Consortium	AA277958
6567 F	5 021NA1GK	M.musculus es64 mRNA {IMAGE:766438}	766438 IMAGE Consortium	AA277998
6568 F	7 021NA1GK	Cytochrome P450, 2e1, ethanol inducible {IMAGE:764009}	764009 IMAGE Consortium	AA285536
6569 F	9 021NA1GK	ESTs {IMAGE:776151}	776151 IMAGE Consortium	AA276180
6570 F	11 021NA1GK	ESTs, Highly similar to N-methyl-D-aspartate receptor glutamate-binding chain [R.norvegicus] {IMAGE:764513}	764513 IMAGE Consortium	AA273669
6571 H	1 021NA1GK	ESTs {IMAGE:775858}	775858 IMAGE Consortium	AA276003
6572 H	3 021NA1GK	GLUCOSE-6-PHOSPHATE ISOMERASE {IMAGE:776210}	776210 IMAGE Consortium	AA276216
6573 H	5 021NA1GK	Mus musculus pleiotropic regulator 1 (PLRG1) mRNA, complete cds {IMAGE:776426}	776426 IMAGE Consortium	AA286018
6574 H	7 021NA1GK	ESTs, Highly similar to hypertension-related protein [R.norvegicus] {IMAGE:776562}	776562 IMAGE Consortium	AA275888
6575 H	9 021NA1GK	ESTs, Weakly similar to D1046.5 [C.elegans] {IMAGE:764921}	764921 IMAGE Consortium	AA273388
6576 H	11 021NA1GK	ESTs {IMAGE:765218}	765218 IMAGE Consortium	AA273282
6577 B	1 021UA1GL	ESTs, Highly similar to UBIQUITIN CARBOXYL-TERMINAL HYDROLASE ISOZYME L1 [Rattus norvegicus] {IMAGE:763647}	763647 IMAGE Consortium	AA285969
6578 B	3 021UA1GL	ESTs {IMAGE:764385}	764385 IMAGE Consortium	AA277451
6579 B	5 021UA1GL	ESTs, Highly similar to SUCCINATE DEHYDROGENASE [Homo sapiens] {IMAGE:776319}	776319 IMAGE Consortium	AA276325
6580 B	7 021UA1GL	IMAGE EST {IMAGE:764634}	764634 IMAGE Consortium	AA273372
6581 B	9 021UA1GL	IMAGE EST {IMAGE:765031}	765031 IMAGE Consortium	AA273607

6582 B	11 021UA1GL	Mouse mRNA for CMP-N-acetylneuraminic acid hydroxylase, complete cds {IMAGE:765066}	765066 IMAGE Consortium	AA275061
6583 D	1 021UA1GL	IMAGE EST {IMAGE:775620}	775620 IMAGE Consortium	AA275745
6584 D	3 021UA1GL	IMAGE EST {IMAGE:763989}	763989 IMAGE Consortium	AA285504
6585 D	5 021UA1GL	ATP-binding cassette 2 {IMAGE:776243}	776243 IMAGE Consortium	AA276156
6586 D	7 021UA1GL	Mus musculus mRNA for glycoprotein-associated amino acid transporter y+LAT1b {IMAGE:776058}	776058 IMAGE Consortium	AA276085
6587 D	9 021UA1GL	Mus musculus mRNA for Guanine Nucleotide Regulatory Protein, partial cds {IMAGE:776403}	776403 IMAGE Consortium	AA285484
6588 D	11 021UA1GL	IMAGE EST {IMAGE:776348}	776348 IMAGE Consortium	AA275663
6589 F	1 021UA1GL	ESTs {IMAGE:764422}	764422 IMAGE Consortium	AA277200
6590 F	3 021UA1GL	IMAGE EST {IMAGE:764649}	764649 IMAGE Consortium	AA273174
6591 F	5 021UA1GL	Mus musculus B-ATF mRNA, complete cds {IMAGE:764444}	764444 IMAGE Consortium	AA273645
6592 F	7 021UA1GL	Mus musculus mRNA for NBAT, complete cds {IMAGE:776770}	776770 IMAGE Consortium	AA273497
6593 F	9 021UA1GL	Transcription factor Dp 1 {IMAGE:765649}	765649 IMAGE Consortium	AA275229
6594 F	11 021UA1GL	ESTs, Highly similar to glycogen phosphorylase [R.norvegicus] {IMAGE:807978}	807978 IMAGE Consortium	AA432928
6595 H	1 021UA1GL	Mus musculus parathion hydrolase (phosphotriesterase)-related protein mRNA, complete cds {IMAGE:803358}	803358 IMAGE Consortium	AA473017
6596 H	3 021UA1GL	Insulin-like growth factor binding protein, acid labile subunit {IMAGE:820122}	820122 IMAGE Consortium	AA423061
6597 H	5 021UA1GL	ESTs {IMAGE:807543}	807543 IMAGE Consortium	AA413050
6598 H	7 021UA1GL	IMAGE EST {IMAGE:807869}	807869 IMAGE Consortium	AA432648
6599 H	9 021UA1GL	ESTs {IMAGE:807836}	807836 IMAGE Consortium	AA466786
6600 H	11 021UA1GL	IMAGE EST {IMAGE:777513}	777513 IMAGE Consortium	AA276642
6601 B	1 0211A1GM	Tissue inhibitor of metalloproteinase 2 {IMAGE:831964}	831964 IMAGE Consortium	AA444490
6602 B	3 0211A1GM	IMAGE EST {IMAGE:808210}	808210 IMAGE Consortium	AA433070
6603 B	5 0211A1GM	IMAGE EST {IMAGE:808947}	808947 IMAGE Consortium	AA467719
6604 B	7 0211A1GM	IMAGE EST {IMAGE:809052}	809052 IMAGE Consortium	AA466044
6605 B	9 0211A1GM	ESTs {IMAGE:818293}	818293 IMAGE Consortium	AA435101
6606 B	11 0211A1GM	Mus musculus sex-determination protein homolog Fem1a gene, complete cds {IMAGE:818916}	818916 IMAGE Consortium	AA434902
6607 D	1 0211A1GM	ESTs, Highly similar to 3-HYDROXYANTHRANILATE 3,4-DIOXYGENASE [Homo sapiens] {IMAGE:777588}	777588 IMAGE Consortium	AA276839
6608 D	3 0211A1GM	IMAGE EST {IMAGE:777549}	777549 IMAGE Consortium	AA286542
6609 D	5 0211A1GM	IMAGE EST {IMAGE:818461}	818461 IMAGE Consortium	AA434761
6610 D	7 0211A1GM	IMAGE EST {IMAGE:818617}	818617 IMAGE Consortium	AA444335
6611 D	9 0211A1GM	ESTs {IMAGE:820204}	820204 IMAGE Consortium	AA423154
6612 D	11 0211A1GM	IMAGE EST {IMAGE:803470}	803470 IMAGE Consortium	AA473285
6613 F	1 0211A1GM	ESTs {IMAGE:777621}	777621 IMAGE Consortium	AA277006
6614 F	3 0211A1GM	IMAGE EST {IMAGE:817962}	817962 IMAGE Consortium	AA466533
6615 F	5 0211A1GM	IMAGE EST {IMAGE:818942}	818942 IMAGE Consortium	AA434916

6616 F	7 0211A1GM	ESTs, Weakly similar to slit protein 2 precursor [D.melanogaster] {IMAGE:820354}	820354 IMAGE Consortium	AA423126
6617 F	9 0211A1GM	ESTs, Highly similar to KIAA0091 gene product is related to subtilisin. [H.sapiens] {IMAGE:820121}	820121 IMAGE Consortium	AA423080
6618 F	11 0211A1GM	IMAGE EST {IMAGE:831918}	831918 IMAGE Consortium	AA444448
6619 H	1 0211A1GM	IMAGE EST {IMAGE:803309}	803309 IMAGE Consortium	AA472994
6620 H	3 0211A1GM	ESTs, Moderately similar to oxidase [R.norvegicus] {IMAGE:803510}	803510 IMAGE Consortium	AA473153
6621 H	5 0211A1GM	ESTs {IMAGE:820550}	820550 IMAGE Consortium	AA423418
6622 H	7 0211A1GM	IMAGE EST {IMAGE:831756}	831756 IMAGE Consortium	AA444946
6623 H	9 0211A1GM	IMAGE EST {IMAGE:808267}	808267 IMAGE Consortium	AA433023
6624 H	11 0211A1GM	Frizzled homolog 4, (Drosophila) {IMAGE:808829}	808829 IMAGE Consortium	AA467534
6625 B	1 0218A1GN	ESTs {IMAGE:807041}	807041 IMAGE Consortium	AA404094
6626 B	3 0218A1GN	ESTs, Moderately similar to CARDIAC PHOSPHOLAMBAN [Homo sapiens] {IMAGE:807663}	807663 IMAGE Consortium	AA470249
6627 B	5 0218A1GN	ESTs, Moderately similar to similar to a human major CRK-binding protein DOCK180. [H.sapiens] {IMAGE:777485}	777485 IMAGE Consortium	AA276470
6628 B	7 0218A1GN	Integrin beta 2 {IMAGE:808785}	808785 IMAGE Consortium	AA467489
6629 B	9 0218A1GN	ESTs {IMAGE:809088}	809088 IMAGE Consortium	AA435418
6630 B	11 0218A1GN	IMAGE EST {IMAGE:809084}	809084 IMAGE Consortium	AA435405
6631 D	1 0218A1GN	ESTs, Moderately similar to CARDIAC PHOSPHOLAMBAN [Homo sapiens] {IMAGE:807862}	807862 IMAGE Consortium	AA466795
6632 D	3 0218A1GN	IMAGE EST {IMAGE:777490}	777490 IMAGE Consortium	AA276669
6633 D	5 0218A1GN	ESTs {IMAGE:808996}	808996 IMAGE Consortium	AA466012
6634 D	7 0218A1GN	ESTs, Highly similar to RBP-MS/type 3 [H.sapiens] {IMAGE:818263}	818263 IMAGE Consortium	AA435387
6635 D	9 0218A1GN	IMAGE EST {IMAGE:820416}	820416 IMAGE Consortium	AA423305
6636 D	11 0218A1GN	Proteasome (prosome, macropain) 28 subunit, 3 {IMAGE:820184}	820184 IMAGE Consortium	AA423209
6637 F	1 0218A1GN	ESTs {IMAGE:818313}	818313 IMAGE Consortium	AA435123
6638 F	3 0218A1GN	ESTs {IMAGE:818792}	818792 IMAGE Consortium	AA467383
6639 F	5 0218A1GN	IMAGE EST {IMAGE:818615}	818615 IMAGE Consortium	AA444323
6640 F	7 0218A1GN	ESTs {IMAGE:820328}	820328 IMAGE Consortium	AA423201
6641 F	9 0218A1GN	ESTs, Moderately similar to COATOMER DELTA SUBUNIT [Homo sapiens] {IMAGE:807044}	807044 IMAGE Consortium	AA404092
6642 F	11 0218A1GN	GTP-BINDING PROTEIN SARA {IMAGE:831879}	831879 IMAGE Consortium	AA444425
6643 H	1 0218A1GN	Insulin-like growth factor binding protein 10 {IMAGE:820167}	820167 IMAGE Consortium	AA423149
6644 H	3 0218A1GN	IMAGE EST {IMAGE:807602}	807602 IMAGE Consortium	AA412821
6645 H	5 0218A1GN	Mus musculus ENDOG mRNA for endonuclease G, complete cds {IMAGE:807619}	807619 IMAGE Consortium	AA412908
6646 H	7 0218A1GN	ESTs {IMAGE:777430}	777430 IMAGE Consortium	AA276585
6647 H	9 0218A1GN	IMAGE EST {IMAGE:777487}	777487 IMAGE Consortium	AA276471
6648 H	11 0218A1GN	ESTs {IMAGE:777851}	777851 IMAGE Consortium	AA286418

6649 B	1 021FA1GO	Mouse mRNA for NDPP-1 protein, complete cds {IMAGE:808083}	808083 IMAGE Consortium	AA444365
6650 B	3 021FA1GO			
6651 B	5 021FA1GO			
6652 B	7 021FA1GO			
6653 B	9 021FA1GO			
6654 B	11 021FA1GO			
6655 D	1 021FA1GO			
6656 D	3 021FA1GO			
6657 D	5 021FA1GO			
6658 D	7 021FA1GO			
6659 D	9 021FA1GO			
6660 D	11 021FA1GO			
6661 F	1 021FA1GO			
6662 F	3 021FA1GO			
6663 F	5 021FA1GO			
6664 F	7 021FA1GO			
6665 F	9 021FA1GO			
6666 F	11 021FA1GO			
6667 H	1 021FA1GO			
6668 H	3 021FA1GO			
6669 H	5 021FA1GO			
6670 H	7 021FA1GO			
6671 H	9 021FA1GO			
6672 H	11 021FA1GO			
6673 B	1 021MA1GP	IMAGE EST {IMAGE:521951}	521951 IMAGE Consortium	AA106263
6674 B	3 021MA1GP	IMAGE EST {IMAGE:533314}	533314 IMAGE Consortium	AA104976
6675 B	5 021MA1GP	ESTs {IMAGE:534015}	534015 IMAGE Consortium	AA072800
6676 B	7 021MA1GP	Transformed mouse 3T3 cell double minute 2 {IMAGE:523982}	523982 IMAGE Consortium	AA072086
6677 B	9 021MA1GP	IMAGE EST {IMAGE:535228}	535228 IMAGE Consortium	AA086610
6678 B	11 021MA1GP	ESTs {IMAGE:535221}	535221 IMAGE Consortium	AA086803
6679 D	1 021MA1GP	ESTs, Highly similar to NADH-CYTOCHROME B5 REDUCTASE [Homo sapiens] {IMAGE:524540}	524540 IMAGE Consortium	AA068440
6680 D	3 021MA1GP	IMAGE EST {IMAGE:551016}	551016 IMAGE Consortium	AA080332
6681 D	5 021MA1GP	ESTs, Moderately similar to PROACTIVATOR POLYPEPTIDE PRECURSOR [Homo sapiens] {IMAGE:522891}	522891 IMAGE Consortium	AA065868
6682 D	7 021MA1GP	IMAGE EST {IMAGE:523123}	523123 IMAGE Consortium	AA067065
6683 D	9 021MA1GP	ESTs {IMAGE:523380}	523380 IMAGE Consortium	AA080169
6684 D	11 021MA1GP	IMAGE EST {IMAGE:524306}	524306 IMAGE Consortium	AA066209
6685 F	1 021MA1GP	IMAGE EST {IMAGE:523572}	523572 IMAGE Consortium	AA097314
6686 F	3 021MA1GP	ESTs {IMAGE:524331}	524331 IMAGE Consortium	AA065728

6687 F	5 021MA1GP	Phosphoribosylglycinamide formyltransferase {IMAGE:535188}	535188 IMAGE Consortium	AA086734
6688 F	7 021MA1GP	ESTs {IMAGE:535706}	535706 IMAGE Consortium	AA106225
6689 F	9 021MA1GP	Mus musculus mRNA for ganglioside-induced differentiation associated protein 2 {IMAGE:524541}	524541 IMAGE Consortium	AA068478
6690 F	11 021MA1GP	IMAGE EST {IMAGE:532250}	532250 IMAGE Consortium	AA066739
6691 H	1 021MA1GP	Mus musculus PAF acetylhydrolase mRNA, complete cds {IMAGE:536464}	536464 IMAGE Consortium	AA073963
6692 H	3 021MA1GP	ESTs, Highly similar to CALCIUM/CALMODULIN-DEPENDENT PROTEIN KINASE TYPE II GAMMA CHAIN [Rattus norvegicus] {IMAGE:532707}	532707 IMAGE Consortium	AA068735
6693 H	5 021MA1GP	IMAGE EST {IMAGE:522713}	522713 IMAGE Consortium	AA087673
6694 H	7 021MA1GP	Small inducible cytokine A3 {IMAGE:533862}	533862 IMAGE Consortium	AA071899
6695 H	9 021MA1GP	IMAGE EST {IMAGE:523795}	523795 IMAGE Consortium	AA067194
6696 H	11 021MA1GP	IMAGE EST {IMAGE:524177}	524177 IMAGE Consortium	AA065624
6697 B	1 021TA1GQ	ESTs, Weakly similar to F45G2.h [C.elegans] {IMAGE:535937}	535937 IMAGE Consortium	AA105444
6698 B	3 021TA1GQ	IMAGE EST {IMAGE:536192}	536192 IMAGE Consortium	AA073402
6699 B	5 021TA1GQ	Mus musculus mRNA for cellular nucleic acid binding protein {IMAGE:533147}	533147 IMAGE Consortium	AA068939
6700 B	7 021TA1GQ	IMAGE EST {IMAGE:523965}	523965 IMAGE Consortium	AA067731
6701 B	9 021TA1GQ	ESTs, Highly similar to SRB7 [H.sapiens] {IMAGE:524274}	524274 IMAGE Consortium	AA067890
6702 B	11 021TA1GQ	ESTs {IMAGE:524705}	524705 IMAGE Consortium	AA068506
6703 D	1 021TA1GQ	ESTs, Highly similar to SIGNAL RECOGNITION PARTICLE 72 KD PROTEIN [Canis familiaris] {IMAGE:524907}	524907 IMAGE Consortium	AA089038
6704 D	3 021TA1GQ	ESTs, Highly similar to PTB-ASSOCIATED SPLICING FACTOR [Homo sapiens] {IMAGE:550916}	550916 IMAGE Consortium	AA098022
6705 D	5 021TA1GQ	IMAGE EST {IMAGE:533405}	533405 IMAGE Consortium	AA105002
6706 D	7 021TA1GQ	IMAGE EST {IMAGE:523303}	523303 IMAGE Consortium	AA080194
6707 D	9 021TA1GQ	ESTs {IMAGE:534178}	534178 IMAGE Consortium	AA072617
6708 D	11 021TA1GQ	IMAGE EST {IMAGE:524338}	524338 IMAGE Consortium	AA065706
6709 F	1 021TA1GQ	ESTs, Highly similar to ERYTHROCYTE ADDUCIN BETA SUBUNIT [Homo sapiens] {IMAGE:524196}	524196 IMAGE Consortium	AA066202
6710 F	3 021TA1GQ	IMAGE EST {IMAGE:390652}	390652 IMAGE Consortium	W71798
6711 F	5 021TA1GQ	IMAGE EST {IMAGE:535892}	535892 IMAGE Consortium	AA105471
6712 F	7 021TA1GQ	Integral membrane protein 2 {IMAGE:532350}	532350 IMAGE Consortium	AA066921
6713 F	9 021TA1GQ	ESTs, Moderately similar to RAS-RELATED PROTEIN RAB-20 [M.musculus] {IMAGE:550869}	550869 IMAGE Consortium	AA098010
6714 F	11 021TA1GQ	Mus musculus Rho-associated, coiled-coil forming protein kinase p160 ROCK-2 mRNA, complete cds {IMAGE:550956}	550956 IMAGE Consortium	AA098168
6715 H	1 021TA1GQ	Cytochrome P450, 2e1, ethanol inducible {IMAGE:656925}	656925 IMAGE Consortium	AA240739
6716 H	3 021TA1GQ	ESTs {IMAGE:663108}	663108 IMAGE Consortium	AA222883
6717 H	5 021TA1GQ	ESTs {IMAGE:732526}	732526 IMAGE Consortium	AA277060
6718 H	7 021TA1GQ	IMAGE EST {IMAGE:656701}	656701 IMAGE Consortium	AA241295

6719 H	9 021TA1GQ	Caudal type homeo box 1 {IMAGE:633635}	633635 IMAGE Consortium	AA184566
6720 H	11 021TA1GQ	ESTs, Moderately similar to RAF SERINE/THREONINE-PROTEIN KINASE TRANSFORMING PROTEIN [Murine sarcoma virus 3611] {IMAGE:651430}	651430 IMAGE Consortium	AA215234
6721 B	1 0210A1GR	Aquaporin 1 {IMAGE:656654}	656654 IMAGE Consortium	AA241281
6722 B	3 0210A1GR	ESTs {IMAGE:661780}	661780 IMAGE Consortium	AA207878
6723 B	5 0210A1GR	ESTs, Highly similar to HM-1 KILLER TOXIN RESISTANCE PROTEIN [Saccharomyces cerevisiae] {IMAGE:652131}	652131 IMAGE Consortium	AA215144
6724 B	7 0210A1GR	IMAGE EST {IMAGE:654228}	654228 IMAGE Consortium	AA197982
6725 B	9 0210A1GR	IMAGE EST {IMAGE:656372}	656372 IMAGE Consortium	AA242145
6726 B	11 0210A1GR	IMAGE EST {IMAGE:656870}	656870 IMAGE Consortium	AA240043
6727 D	1 0210A1GR	IMAGE EST {IMAGE:654099}	654099 IMAGE Consortium	AA197818
6728 D	3 0210A1GR	Mus musculus receptor tyrosine kinase mRNA, complete cds {IMAGE:905055}	905055 IMAGE Consortium	AA518290
6729 D	5 0210A1GR	ESTs, Moderately similar to GLYPICAN-3 PRECURSOR [Rattus norvegicus] {IMAGE:657264}	657264 IMAGE Consortium	AA240923
6730 D	7 0210A1GR	ESTs {IMAGE:662476}	662476 IMAGE Consortium	AA207912
6731 D	9 0210A1GR	IMAGE EST {IMAGE:651692}	651692 IMAGE Consortium	AA212418
6732 D	11 0210A1GR	IMAGE EST {IMAGE:917654}	917654 IMAGE Consortium	AA492963
6733 F	1 0210A1GR	IMAGE EST {IMAGE:656063}	656063 IMAGE Consortium	AA239753
6734 F	3 0210A1GR	Mus musculus FK506-binding protein (FKBP23) mRNA, complete cds {IMAGE:656375}	656375 IMAGE Consortium	AA242149
6735 F	5 0210A1GR	IMAGE EST {IMAGE:608351}	608351 IMAGE Consortium	AA162065
6736 F	7 0210A1GR	IMAGE EST {IMAGE:661607}	661607 IMAGE Consortium	AA208314
6737 F	9 0210A1GR	Glutathione S-transferase, theta 2 {IMAGE:663169}	663169 IMAGE Consortium	AA230348
6738 F	11 0210A1GR	IMAGE EST {IMAGE:652207}	652207 IMAGE Consortium	AA215024
6739 H	1 0210A1GR			
6740 H	3 0210A1GR			
6741 H	5 0210A1GR			
6742 H	7 0210A1GR			
6743 H	9 0210A1GR			
6744 H	11 0210A1GR			
6745 B	1 021Q9974	Control: Ratio 1:3 (Cy3:Cy5)	Ratio_A	Controls
6746 B	3 021Q9974	Control: Ratio 3:1 (Cy3:Cy5)	Ratio_B	Controls
6747 B	5 021Q9974	Control: Ratio 1:10 (Cy3:Cy5)	Ratio_C	Controls
6748 B	7 021Q9974	Control: Ratio 10:1 (Cy3:Cy5)	Ratio_D	Controls
6749 B	9 021Q9974	Control: Ratio 1:25 (Cy3:Cy5)	Ratio_E	Controls
6750 B	11 021Q9974	Control: Ratio 25:1 (Cy3:Cy5)	Ratio_F	Controls
6751 D	1 021Q9974	Control: Sensitivity 2pg	Sensitivity_A	Controls
6752 D	3 021Q9974	Control: Sensitivity 20pg	Sensitivity_B	Controls
6753 D	5 021Q9974	Control: Sensitivity 200pg	Sensitivity_C	Controls
6754 D	7 021Q9974	Control: Sensitivity 2000pg	Sensitivity_D	Controls
6755 D	9 021Q9974	Control: Buffer Only	Buffer_Only_Control	Controls



6756 D	11	021Q9974	Control: Buffer Only	Buffer_Only_Control	Controls		
6757 F	1	021Q9974	Control: Internal	Internal_Control_A	Controls		
6758 F	3	021Q9974	Control: Buffer Only	Buffer_Only_Control	Controls		
6759 F	5	021Q9974	Control: Internal	Internal_Control_B	Controls		
6760 F	7	021Q9974	Control: Buffer Only	Buffer_Only_Control	Controls		
6761 F	9	021Q9974	Control: Internal	Internal_Control_C	Controls		
6762 F	11	021Q9974	Control: Internal	Internal_Control_D	Controls		
6763 H	1	021Q9974	Control: Complex Target (Mus musculus)	RNA_Quality_MM	Controls		
6764 H	3	021Q9974	Control: Complex Target (Mus musculus)	RNA_Quality_MM	Controls		
6765 H	5	021Q9974	Control: Complex Target (Mus musculus)	RNA_Quality_MM	Controls		
6766 H	7	021Q9974	Control: Housekeeping (Alpha Tubulin)	Housekeeping_B	Controls		
6767 H	9	021Q9974	Control: Housekeeping (Ribosomal S9)	Housekeeping_A	Controls		
6768 H	11	021Q9974	Control: Housekeeping (23kD HBP)	Housekeeping_C	Controls		
6769 B	2	2159971	Internal_Control_E	Internal_Control_E	Controls		
6770 B	4	2159971	Internal_Control_F	Internal_Control_F	Controls		
6771 B	6	2159971	Internal_Control_G	Internal_Control_G	Controls		
6772 B	8	2159971	Internal_Control_H	Internal_Control_H	Controls		
6773 B	10	2159971	Internal_Control_I	Internal_Control_I	Controls		
6774 B	12	2159971	Internal_Control_J	Internal_Control_J	Controls		
6775 D	2	2159971	Internal_Control_K	Internal_Control_K	Controls		
6776 D	4	2159971	Internal_Control_L	Internal_Control_L	Controls		
6777 D	6	2159971	Internal_Control_M	Internal_Control_M	Controls		
6778 D	8	2159971	Internal_Control_N	Internal_Control_N	Controls		
6779 D	10	2159971	Internal_Control_O	Internal_Control_O	Controls		
6780 D	12	2159971	Internal_Control_P	Internal_Control_P	Controls		
6781 F	2	2159971	Internal_Control_Q	Internal_Control_Q	Controls		
6782 F	4	2159971	Internal_Control_R	Internal_Control_R	Controls		
6783 F	6	2159971	Internal_Control_S	Internal_Control_S	Controls		
6784 F	8	2159971	Internal_Control_T	Internal_Control_T	Controls		
6785 F	10	2159971	Internal_Control_U	Internal_Control_U	Controls		
6786 F	12	2159971	Internal_Control_V	Internal_Control_V	Controls		
6787 H	2	2159971	Internal_Control_W	Internal_Control_W	Controls		
6788 H	4	2159971	Internal_Control_X	Internal_Control_X	Controls		
6789 H	6	2159971	Internal_Control_Y	Internal_Control_Y	Controls		
6790 H	8	2159971	Internal_Control_Z	Internal_Control_Z	Controls		
6791 H	10	2159971	Internal_Control_AA	Internal_Control_AA	Controls		
6792 H	12	2159971	Internal_Control_AB	Internal_Control_AB	Controls		
6793 B	2	021MA1E8	IMAGE EST {IMAGE:893856}	893856	IMAGE Consortium	AA517194	
6794 B	4	021MA1E8	IMAGE EST {IMAGE:760918}	760918	IMAGE Consortium	AA386814	
6795 B	6	021MA1E8	IMAGE EST {IMAGE:760972}	760972	IMAGE Consortium	AA387448	
6796 B	8	021MA1E8	IMAGE EST {IMAGE:761158}	761158	IMAGE Consortium	AA387854	
6797 B	10	021MA1E8	ESTs {IMAGE:804773}	804773	IMAGE Consortium	AA472816	

6798 B	12 021MA1E8	ESTs {IMAGE:934425}	934425 IMAGE Consortium	AA545046
6799 D	2 021MA1E8	IMAGE EST {IMAGE:805419}	805419 IMAGE Consortium	AA474499
6800 D	4 021MA1E8	ESTs {IMAGE:805579}	805579 IMAGE Consortium	AA397280
6801 D	6 021MA1E8	ESTs {IMAGE:834225}	834225 IMAGE Consortium	AA413194
6802 D	8 021MA1E8	IMAGE EST {IMAGE:760858}	760858 IMAGE Consortium	AA386685
6803 D	10 021MA1E8	Mus musculus serine palmitoyltransferase LCB1 subunit mRNA, complete cds {IMAGE:760865}	760865 IMAGE Consortium	AA386799
6804 D	12 021MA1E8	IMAGE EST {IMAGE:761298}	761298 IMAGE Consortium	AA388321
6805 F	2 021MA1E8	ESTs {IMAGE:761346}	761346 IMAGE Consortium	AA388142
6806 F	4 021MA1E8	ESTs {IMAGE:804778}	804778 IMAGE Consortium	AA472810
6807 F	6 021MA1E8	IMAGE EST {IMAGE:805613}	805613 IMAGE Consortium	AA399831
6808 F	8 021MA1E8	ESTs, Highly similar to retinoic acid-regulated protein pH 34 [M.musculus] {IMAGE:944963}	944963 IMAGE Consortium	AA537161
6809 F	10 021MA1E8	ESTs {IMAGE:834159}	834159 IMAGE Consortium	AA413119
6810 F	12 021MA1E8	ESTs {IMAGE:835159}	835159 IMAGE Consortium	AA467238
6811 H	2 021MA1E8	ESTs {IMAGE:805046}	805046 IMAGE Consortium	AA397197
6812 H	4 021MA1E8	IMAGE EST {IMAGE:805822}	805822 IMAGE Consortium	AA399850
6813 H	6 021MA1E8	ESTs, Highly similar to CAD PROTEIN [Homo sapiens] {IMAGE:834541}	834541 IMAGE Consortium	AA466758
6814 H	8 021MA1E8	Kinesin heavy chain member 5B, ubiquitous {IMAGE:760837}	760837 IMAGE Consortium	AA386680
6815 H	10 021MA1E8	ESTs, Moderately similar to PHOSPHATIDYLINOSITOL [Homo sapiens] {IMAGE:761041}	761041 IMAGE Consortium	AA387401
6816 H	12 021MA1E8	IMAGE EST {IMAGE:932884}	932884 IMAGE Consortium	AA541870
6817 B	2 021TA1E9	ESTs, Weakly similar to HIRA protein [M.musculus] {IMAGE:761060}	761060 IMAGE Consortium	AA387585
6818 B	4 021TA1E9	ESTs, Highly similar to hNop56 [H.sapiens] {IMAGE:933491}	933491 IMAGE Consortium	AA542348
6819 B	6 021TA1E9	ESTs {IMAGE:804791}	804791 IMAGE Consortium	AA472821
6820 B	8 021TA1E9	Peroxisomal membrane protein, 70 kDa {IMAGE:805835}	805835 IMAGE Consortium	AA437485
6821 B	10 021TA1E9	ESTs {IMAGE:833829}	833829 IMAGE Consortium	AA412925
6822 B	12 021TA1E9	IMAGE EST {IMAGE:945689}	945689 IMAGE Consortium	AA546195
6823 D	2 021TA1E9	Mus musculus ATP-dependent metalloprotease FtsH1 mRNA, complete cds {IMAGE:833670}	833670 IMAGE Consortium	AA466714
6824 D	4 021TA1E9	ESTs {IMAGE:834572}	834572 IMAGE Consortium	AA470303
6825 D	6 021TA1E9	M.musculus mRNA for SOX15 protein {IMAGE:836802}	836802 IMAGE Consortium	AA426845
6826 D	8 021TA1E9	IMAGE EST {IMAGE:760917}	760917 IMAGE Consortium	AA386815
6827 D	10 021TA1E9	IMAGE EST {IMAGE:761009}	761009 IMAGE Consortium	AA387552
6828 D	12 021TA1E9	IMAGE EST {IMAGE:775154}	775154 IMAGE Consortium	AA387003
6829 F	2 021TA1E9	IMAGE EST {IMAGE:774843}	774843 IMAGE Consortium	AA386749
6830 F	4 021TA1E9	ESTs, Highly similar to SODIUM-DEPENDENT CHOLINE TRANSPORTER [Rattus norvegicus] {IMAGE:805483}	805483 IMAGE Consortium	AA473227
6831 F	6 021TA1E9	ESTs {IMAGE:833720}	833720 IMAGE Consortium	AA466731
6832 F	8 021TA1E9	ESTs, Moderately similar to JAK3 [M.musculus] {IMAGE:834549}	834549 IMAGE Consortium	AA470301

6833 F	10 021TA1E9	ESTs {IMAGE:834783}	834783 IMAGE Consortium	AA467436
6834 F	12 021TA1E9	ESTs {IMAGE:837565}	837565 IMAGE Consortium	AA432818
6835 H	2 021TA1E9	IMAGE EST {IMAGE:775092}	775092 IMAGE Consortium	AA386832
6836 H	4 021TA1E9	IMAGE EST {IMAGE:775507}	775507 IMAGE Consortium	AA387272
6837 H	6 021TA1E9	ESTs, Highly similar to Diff40 gene product [H.sapiens] {IMAGE:775158}	775158 IMAGE Consortium	AA387005
6838 H	8 021TA1E9	IMAGE EST {IMAGE:775394}	775394 IMAGE Consortium	AA387250
6839 H	10 021TA1E9	IMAGE EST {IMAGE:775555}	775555 IMAGE Consortium	AA387288
6840 H	12 021TA1E9	IMAGE EST {IMAGE:774811}	774811 IMAGE Consortium	AA386714
6841 B	2 021DA1EA	ESTs, Moderately similar to weak similarity to Arabidopsis thaliana ubiquitin-like protein 8 [C.elegans] {IMAGE:780019}	780019 IMAGE Consortium	AA414936
6842 B	4 021DA1EA	IMAGE EST {IMAGE:804255}	804255 IMAGE Consortium	AA474427
6843 B	6 021DA1EA	ESTs, Highly similar to DTDP-4-DEHYDRORHAMNOSE REDUCTASE [Salmonella typhimurium] {IMAGE:835756}	835756 IMAGE Consortium	AA422576
6844 B	8 021DA1EA	Cyclin B1, related sequence 1 {IMAGE:836311}	836311 IMAGE Consortium	AA426917
6845 B	10 021DA1EA	Activating transcription factor 2 {IMAGE:778942}	778942 IMAGE Consortium	AA414544
6846 B	12 021DA1EA	ESTs, Highly similar to AMIDOPHOSPHORIBOSYLTRANSFERASE PRECURSOR [Rattus norvegicus] {IMAGE:779343}	779343 IMAGE Consortium	AA414106
6847 D	2 021DA1EA	Mus musculus metal response element DNA-binding protein M96 mRNA, complete cds {IMAGE:779876}	779876 IMAGE Consortium	AA414761
6848 D	4 021DA1EA	ESTs {IMAGE:804212}	804212 IMAGE Consortium	AA474452
6849 D	6 021DA1EA	ESTs {IMAGE:836414}	836414 IMAGE Consortium	AA426924
6850 D	8 021DA1EA	IMAGE EST {IMAGE:792923}	792923 IMAGE Consortium	AA415580
6851 D	10 021DA1EA	IMAGE EST {IMAGE:779598}	779598 IMAGE Consortium	AA413691
6852 D	12 021DA1EA	IMAGE EST {IMAGE:803622}	803622 IMAGE Consortium	AA422792
6853 F	2 021DA1EA	ESTs {IMAGE:833200}	833200 IMAGE Consortium	AA432985
6854 F	4 021DA1EA	ESTs {IMAGE:793591}	793591 IMAGE Consortium	AA415557
6855 F	6 021DA1EA	ESTs {IMAGE:833160}	833160 IMAGE Consortium	AA467585
6856 F	8 021DA1EA	ESTs {IMAGE:836420}	836420 IMAGE Consortium	AA426926
6857 F	10 021DA1EA	ESTs, Highly similar to tpr protein [H.sapiens] {IMAGE:793129}	793129 IMAGE Consortium	AA437983
6858 F	12 021DA1EA	Activating transcription factor 2 {IMAGE:779278}	779278 IMAGE Consortium	AA445861
6859 H	2 021DA1EA	ESTs, Weakly similar to 112.3 KD PROTEIN IN PYK1-SNC1 INTERGENIC REGION [S.cerevisiae] {IMAGE:944355}	944355 IMAGE Consortium	AA547684
6860 H	4 021DA1EA	ESTs {IMAGE:961114}	961114 IMAGE Consortium	AA547315
6861 H	6 021DA1EA	RAD21 homolog (S. pombe) {IMAGE:959708}	959708 IMAGE Consortium	AA546839
6862 H	8 021DA1EA	IMAGE EST {IMAGE:959804}	959804 IMAGE Consortium	AA546948
6863 H	10 021DA1EA	IMAGE EST {IMAGE:961056}	961056 IMAGE Consortium	AA547175
6864 H	12 021DA1EA	IMAGE EST {IMAGE:961224}	961224 IMAGE Consortium	AA547514
6865 B	2 021KA1EB	Mus musculus apoptotic protease activating factor 1 mRNA, complete cds {IMAGE:775111}	775111 IMAGE Consortium	AA387058
6866 B	4 021KA1EB	ESTs {IMAGE:775135}	775135 IMAGE Consortium	AA387002

6867 B	6 021KA1EB	IMAGE EST {IMAGE:775472}	775472 IMAGE Consortium	AA388125
6868 B	8 021KA1EB	IMAGE EST {IMAGE:775408}	775408 IMAGE Consortium	AA386915
6869 B	10 021KA1EB	IMAGE EST {IMAGE:775420}	775420 IMAGE Consortium	AA387342
6870 B	12 021KA1EB	IMAGE EST {IMAGE:789806}	789806 IMAGE Consortium	AA388019
6871 D	2 021KA1EB	ESTs {IMAGE:778359}	778359 IMAGE Consortium	AA413552
6872 D	4 021KA1EB	ESTs {IMAGE:778456}	778456 IMAGE Consortium	AA413505
6873 D	6 021KA1EB	ESTs, Weakly similar to butyrophilin [H.sapiens] {IMAGE:778266}	778266 IMAGE Consortium	AA413760
6874 D	8 021KA1EB	SAPK/Erk/kinase 1 {IMAGE:778459}	778459 IMAGE Consortium	AA413508
6875 D	10 021KA1EB	IMAGE EST {IMAGE:778697}	778697 IMAGE Consortium	AA413752
6876 D	12 021KA1EB	Protein kinase C substrate 80K-H {IMAGE:793127}	793127 IMAGE Consortium	AA437882
6877 F	2 021KA1EB	ESTs {IMAGE:791079}	791079 IMAGE Consortium	AA415185
6878 F	4 021KA1EB	CLARA CELL PHOSPHOLIPID-BINDING PROTEIN PRECURSOR {IMAGE:833469}	833469 IMAGE Consortium	AA432689
6879 F	6 021KA1EB	ESTs, Moderately similar to DMR-N9 PROTEIN [M.musculus] {IMAGE:836047}	836047 IMAGE Consortium	AA426739
6880 F	8 021KA1EB	ESTs, Highly similar to herpesvirus associated ubiquitin-specific protease [H.sapiens] {IMAGE:793038}	793038 IMAGE Consortium	AA444224
6881 F	10 021KA1EB	Max protein {IMAGE:793537}	793537 IMAGE Consortium	AA415602
6882 F	12 021KA1EB	ESTs, Weakly similar to HYPOTHETICAL 19.4 KD PROTEIN D2007.4 IN CHROMOSOME III [C.elegans] {IMAGE:803817}	803817 IMAGE Consortium	AA422973
6883 H	2 021KA1EB	ESTs {IMAGE:791176}	791176 IMAGE Consortium	AA415516
6884 H	4 021KA1EB	ESTs {IMAGE:791570}	791570 IMAGE Consortium	AA437456
6885 H	6 021KA1EB	ESTs, Highly similar to HYPOTHETICAL 51.6 KD PROTEIN F59B2.5 IN CHROMOSOME III [Caenorhabditis elegans] {IMAGE:833508}	833508 IMAGE Consortium	AA465980
6886 H	8 021KA1EB	ESTs {IMAGE:836133}	836133 IMAGE Consortium	AA426897
6887 H	10 021KA1EB	ESTs, Moderately similar to No similarities to any reported proteins [H.sapiens] {IMAGE:793570}	793570 IMAGE Consortium	AA415241
6888 H	12 021KA1EB	RAB7, member RAS oncogene family {IMAGE:779604}	779604 IMAGE Consortium	AA413694
6889 B	2 021RA1EC	TUBULIN ALPHA-4 CHAIN {IMAGE:329726}	329726 IMAGE Consortium	W11746
6890 B	4 021RA1EC	ESTs, Moderately similar to SCABROUS PROTEIN PRECURSOR [Drosophila melanogaster] {IMAGE:329741}	329741 IMAGE Consortium	W13905
6891 B	6 021RA1EC	ESTs, Moderately similar to acetolactate synthase homolog [H.sapiens] {IMAGE:330168}	330168 IMAGE Consortium	W11619
6892 B	8 021RA1EC	Thymus cell antigen 1, theta {IMAGE:316409}	316409 IMAGE Consortium	W13151
6893 B	10 021RA1EC	ESTs {IMAGE:330825}	330825 IMAGE Consortium	W14925
6894 B	12 021RA1EC	ESTs {IMAGE:317286}	317286 IMAGE Consortium	W13718
6895 D	2 021RA1EC	Surfeit gene 4 {IMAGE:316113}	316113 IMAGE Consortium	W12260
6896 D	4 021RA1EC	ESTs {IMAGE:330222}	330222 IMAGE Consortium	W13932
6897 D	6 021RA1EC	IMAGE EST {IMAGE:330402}	330402 IMAGE Consortium	W14992
6898 D	8 021RA1EC	ESTs {IMAGE:317409}	317409 IMAGE Consortium	W13357
6899 D	10 021RA1EC	Insulin-like growth factor 1 {IMAGE:313322}	313322 IMAGE Consortium	W10072

6900 D	12 021RA1EC	ESTs {IMAGE:313430}	313430 IMAGE Consortium	W10025
6901 F	2 021RA1EC	ESTs {IMAGE:330128}	330128 IMAGE Consortium	W11759
6902 F	4 021RA1EC	Receptor-associated protein of the synapse, 43 kDa {IMAGE:313466}	313466 IMAGE Consortium	W10155
6903 F	6 021RA1EC	ESTs {IMAGE:317597}	317597 IMAGE Consortium	W11380
6904 F	8 021RA1EC	Phosphodiesterase 7A {IMAGE:314384}	314384 IMAGE Consortium	W10680
6905 F	10 021RA1EC	Glypican 4 {IMAGE:314639}	314639 IMAGE Consortium	W12478
6906 F	12 021RA1EC	Mus musculus developmentally regulated repeat (Dr-3), mRNA {IMAGE:318922}	318922 IMAGE Consortium	W34507
6907 H	2 021RA1EC	ESTs {IMAGE:317958}	317958 IMAGE Consortium	W11432
6908 H	4 021RA1EC	IMAGE EST {IMAGE:317848}	317848 IMAGE Consortium	W13063
6909 H	6 021RA1EC	Creatine kinase, brain {IMAGE:315775}	315775 IMAGE Consortium	W09925
6910 H	8 021RA1EC	ESTs {IMAGE:329816}	329816 IMAGE Consortium	W13971
6911 H	10 021RA1EC	ESTs, Weakly similar to HYPOTHETICAL 11.4 KD PROTEIN C13G6.04 IN CHROMOSOME I [Schizosaccharomyces pombe] {IMAGE:330146}	330146 IMAGE Consortium	W11535
6912 H	12 021RA1EC	ESTs, Highly similar to CYCLIN-DEPENDENT KINASES REGULATORY SUBUNIT 2 [Homo sapiens] {IMAGE:316411}	316411 IMAGE Consortium	W13152
6913 B	2 021YA1ED	Kallikrein-3, plasma {IMAGE:329866}	329866 IMAGE Consortium	W14138
6914 B	4 021YA1ED	Mus musculus non-muscle myosin light chain 3 (MLC3nm) mRNA, partial cds {IMAGE:330370}	330370 IMAGE Consortium	W14559
6915 B	6 021YA1ED	IMAGE EST {IMAGE:316805}	316805 IMAGE Consortium	W11325
6916 B	8 021YA1ED	ESTs, Weakly similar to cDNA EST EMBL:T01157 comes from this gene [C.elegans] {IMAGE:313588}	313588 IMAGE Consortium	W10174
6917 B	10 021YA1ED	ESTs, Highly similar to SARCOPLASMIC RETICULUM HISTIDINE-RICH CALCIUM-BINDING PROTEIN PRECURSOR [Homo sapiens] {IMAGE:317769}	317769 IMAGE Consortium	W13030
6918 B	12 021YA1ED	ESTs {IMAGE:317828}	317828 IMAGE Consortium	W13059
6919 D	2 021YA1ED	IMAGE EST {IMAGE:317268}	317268 IMAGE Consortium	W13212
6920 D	4 021YA1ED	IMAGE EST {IMAGE:317712}	317712 IMAGE Consortium	W11549
6921 D	6 021YA1ED	ESTs {IMAGE:317662}	317662 IMAGE Consortium	W12778
6922 D	8 021YA1ED	IMAGE EST {IMAGE:314627}	314627 IMAGE Consortium	W10904
6923 D	10 021YA1ED	ESTs, Weakly similar to S100 calcium-binding protein A13 [M.musculus] {IMAGE:318796}	318796 IMAGE Consortium	W34209
6924 D	12 021YA1ED	ESTs, Weakly similar to Ydr531wp [S.cerevisiae] {IMAGE:315704}	315704 IMAGE Consortium	W09970
6925 F	2 021YA1ED	IMAGE EST {IMAGE:314509}	314509 IMAGE Consortium	W10776
6926 F	4 021YA1ED	Synaptotagmin 4 {IMAGE:329565}	329565 IMAGE Consortium	W42224
6927 F	6 021YA1ED	ESTs {IMAGE:316122}	316122 IMAGE Consortium	W12606
6928 F	8 021YA1ED	IMAGE EST {IMAGE:316405}	316405 IMAGE Consortium	W11146
6929 F	10 021YA1ED	IMAGE EST {IMAGE:330620}	330620 IMAGE Consortium	W14275
6930 F	12 021YA1ED	IMAGE EST {IMAGE:317487}	317487 IMAGE Consortium	W13561
6931 H	2 021YA1ED	IMAGE EST {IMAGE:317735}	317735 IMAGE Consortium	W13012

6932 H	4 021YA1ED	ESTs, Moderately similar to CYSTEINYL-TRNA SYNTHETASE [H.sapiens] {IMAGE:318129}	318129 IMAGE Consortium	W11649
6933 H	6 021YA1ED	Malate dehydrogenase, soluble {IMAGE:318346}	318346 IMAGE Consortium	W13686
6934 H	8 021YA1ED	ESTs, Highly similar to UDP-GLUCOSE 4-EPIMERASE [Rattus norvegicus] {IMAGE:318810}	318810 IMAGE Consortium	W34357
6935 H	10 021YA1ED	ESTs, Highly similar to CITRATE SYNTHASE, MITOCHONDRIAL PRECURSOR [Sus scrofa] {IMAGE:329884}	329884 IMAGE Consortium	W14146
6936 H	12 021YA1ED	Cadherin 3 {IMAGE:329780}	329780 IMAGE Consortium	W12889
6937 B	2 0215A1EE	IMAGE EST {IMAGE:336463}	336463 IMAGE Consortium	W18366
6938 B	4 0215A1EE	IMAGE EST {IMAGE:336542}	336542 IMAGE Consortium	W18521
6939 B	6 0215A1EE	ESTs {IMAGE:333063}	333063 IMAGE Consortium	W15566
6940 B	8 0215A1EE	ESTs, Weakly similar to PUTATIVE NICOTINATE-NUCLEOTIDE PYROPHOSPHORYLASE [Saccharomyces cerevisiae] {IMAGE:334067}	334067 IMAGE Consortium	W16187
6941 B	10 0215A1EE	TALIN {IMAGE:334872}	334872 IMAGE Consortium	W17813
6942 B	12 0215A1EE	IMAGE EST {IMAGE:335101}	335101 IMAGE Consortium	W17959
6943 D	2 0215A1EE	IMAGE EST {IMAGE:335017}	335017 IMAGE Consortium	W17919
6944 D	4 0215A1EE	Acid beta glucosidase {IMAGE:331646}	331646 IMAGE Consortium	W08086
6945 D	6 0215A1EE	Paraoxonase 1 {IMAGE:335883}	335883 IMAGE Consortium	W17967
6946 D	8 0215A1EE	ESTs, Highly similar to HYPOTHETICAL 64.5 KD PROTEIN ZK652.9 IN CHROMOSOME III [Caenorhabditis elegans] {IMAGE:336042}	336042 IMAGE Consortium	W36403
6947 D	10 0215A1EE	ESTs, Highly similar to DNA POLYMERASE BETA [Rattus norvegicus] {IMAGE:336464}	336464 IMAGE Consortium	W18497
6948 D	12 0215A1EE	IMAGE EST {IMAGE:333117}	333117 IMAGE Consortium	W15847
6949 F	2 0215A1EE	Natriuretic peptide precursor type A {IMAGE:331803}	331803 IMAGE Consortium	W14325
6950 F	4 0215A1EE	Mus musculus carbonic anhydrase IV gene, complete cds {IMAGE:332285}	332285 IMAGE Consortium	W08694
6951 F	6 0215A1EE	ESTs {IMAGE:332325}	332325 IMAGE Consortium	W08451
6952 F	8 0215A1EE	IMAGE EST {IMAGE:332601}	332601 IMAGE Consortium	W08512
6953 F	10 0215A1EE	IMAGE EST {IMAGE:336673}	336673 IMAGE Consortium	W36801
6954 F	12 0215A1EE	ESTs, Weakly similar to F15G9.4 [C.elegans] {IMAGE:337206}	337206 IMAGE Consortium	W36861
6955 H	2 0215A1EE	ESTs, Moderately similar to motor protein [H.sapiens] {IMAGE:336021}	336021 IMAGE Consortium	W36382
6956 H	4 0215A1EE	ESTs, Weakly similar to S100 calcium-binding protein A13 [M.musculus] {IMAGE:336593}	336593 IMAGE Consortium	W20659
6957 H	6 0215A1EE	M.musculus MMLP gene for muscle LIM protein {IMAGE:332950}	332950 IMAGE Consortium	W08774
6958 H	8 0215A1EE	IMAGE EST {IMAGE:333943}	333943 IMAGE Consortium	W16132
6959 H	10 0215A1EE	IMAGE EST {IMAGE:337567}	337567 IMAGE Consortium	W29300
6960 H	12 0215A1EE	IMAGE EST {IMAGE:334347}	334347 IMAGE Consortium	W37018
6961 B	2 021CA1EF	Mus musculus heterogenous nuclear ribonucleoprotein A2/B1	315740 IMAGE Consortium	W12375

		(hnRNP A2/B1) mRNA, complete cds {IMAGE:315740}		
6962 B	4 021CA1EF	ESTs {IMAGE:316112}	316112 IMAGE Consortium	W12256
6963 B	6 021CA1EF	ESTs {IMAGE:316779}	316779 IMAGE Consortium	W11312
6964 B	8 021CA1EF	ESTs {IMAGE:330910}	330910 IMAGE Consortium	W13947
6965 B	10 021CA1EF	ESTs, Highly similar to DESTRAIN [Homo sapiens; Sus scrofa] {IMAGE:335112}	335112 IMAGE Consortium	W17549
6966 B	12 021CA1EF	Transthyretin {IMAGE:335319}	335319 IMAGE Consortium	W17647
6967 D	2 021CA1EF	IMAGE EST {IMAGE:348028}	348028 IMAGE Consortium	W29784
6968 D	4 021CA1EF	ESTs {IMAGE:331399}	331399 IMAGE Consortium	W14955
6969 D	6 021CA1EF	Excision repair 1 {IMAGE:331884}	331884 IMAGE Consortium	W08125
6970 D	8 021CA1EF	ESTs, Highly similar to ERYTHROCYTE ADDUCIN BETA SUBUNIT [Homo sapiens] {IMAGE:336181}	336181 IMAGE Consortium	W18074
6971 D	10 021CA1EF	IMAGE EST {IMAGE:332667}	332667 IMAGE Consortium	W08748
6972 D	12 021CA1EF	IMAGE EST {IMAGE:332534}	332534 IMAGE Consortium	W08556
6973 F	2 021CA1EF	ESTs {IMAGE:335851}	335851 IMAGE Consortium	W34306
6974 F	4 021CA1EF	IMAGE EST {IMAGE:336509}	336509 IMAGE Consortium	W18397
6975 F	6 021CA1EF	ESTs {IMAGE:333214}	333214 IMAGE Consortium	W18229
6976 F	8 021CA1EF	GLUCOSE-6-PHOSPHATE ISOMERASE {IMAGE:337413}	337413 IMAGE Consortium	W29397
6977 F	10 021CA1EF	ESTs {IMAGE:337572}	337572 IMAGE Consortium	W29432
6978 F	12 021CA1EF	ESTs {IMAGE:334591}	334591 IMAGE Consortium	W18348
6979 H	2 021CA1EF	ESTs {IMAGE:336422}	336422 IMAGE Consortium	W36700
6980 H	4 021CA1EF	IMAGE EST {IMAGE:336310}	336310 IMAGE Consortium	W18449
6981 H	6 021CA1EF	ESTs {IMAGE:336684}	336684 IMAGE Consortium	W36781
6982 H	8 021CA1EF	IMAGE EST {IMAGE:337132}	337132 IMAGE Consortium	W36837
6983 H	10 021CA1EF	IMAGE EST {IMAGE:337240}	337240 IMAGE Consortium	W29156
6984 H	12 021CA1EF	ESTs {IMAGE:337127}	337127 IMAGE Consortium	W20777
6985 B	2 021JA1EG	IMAGE EST {IMAGE:336323}	336323 IMAGE Consortium	AA052208
6986 B	4 021JA1EG	ESTs {IMAGE:332687}	332687 IMAGE Consortium	W08673
6987 B	6 021JA1EG	Mus musculus Bet1p homolog (mbet1) mRNA, complete cds {IMAGE:333232}	333232 IMAGE Consortium	W18376
6988 B	8 021JA1EG	Suppressor of Ty 6 (S. cerevisiae) homolog {IMAGE:333778}	333778 IMAGE Consortium	W15976
6989 B	10 021JA1EG	IMAGE EST {IMAGE:334273}	334273 IMAGE Consortium	AA052391
6990 B	12 021JA1EG	ESTs, Moderately similar to PHAS-II [M.musculus] {IMAGE:348098}	348098 IMAGE Consortium	W18851
6991 D	2 021JA1EG	IMAGE EST {IMAGE:337282}	337282 IMAGE Consortium	W20942
6992 D	4 021JA1EG	Mus musculus sulfotransferase-related protein (SULT-X1) mRNA, complete cds {IMAGE:334058}	334058 IMAGE Consortium	W16116
6993 D	6 021JA1EG	ESTs {IMAGE:334002}	334002 IMAGE Consortium	W17528
6994 D	8 021JA1EG	M.musculus mRNA for s17 protein {IMAGE:348123}	348123 IMAGE Consortium	W18922
6995 D	10 021JA1EG	ESTs, Highly similar to FIBRINOGEN BETA CHAIN PRECURSOR [Homo sapiens] {IMAGE:348037}	348037 IMAGE Consortium	W29662
6996 D	12 021JA1EG	Cytochrome P450, 2e1, ethanol inducible {IMAGE:331386}	331386 IMAGE Consortium	W14813
6997 F	2 021JA1EG	B-cell translocation gene 2, anti-proliferative {IMAGE:334106}	334106 IMAGE Consortium	W16205

6998 F	4 021JA1EG	IMAGE EST {IMAGE:337957}	337957 IMAGE Consortium	W18886
6999 F	6 021JA1EG	ESTs, Highly similar to TYROSINE-PROTEIN KINASE JAK1 [Homo sapiens] {IMAGE:347921}	347921 IMAGE Consortium	W29699
7000 F	8 021JA1EG	ESTs, Moderately similar to SAS [H.sapiens] {IMAGE:348438}	348438 IMAGE Consortium	W33323
7001 F	10 021JA1EG	ESTs, Highly similar to NADH-UBIQUINONE OXIDOREDUCTASE B12 SUBUNIT [Bos taurus] {IMAGE:352579}	352579 IMAGE Consortium	W34455
7002 F	12 021JA1EG	IMAGE EST {IMAGE:353032}	353032 IMAGE Consortium	W34775
7003 H	2 021JA1EG	IMAGE EST {IMAGE:354101}	354101 IMAGE Consortium	AA052574
7004 H	4 021JA1EG	ESTs, Highly similar to GLUTAMINASE, LIVER ISOFORM PRECURSOR [Rattus norvegicus] {IMAGE:351733}	351733 IMAGE Consortium	W41254
7005 H	6 021JA1EG	IMAGE EST {IMAGE:352866}	352866 IMAGE Consortium	W36046
7006 H	8 021JA1EG	ESTs {IMAGE:350182}	350182 IMAGE Consortium	W34722
7007 H	10 021JA1EG	IMAGE EST {IMAGE:354238}	354238 IMAGE Consortium	W44162
7008 H	12 021JA1EG	IMAGE EST {IMAGE:351810}	351810 IMAGE Consortium	W41250
7009 B	2 021QA1EH	IMAGE EST {IMAGE:354575}	354575 IMAGE Consortium	W46016
7010 B	4 021QA1EH	IMAGE EST {IMAGE:352406}	352406 IMAGE Consortium	W33774
7011 B	6 021QA1EH	IMAGE EST {IMAGE:353094}	353094 IMAGE Consortium	W34856
7012 B	8 021QA1EH	ESTs, Highly similar to C-PROTEIN, SKELETAL MUSCLE SLOW-ISOFORM [Homo sapiens] {IMAGE:350088}	350088 IMAGE Consortium	W34541
7013 B	10 021QA1EH	IMAGE EST {IMAGE:350254}	350254 IMAGE Consortium	W34670
7014 B	12 021QA1EH	ESTs {IMAGE:351039}	351039 IMAGE Consortium	W35079
7015 D	2 021QA1EH	IMAGE EST {IMAGE:350304}	350304 IMAGE Consortium	W34682
7016 D	4 021QA1EH	IMAGE EST {IMAGE:350245}	350245 IMAGE Consortium	W35945
7017 D	6 021QA1EH	IMAGE EST {IMAGE:350327}	350327 IMAGE Consortium	W40724
7018 D	8 021QA1EH	ESTs, Weakly similar to PACSIN [M.musculus] {IMAGE:354625}	354625 IMAGE Consortium	W46097
7019 D	10 021QA1EH	IMAGE EST {IMAGE:351844}	351844 IMAGE Consortium	W40590
7020 D	12 021QA1EH	ESTs, Highly similar to ATP-CITRATE [Rattus norvegicus] {IMAGE:352146}	352146 IMAGE Consortium	W33415
7021 F	2 021QA1EH	ESTs {IMAGE:352943}	352943 IMAGE Consortium	W34927
7022 F	4 021QA1EH	Coagulation factor II {IMAGE:350344}	350344 IMAGE Consortium	W40725
7023 F	6 021QA1EH	ESTs, Moderately similar to The ha3523 gene product is related to S.cerevisiae gene product located in chromosome III. [H.sapiens] {IMAGE:354181}	354181 IMAGE Consortium	AA052369
7024 F	8 021QA1EH	ESTs {IMAGE:352219}	352219 IMAGE Consortium	W36074
7025 F	10 021QA1EH	ESTs {IMAGE:352481}	352481 IMAGE Consortium	W33853
7026 F	12 021QA1EH	Mus musculus endophilin I mRNA, complete cds {IMAGE:352836}	352836 IMAGE Consortium	W34672
7027 H	2 021QA1EH	RAR-related orphan receptor alpha {IMAGE:350310}	350310 IMAGE Consortium	W34685
7028 H	4 021QA1EH	IMAGE EST {IMAGE:350525}	350525 IMAGE Consortium	W35023
7029 H	6 021QA1EH	IMAGE EST {IMAGE:350959}	350959 IMAGE Consortium	AA052449



7030 H	8 021QA1EH	Mus musculus eIF-1A (eIF-1A) mRNA, complete cds {IMAGE:351631}	351631 IMAGE Consortium	W41459
7031 H	10 021QA1EH	IMAGE EST {IMAGE:352090}	352090 IMAGE Consortium	W33456
7032 H	12 021QA1EH	IMAGE EST {IMAGE:352909}	352909 IMAGE Consortium	W34831
7033 B	2 021XA1EI	Heparan sulfate glucosaminyl 3-O-sulfotransferase {IMAGE:373942}	373942 IMAGE Consortium	W62484
7034 B	4 021XA1EI	ESTs, Highly similar to EOSINOPHIL GRANULE MAJOR BASIC PROTEIN PRECURSOR [Homo sapiens] {IMAGE:354861}	354861 IMAGE Consortium	W45834
7035 B	6 021XA1EI	ESTs, Highly similar to p16-Arc [H.sapiens] {IMAGE:355211}	355211 IMAGE Consortium	W47753
7036 B	8 021XA1EI	IMAGE EST {IMAGE:355582}	355582 IMAGE Consortium	W48894
7037 B	10 021XA1EI	ESTs {IMAGE:372368}	372368 IMAGE Consortium	W53544
7038 B	12 021XA1EI	ESTs {IMAGE:367514}	367514 IMAGE Consortium	W54611
7039 D	2 021XA1EI	ESTs {IMAGE:373505}	373505 IMAGE Consortium	W59073
7040 D	4 021XA1EI	ESTs, Weakly similar to HIRA protein [M.musculus] {IMAGE:373854}	373854 IMAGE Consortium	W64706
7041 D	6 021XA1EI	IMAGE EST {IMAGE:374041}	374041 IMAGE Consortium	W59657
7042 D	8 021XA1EI	GUANINE NUCLEOTIDE DISSOCIATION STIMULATOR RALGDSA {IMAGE:355112}	355112 IMAGE Consortium	W48230
7043 D	10 021XA1EI	ESTs {IMAGE:367014}	367014 IMAGE Consortium	W54365
7044 D	12 021XA1EI	ESTs {IMAGE:367540}	367540 IMAGE Consortium	W54620
7045 F	2 021XA1EI	Hexokinase 1 {IMAGE:389177}	389177 IMAGE Consortium	W74835
7046 F	4 021XA1EI	IMAGE EST {IMAGE:389324}	389324 IMAGE Consortium	W71347
7047 F	6 021XA1EI	ESTs {IMAGE:386218}	386218 IMAGE Consortium	W65070
7048 F	8 021XA1EI	ESTs, Highly similar to INTESTINAL MEMBRANE A4 PROTEIN [Homo sapiens] {IMAGE:386254}	386254 IMAGE Consortium	W64735
7049 F	10 021XA1EI	ESTs {IMAGE:386682}	386682 IMAGE Consortium	W64844
7050 F	12 021XA1EI	ESTs {IMAGE:387200}	387200 IMAGE Consortium	W65895
7051 H	2 021XA1EI	IMAGE EST {IMAGE:387395}	387395 IMAGE Consortium	W65869
7052 H	4 021XA1EI	IMAGE EST {IMAGE:385039}	385039 IMAGE Consortium	W64745
7053 H	6 021XA1EI	ESTs, Highly similar to YSA1 PROTEIN [Saccharomyces cerevisiae] {IMAGE:385358}	385358 IMAGE Consortium	W70837
7054 H	8 021XA1EI	Mus musculus MAST205 protein kinase mRNA, complete cds {IMAGE:385690}	385690 IMAGE Consortium	W61408
7055 H	10 021XA1EI	IMAGE EST {IMAGE:389034}	389034 IMAGE Consortium	W67049
7056 H	12 021XA1EI	NUCLEOBINDIN PRECURSOR {IMAGE:390016}	390016 IMAGE Consortium	W71629
7057 B	2 0214A1EJ	Tumor necrosis factor receptor superfamily, member 1b {IMAGE:354773}	354773 IMAGE Consortium	W45975
7058 B	4 0214A1EJ	IMAGE EST {IMAGE:356215}	356215 IMAGE Consortium	W50731
7059 B	6 0214A1EJ	IMAGE EST {IMAGE:366992}	366992 IMAGE Consortium	W54295
7060 B	8 0214A1EJ	IMAGE EST {IMAGE:372939}	372939 IMAGE Consortium	W59584
7061 B	10 0214A1EJ	Myeloperoxidase {IMAGE:367693}	367693 IMAGE Consortium	W55655
7062 B	12 0214A1EJ	ESTs {IMAGE:368026}	368026 IMAGE Consortium	W53778

7063 D	2 0214A1EJ	IMAGE EST {IMAGE:355425}	355425 IMAGE Consortium	W48167
7064 D	4 0214A1EJ	IMAGE EST {IMAGE:355939}	355939 IMAGE Consortium	W49208
7065 D	6 0214A1EJ	ESTs {IMAGE:367329}	367329 IMAGE Consortium	W53944
7066 D	8 0214A1EJ	ESTs {IMAGE:373036}	373036 IMAGE Consortium	W62448
7067 D	10 0214A1EJ	ESTs {IMAGE:373489}	373489 IMAGE Consortium	W57174
7068 D	12 0214A1EJ	IMAGE EST {IMAGE:373596}	373596 IMAGE Consortium	W57258
7069 F	2 0214A1EJ	M.musculus trop2 gene {IMAGE:373716}	373716 IMAGE Consortium	W64550
7070 F	4 0214A1EJ	ESTs {IMAGE:368189}	368189 IMAGE Consortium	W53913
7071 F	6 0214A1EJ	ESTs {IMAGE:355138}	355138 IMAGE Consortium	W48262
7072 F	8 0214A1EJ	RIBOSOMAL PROTEIN S6 KINASE II ALPHA 1 {IMAGE:355755}	355755 IMAGE Consortium	W47785
7073 F	10 0214A1EJ	DiGeorge syndrome chromosome region 6 {IMAGE:356181}	356181 IMAGE Consortium	W54234
7074 F	12 0214A1EJ	IMAGE EST {IMAGE:367802}	367802 IMAGE Consortium	W53994
7075 H	2 0214A1EJ	ESTs, Moderately similar to hypothetical protein [M.musculus] {IMAGE:355426}	355426 IMAGE Consortium	W48161
7076 H	4 0214A1EJ	ESTs {IMAGE:355517}	355517 IMAGE Consortium	W47974
7077 H	6 0214A1EJ	ESTs {IMAGE:356159}	356159 IMAGE Consortium	W54229
7078 H	8 0214A1EJ	IMAGE EST {IMAGE:367333}	367333 IMAGE Consortium	W53946
7079 H	10 0214A1EJ	ESTs {IMAGE:373305}	373305 IMAGE Consortium	W62436
7080 H	12 0214A1EJ	Mus musculus DTEF-1 mRNA, complete cds {IMAGE:367947}	367947 IMAGE Consortium	W54739
7081 B	2 021BA1EK	IMAGE EST {IMAGE:388217}	388217 IMAGE Consortium	W66600
7082 B	4 021BA1EK	ESTs {IMAGE:386112}	386112 IMAGE Consortium	W62198
7083 B	6 021BA1EK	ESTs {IMAGE:386100}	386100 IMAGE Consortium	W62403
7084 B	8 021BA1EK	Mus musculus mRNA for protein L, partial cds {IMAGE:386234}	386234 IMAGE Consortium	W65080
7085 B	10 021BA1EK	IMAGE EST {IMAGE:386533}	386533 IMAGE Consortium	W64997
7086 B	12 021BA1EK	IMAGE EST {IMAGE:390303}	390303 IMAGE Consortium	W70775
7087 D	2 021BA1EK	ESTs {IMAGE:386101}	386101 IMAGE Consortium	W62196
7088 D	4 021BA1EK	IMAGE EST {IMAGE:386198}	386198 IMAGE Consortium	W65056
7089 D	6 021BA1EK	Mus musculus SHYC (Shyc) mRNA, complete cds {IMAGE:386790}	386790 IMAGE Consortium	W64430
7090 D	8 021BA1EK	ESTs, Highly similar to SODIUM/POTASSIUM- TRANSPORTING ATPASE ALPHA-1 CHAIN [Rattus norvegicus] {IMAGE:374487}	374487 IMAGE Consortium	W65246
7091 D	10 021BA1EK	ESTs, Weakly similar to similar to Zinc finger, C3HC4 type [C.elegans] {IMAGE:387446}	387446 IMAGE Consortium	W65834
7092 D	12 021BA1EK	ESTs {IMAGE:388330}	388330 IMAGE Consortium	W70906
7093 F	2 021BA1EK	IMAGE EST {IMAGE:387408}	387408 IMAGE Consortium	W65872
7094 F	4 021BA1EK	Cytochrome P450, 2e1, ethanol inducible {IMAGE:388027}	388027 IMAGE Consortium	W66885
7095 F	6 021BA1EK	ESTs {IMAGE:388212}	388212 IMAGE Consortium	W67062
7096 F	8 021BA1EK	ESTs, Highly similar to BRAIN PROTEIN H5 [Mus musculus] {IMAGE:385729}	385729 IMAGE Consortium	W62231

7097 F	10 021BA1EK	ESTs {IMAGE:385525}	385525 IMAGE Consortium	W61449
7098 F	12 021BA1EK	ESTs {IMAGE:385853}	385853 IMAGE Consortium	W62295
7099 H	2 021BA1EK	Mus musculus rho7 (rho7) mRNA, partial cds {IMAGE:385746}	385746 IMAGE Consortium	W64242
7100 H	4 021BA1EK	IMAGE EST {IMAGE:389580}	389580 IMAGE Consortium	W71435
7101 H	6 021BA1EK	Mus musculus RNaseP protein p30 (Rpp30) mRNA, complete cds {IMAGE:389279}	389279 IMAGE Consortium	W71337
7102 H	8 021BA1EK	IMAGE EST {IMAGE:390118}	390118 IMAGE Consortium	W70355
7103 H	10 021BA1EK	IMAGE EST {IMAGE:387291}	387291 IMAGE Consortium	W65810
7104 H	12 021BA1EK	IMAGE EST {IMAGE:387227}	387227 IMAGE Consortium	W65830
7105 B	2 021IA1EL	IMAGE EST {IMAGE:385986}	385986 IMAGE Consortium	W64276
7106 B	4 021IA1EL	Mus musculus mRNA for Ftp-1, complete cds {IMAGE:386456}	386456 IMAGE Consortium	W64415
7107 B	6 021IA1EL	IMAGE EST {IMAGE:390433}	390433 IMAGE Consortium	W71838
7108 B	8 021IA1EL	Growth factor independent 1B {IMAGE:387315}	387315 IMAGE Consortium	W65842
7109 B	10 021IA1EL	ESTs {IMAGE:387594}	387594 IMAGE Consortium	W66845
7110 B	12 021IA1EL	Regulatory factor (trans-acting) 2 {IMAGE:388321}	388321 IMAGE Consortium	W70909
7111 D	2 021IA1EL	IMAGE EST {IMAGE:387459}	387459 IMAGE Consortium	W65788
7112 D	4 021IA1EL	IMAGE EST {IMAGE:388394}	388394 IMAGE Consortium	W70958
7113 D	6 021IA1EL	Epimorphin {IMAGE:385220}	385220 IMAGE Consortium	W64796
7114 D	8 021IA1EL	IMAGE EST {IMAGE:385723}	385723 IMAGE Consortium	W64238
7115 D	10 021IA1EL	ESTs {IMAGE:386142}	386142 IMAGE Consortium	W62404
7116 D	12 021IA1EL	IMAGE EST {IMAGE:386300}	386300 IMAGE Consortium	W64920
7117 F	2 021IA1EL	ESTs, Moderately similar to CYANELLE 30S RIBOSOMAL PROTEIN S11 [Cyanophora paradoxa] {IMAGE:386872}	386872 IMAGE Consortium	W65559
7118 F	4 021IA1EL	Myosin Va {IMAGE:402475}	402475 IMAGE Consortium	W80031
7119 F	6 021IA1EL	ESTs {IMAGE:402923}	402923 IMAGE Consortium	W80210
7120 F	8 021IA1EL	House mouse; Musculus domesticus mRNA for LN1, complete cds {IMAGE:403728}	403728 IMAGE Consortium	W82212
7121 F	10 021IA1EL	Mus musculus Sil mRNA, complete cds {IMAGE:400735}	400735 IMAGE Consortium	W75881
7122 F	12 021IA1EL	IMAGE EST {IMAGE:400530}	400530 IMAGE Consortium	W76728
7123 H	2 021IA1EL	ESTs {IMAGE:403681}	403681 IMAGE Consortium	W82203
7124 H	4 021IA1EL	IMAGE EST {IMAGE:400592}	400592 IMAGE Consortium	W75803
7125 H	6 021IA1EL	Mouse small nuclear RNA (Rnu1a-1) mRNA, complete cds {IMAGE:400554}	400554 IMAGE Consortium	W76735
7126 H	8 021IA1EL	IMAGE EST {IMAGE:401167}	401167 IMAGE Consortium	W98060
7127 H	10 021IA1EL	Keratin complex 2, basic, gene 4 {IMAGE:401327}	401327 IMAGE Consortium	W98341
7128 H	12 021IA1EL	ESTs {IMAGE:401820}	401820 IMAGE Consortium	W78555
7129 B	2 021PA1EM	ESTs {IMAGE:402800}	402800 IMAGE Consortium	W80005
7130 B	4 021PA1EM	ESTs {IMAGE:402934}	402934 IMAGE Consortium	W80302
7131 B	6 021PA1EM	ESTs {IMAGE:403413}	403413 IMAGE Consortium	W82905
7132 B	8 021PA1EM	ESTs, Highly similar to HYPOTHETICAL 43.3 KD PROTEIN K06H7.5 IN CHROMOSOME III [Caenorhabditis elegans]	400613 IMAGE Consortium	W75669

		{IMAGE:400613}		
7133 B	10 021PA1EM	ESTs {IMAGE:400408}	400408 IMAGE Consortium	W77278
7134 B	12 021PA1EM	ESTs, Highly similar to NADH-UBIQUINONE OXIDOREDUCTASE ASHI SUBUNIT PRECURSOR [Bos taurus] {IMAGE:404593}	404593 IMAGE Consortium	W83085
7135 D	2 021PA1EM	M.musculus mRNA for NADH dehydrogenase {IMAGE:404499}	404499 IMAGE Consortium	W83104
7136 D	4 021PA1EM	ESTs, Weakly similar to ENDOSOMAL P24A PROTEIN PRECURSOR [Saccharomyces cerevisiae] {IMAGE:404535}	404535 IMAGE Consortium	W83187
7137 D	6 021PA1EM	Dlk1-like homolog (Drosophila) {IMAGE:402244}	402244 IMAGE Consortium	W76877
7138 D	8 021PA1EM	IMAGE EST {IMAGE:402162}	402162 IMAGE Consortium	W76849
7139 D	10 021PA1EM	ESTs {IMAGE:403034}	403034 IMAGE Consortium	W81908
7140 D	12 021PA1EM	IMAGE EST {IMAGE:403072}	403072 IMAGE Consortium	W81873
7141 F	2 021PA1EM	IMAGE EST {IMAGE:403274}	403274 IMAGE Consortium	W82668
7142 F	4 021PA1EM	Mus musculus c-Cbl associated protein CAP mRNA, complete cds {IMAGE:403656}	403656 IMAGE Consortium	W82192
7143 F	6 021PA1EM	IMAGE EST {IMAGE:400495}	400495 IMAGE Consortium	W77454
7144 F	8 021PA1EM	IMAGE EST {IMAGE:404165}	404165 IMAGE Consortium	W82472
7145 F	10 021PA1EM	ESTs, Moderately similar to similar to the DNA/RNA non-specific endoneuclease family [C.elegans] {IMAGE:404597}	404597 IMAGE Consortium	W83087
7146 F	12 021PA1EM	Tuberous sclerosis 2 {IMAGE:404321}	404321 IMAGE Consortium	W82751
7147 H	2 021PA1EM	IMAGE EST {IMAGE:408898}	408898 IMAGE Consortium	W85623
7148 H	4 021PA1EM	ESTs {IMAGE:408934}	408934 IMAGE Consortium	W85540
7149 H	6 021PA1EM	IMAGE EST {IMAGE:418548}	418548 IMAGE Consortium	W99118
7150 H	8 021PA1EM	ESTs {IMAGE:405779}	405779 IMAGE Consortium	W83025
7151 H	10 021PA1EM	ESTs, Weakly similar to predicted using Genefinder [C.elegans] {IMAGE:406476}	406476 IMAGE Consortium	W83499
7152 H	12 021PA1EM	ESTs {IMAGE:419086}	419086 IMAGE Consortium	W89966
7153 B	2 021WA1EN	ESTs {IMAGE:404169}	404169 IMAGE Consortium	W82521
7154 B	4 021WA1EN	ESTs {IMAGE:404057}	404057 IMAGE Consortium	W82577
7155 B	6 021WA1EN	ESTs {IMAGE:401557}	401557 IMAGE Consortium	W78448
7156 B	8 021WA1EN	Mus musculus small GTPase Rab10 (Rab10) mRNA, complete cds {IMAGE:401878}	401878 IMAGE Consortium	W82973
7157 B	10 021WA1EN	IMAGE EST {IMAGE:402080}	402080 IMAGE Consortium	W75911
7158 B	12 021WA1EN	ESTs, Moderately similar to PROBABLE PEPTIDYL-PROLYL CIS-TRANS ISOMERASE C21E11.05C [Schizosaccharomyces pombe] {IMAGE:402748}	402748 IMAGE Consortium	W80072
7159 D	2 021WA1EN	Secretogranin III {IMAGE:402105}	402105 IMAGE Consortium	W75931
7160 D	4 021WA1EN	ESTs {IMAGE:402429}	402429 IMAGE Consortium	W79991
7161 D	6 021WA1EN	IMAGE EST {IMAGE:402979}	402979 IMAGE Consortium	W80316
7162 D	8 021WA1EN	ESTs, Moderately similar to ATP SYNTHASE F CHAIN, MITOCHONDRIAL [M.musculus] {IMAGE:403660}	403660 IMAGE Consortium	W82194
7163 D	10 021WA1EN	IMAGE EST {IMAGE:403525}	403525 IMAGE Consortium	W82183

7164 D	12 021WA1EN	ESTs {IMAGE:401288}	401288 IMAGE Consortium	W98234
7165 F	2 021WA1EN	IMAGE EST {IMAGE:400500}	400500 IMAGE Consortium	W77481
7166 F	4 021WA1EN	ESTs, Weakly similar to predicted using Genefinder [C.elegans] {IMAGE:403932}	403932 IMAGE Consortium	W82174
7167 F	6 021WA1EN	ESTs, Highly similar to BETA CRYSTALLIN A4 [Bos taurus] {IMAGE:403869}	403869 IMAGE Consortium	W82104
7168 F	8 021WA1EN	ESTs, Moderately similar to HYPOTHETICAL TRP-ASP REPEATS CONTAINING PROTEIN IN CPR4-SOL2 INTERGENIC REGION [Saccharomyces cerevisiae] {IMAGE:404268}	404268 IMAGE Consortium	W82627
7169 F	10 021WA1EN	ESTs {IMAGE:404589}	404589 IMAGE Consortium	W82894
7170 F	12 021WA1EN	Galanin {IMAGE:405989}	405989 IMAGE Consortium	W83304
7171 H	2 021WA1EN	IMAGE EST {IMAGE:404613}	404613 IMAGE Consortium	W82945
7172 H	4 021WA1EN	ESTs {IMAGE:401966}	401966 IMAGE Consortium	W77499
7173 H	6 021WA1EN	IMAGE EST {IMAGE:402696}	402696 IMAGE Consortium	W80058
7174 H	8 021WA1EN	IMAGE EST {IMAGE:402814}	402814 IMAGE Consortium	W80088
7175 H	10 021WA1EN	IMAGE EST {IMAGE:403220}	403220 IMAGE Consortium	W82016
7176 H	12 021WA1EN	ESTs {IMAGE:403663}	403663 IMAGE Consortium	W82199
7177 B	2 0213A1EO	ESTs {IMAGE:420131}	420131 IMAGE Consortium	W89800
7178 B	4 0213A1EO	IMAGE EST {IMAGE:420015}	420015 IMAGE Consortium	W91259
7179 B	6 0213A1EO	ESTs, Highly similar to INTEGRAL MEMBRANE GLYCOPROTEIN GP210 PRECURSOR [Rattus norvegicus] {IMAGE:418576}	418576 IMAGE Consortium	W99187
7180 B	8 0213A1EO	ESTs {IMAGE:418462}	418462 IMAGE Consortium	W87055
7181 B	10 0213A1EO	ESTs {IMAGE:406609}	406609 IMAGE Consortium	W83605
7182 B	12 0213A1EO	Mouse binding protein for T-cell receptor (TCR-ATF1) mRNA, complete cds {IMAGE:419424}	419424 IMAGE Consortium	W87965
7183 D	2 0213A1EO	ESTs, Moderately similar to p53BP2 [M.musculus] {IMAGE:418440}	418440 IMAGE Consortium	W85569
7184 D	4 0213A1EO	IMAGE EST {IMAGE:418495}	418495 IMAGE Consortium	W85526
7185 D	6 0213A1EO	Calcium channel beta 3 subunit {IMAGE:420487}	420487 IMAGE Consortium	W90827
7186 D	8 0213A1EO	IMAGE EST {IMAGE:418582}	418582 IMAGE Consortium	W99190
7187 D	10 0213A1EO	Mus musculus RaIBP1-associated EH domain protein Reps1 (reps1) mRNA, complete cds {IMAGE:418926}	418926 IMAGE Consortium	AA052176
7188 D	12 0213A1EO	IMAGE EST {IMAGE:418861}	418861 IMAGE Consortium	W89337
7189 F	2 0213A1EO	Plasminogen {IMAGE:418537}	418537 IMAGE Consortium	W99028
7190 F	4 0213A1EO	ESTs {IMAGE:418453}	418453 IMAGE Consortium	W87014
7191 F	6 0213A1EO	ESTs, Highly similar to HYPOTHETICAL 34.7 KD PROTEIN IN SPT10-GCD14 INTERGENIC REGION [Saccharomyces cerevisiae] {IMAGE:406154}	406154 IMAGE Consortium	W83856
7192 F	8 0213A1EO	IMAGE EST {IMAGE:419070}	419070 IMAGE Consortium	W89354
7193 F	10 0213A1EO	ESTs {IMAGE:419030}	419030 IMAGE Consortium	W89400
7194 F	12 0213A1EO	IMAGE EST {IMAGE:419544}	419544 IMAGE Consortium	W88154

7195 H	2 0213A1EO	Lysyl oxidase {IMAGE:418645}	418645 IMAGE Consortium	W96914
7196 H	4 0213A1EO	ESTs {IMAGE:405902}	405902 IMAGE Consortium	W83057
7197 H	6 0213A1EO	IMAGE EST {IMAGE:406218}	406218 IMAGE Consortium	W84060
7198 H	8 0213A1EO	ESTs {IMAGE:406708}	406708 IMAGE Consortium	W83671
7199 H	10 0213A1EO	House mouse; Musculus domesticus mRNA for membrane glycoprotein, complete cds {IMAGE:406913}	406913 IMAGE Consortium	W83922
7200 H	12 0213A1EO	IMAGE EST {IMAGE:419580}	419580 IMAGE Consortium	AA052081
7201 B	2 021AA1EP	IMAGE EST {IMAGE:418473}	418473 IMAGE Consortium	W87034
7202 B	4 021AA1EP	IMAGE EST {IMAGE:420727}	420727 IMAGE Consortium	W89659
7203 B	6 021AA1EP	IMAGE EST {IMAGE:406323}	406323 IMAGE Consortium	W83186
7204 B	8 021AA1EP	Insulin-like growth factor binding protein 5 {IMAGE:418952}	418952 IMAGE Consortium	W89253
7205 B	10 021AA1EP	IMAGE EST {IMAGE:418888}	418888 IMAGE Consortium	W89392
7206 B	12 021AA1EP	IMAGE EST {IMAGE:419237}	419237 IMAGE Consortium	W89623
7207 D	2 021AA1EP	ESTs {IMAGE:418737}	418737 IMAGE Consortium	W97078
7208 D	4 021AA1EP	ESTs, Highly similar to TRANSCRIPTION FACTOR IIIA [ <i>Xenopus laevis</i> ] {IMAGE:418670}	418670 IMAGE Consortium	W96919
7209 D	6 021AA1EP	IMAGE EST {IMAGE:406144}	406144 IMAGE Consortium	W83315
7210 D	8 021AA1EP	ESTs {IMAGE:418535}	418535 IMAGE Consortium	W87122
7211 D	10 021AA1EP	ESTs, Weakly similar to C32A3.3 [ <i>C.elegans</i> ] {IMAGE:406566}	406566 IMAGE Consortium	W83390
7212 D	12 021AA1EP	Survival motor neuron {IMAGE:419573}	419573 IMAGE Consortium	W88179
7213 F	2 021AA1EP	ESTs {IMAGE:405905}	405905 IMAGE Consortium	W83477
7214 F	4 021AA1EP	ESTs, Highly similar to DEAD-box protein p72 [ <i>H.sapiens</i> ] {IMAGE:418920}	418920 IMAGE Consortium	AA052175
7215 F	6 021AA1EP	Mus musculus pericentrin mRNA, complete cds {IMAGE:419050}	419050 IMAGE Consortium	W89405
7216 F	8 021AA1EP	ESTs, Weakly similar to KIAA0349 [ <i>H.sapiens</i> ] {IMAGE:406894}	406894 IMAGE Consortium	W83898
7217 F	10 021AA1EP	Mus musculus Rho-GDI3 mRNA, complete cds {IMAGE:407050}	407050 IMAGE Consortium	W83964
7218 F	12 021AA1EP	ESTs {IMAGE:407354}	407354 IMAGE Consortium	W87102
7219 H	2 021AA1EP	IMAGE EST {IMAGE:419184}	419184 IMAGE Consortium	W87950
7220 H	4 021AA1EP	ESTs {IMAGE:419146}	419146 IMAGE Consortium	W88005
7221 H	6 021AA1EP	GUANINE NUCLEOTIDE-BINDING PROTEIN, ALPHA-13 SUBUNIT {IMAGE:419058}	419058 IMAGE Consortium	W89964
7222 H	8 021AA1EP	ESTs, Weakly similar to KIAA0009 [ <i>H.sapiens</i> ] {IMAGE:406955}	406955 IMAGE Consortium	AA052810
7223 H	10 021AA1EP	ESTs {IMAGE:407428}	407428 IMAGE Consortium	AA052168
7224 H	12 021AA1EP	IMAGE EST {IMAGE:408747}	408747 IMAGE Consortium	W85461
7225 B	2 021HA1EQ	EST {IMAGE:423422}	423422 IMAGE Consortium	W98898
7226 B	4 021HA1EQ	IMAGE EST {IMAGE:420591}	420591 IMAGE Consortium	W89466
7227 B	6 021HA1EQ	ESTs {IMAGE:422403}	422403 IMAGE Consortium	W97732
7228 B	8 021HA1EQ	IMAGE EST {IMAGE:421157}	421157 IMAGE Consortium	AA052083

7229 B	10 021HA1EQ	ESTs {IMAGE:421145}	421145 IMAGE Consortium	W98975
7230 B	12 021HA1EQ	IMAGE EST {IMAGE:423050}	423050 IMAGE Consortium	W98025
7231 D	2 021HA1EQ	ESTs {IMAGE:422959}	422959 IMAGE Consortium	W97983
7232 D	4 021HA1EQ	IMAGE EST {IMAGE:422799}	422799 IMAGE Consortium	W96991
7233 D	6 021HA1EQ	ESTs, Highly similar to C-PROTEIN, SKELETAL MUSCLE SLOW-ISOFORM [Homo sapiens] {IMAGE:421758}	421758 IMAGE Consortium	W98944
7234 D	8 021HA1EQ	IMAGE EST {IMAGE:423179}	423179 IMAGE Consortium	W97304
7235 D	10 021HA1EQ	ESTs {IMAGE:423070}	423070 IMAGE Consortium	W97244
7236 D	12 021HA1EQ	ESTs {IMAGE:423473}	423473 IMAGE Consortium	W98815
7237 F	2 021HA1EQ	Paraoxonase 2 {IMAGE:423266}	423266 IMAGE Consortium	W98586
7238 F	4 021HA1EQ	IMAGE EST {IMAGE:423203}	423203 IMAGE Consortium	W97311
7239 F	6 021HA1EQ	ESTs {IMAGE:423327}	423327 IMAGE Consortium	W98730
7240 F	8 021HA1EQ	IMAGE EST {IMAGE:422162}	422162 IMAGE Consortium	W97381
7241 F	10 021HA1EQ	ESTs, Highly similar to OOCYTE ZINC FINGER PROTEIN XLCOF28 [Xenopus laevis] {IMAGE:423593}	423593 IMAGE Consortium	W98657
7242 F	12 021HA1EQ	IMAGE EST {IMAGE:422254}	422254 IMAGE Consortium	W97621
7243 H	2 021HA1EQ	ESTs, Weakly similar to HYPOTHETICAL 97.1 KD PROTEIN R05D3.4 IN CHROMOSOME III [Caenorhabditis elegans] {IMAGE:422262}	422262 IMAGE Consortium	W97464
7244 H	4 021HA1EQ	IMAGE EST {IMAGE:420544}	420544 IMAGE Consortium	W89695
7245 H	6 021HA1EQ	Mus musculus S6 kinase 2 {IMAGE:421114}	421114 IMAGE Consortium	W91436
7246 H	8 021HA1EQ	ESTs {IMAGE:421129}	421129 IMAGE Consortium	W91029
7247 H	10 021HA1EQ	ESTs {IMAGE:422875}	422875 IMAGE Consortium	W97952
7248 H	12 021HA1EQ	Acid beta glucosidase {IMAGE:421616}	421616 IMAGE Consortium	W99211
7249 B	2 021OA1ER	Mannan-binding lectin serine protease 1 {IMAGE:419455}	419455 IMAGE Consortium	W88086
7250 B	4 021OA1ER	IMAGE EST {IMAGE:407620}	407620 IMAGE Consortium	W89239
7251 B	6 021OA1ER	Telomerase associated protein 1 {IMAGE:407351}	407351 IMAGE Consortium	W87045
7252 B	8 021OA1ER	IMAGE EST {IMAGE:419895}	419895 IMAGE Consortium	W89566
7253 B	10 021OA1ER	IMAGE EST {IMAGE:419755}	419755 IMAGE Consortium	W89552
7254 B	12 021OA1ER	ESTs {IMAGE:420049}	420049 IMAGE Consortium	W91158
7255 D	2 021OA1ER	ESTs {IMAGE:418955}	418955 IMAGE Consortium	W89258
7256 D	4 021OA1ER	ESTs, Moderately similar to the KIAA0199 gene is expressed ubiquitously. [H.sapiens] {IMAGE:407041}	407041 IMAGE Consortium	W83967
7257 D	6 021OA1ER	ESTs {IMAGE:419490}	419490 IMAGE Consortium	W88200
7258 D	8 021OA1ER	ESTs {IMAGE:408233}	408233 IMAGE Consortium	W85553
7259 D	10 021OA1ER	IMAGE EST {IMAGE:420169}	420169 IMAGE Consortium	W89835
7260 D	12 021OA1ER	Mus musculus DNA polymerase gamma mRNA, nuclear gene encoding mitochondrial protein, complete cds {IMAGE:420278}	420278 IMAGE Consortium	W89816
7261 F	2 021OA1ER	Mus musculus unknown mRNA, complete cds {IMAGE:421645}	421645 IMAGE Consortium	W99217
7262 F	4 021OA1ER	IMAGE EST {IMAGE:423210}	423210 IMAGE Consortium	W97334
7263 F	6 021OA1ER	IMAGE EST {IMAGE:422069}	422069 IMAGE Consortium	W96877

7264 F	8 021OA1ER	IMAGE EST {IMAGE:422174}	422174 IMAGE Consortium	W96828
7265 F	10 021OA1ER	IMAGE EST {IMAGE:422393}	422393 IMAGE Consortium	W97514
7266 F	12 021OA1ER	ESTs {IMAGE:423150}	423150 IMAGE Consortium	W97057
7267 H	2 021OA1ER	Cortactin {IMAGE:421324}	421324 IMAGE Consortium	W96939
7268 H	4 021OA1ER	ESTs, Weakly similar to GOLIATH PROTEIN [Drosophila melanogaster] {IMAGE:421432}	421432 IMAGE Consortium	W99150
7269 H	6 021OA1ER	CD24a antigen {IMAGE:421150}	421150 IMAGE Consortium	W98974
7270 H	8 021OA1ER	IMAGE EST {IMAGE:423174}	423174 IMAGE Consortium	W97212
7271 H	10 021OA1ER	ESTs {IMAGE:423393}	423393 IMAGE Consortium	W98894
7272 H	12 021OA1ER	EST {IMAGE:423134}	423134 IMAGE Consortium	W97869
7273 B	2 021VA1ES	Procollagen, type XI, alpha 1 {IMAGE:423028}	423028 IMAGE Consortium	W98013
7274 B	4 021VA1ES	ESTs {IMAGE:421537}	421537 IMAGE Consortium	W97113
7275 B	6 021VA1ES	IMAGE EST {IMAGE:423081}	423081 IMAGE Consortium	W98010
7276 B	8 021VA1ES	ESTs {IMAGE:423276}	423276 IMAGE Consortium	W98720
7277 B	10 021VA1ES	ESTs {IMAGE:421653}	421653 IMAGE Consortium	W98665
7278 B	12 021VA1ES	ESTs, Highly similar to transcription factor E2F-4 [M.musculus] {IMAGE:422188}	422188 IMAGE Consortium	W97445
7279 D	2 021VA1ES	ESTs {IMAGE:421973}	421973 IMAGE Consortium	W91374
7280 D	4 021VA1ES	IMAGE EST {IMAGE:420641}	420641 IMAGE Consortium	W90920
7281 D	6 021VA1ES	IMAGE EST {IMAGE:422060}	422060 IMAGE Consortium	W96845
7282 D	8 021VA1ES	Homeo box B9 {IMAGE:422746}	422746 IMAGE Consortium	W97853
7283 D	10 021VA1ES	ESTs {IMAGE:423125}	423125 IMAGE Consortium	W98042
7284 D	12 021VA1ES	Myelin transcription factor 1-like {IMAGE:423114}	423114 IMAGE Consortium	W97257
7285 F	2 021VA1ES	Eyes absent 2 homolog (Drosophila) {IMAGE:426524}	426524 IMAGE Consortium	AA002292
7286 F	4 021VA1ES	ACETYLCHOLINE RECEPTOR PROTEIN, GAMMA CHAIN PRECURSOR {IMAGE:425717}	425717 IMAGE Consortium	AA000299
7287 F	6 021VA1ES	ESTs {IMAGE:423599}	423599 IMAGE Consortium	W98659
7288 F	8 021VA1ES	ESTs, Weakly similar to HYPOTHETICAL PROTEIN F44G4.1 IN CHROMOSOME II [C.elegans] {IMAGE:424052}	424052 IMAGE Consortium	AA052838
7289 F	10 021VA1ES	IMAGE EST {IMAGE:425849}	425849 IMAGE Consortium	AA000709
7290 F	12 021VA1ES	ESTs {IMAGE:425878}	425878 IMAGE Consortium	AA000714
7291 H	2 021VA1ES	Mast cell protease 5 {IMAGE:425653}	425653 IMAGE Consortium	AA000304
7292 H	4 021VA1ES	ESTs, Moderately similar to NECDIN [Mus musculus] {IMAGE:423743}	423743 IMAGE Consortium	W98359
7293 H	6 021VA1ES	ESTs {IMAGE:424076}	424076 IMAGE Consortium	W91496
7294 H	8 021VA1ES	IMAGE EST {IMAGE:424494}	424494 IMAGE Consortium	W98636
7295 H	10 021VA1ES	ESTs, Weakly similar to LATENT TRANSFORMING GROWTH FACTOR BETA BINDING PROTEIN 1 PRECURSOR [Rattus norvegicus] {IMAGE:424525}	424525 IMAGE Consortium	W98208
7296 H	12 021VA1ES	IMAGE EST {IMAGE:424610}	424610 IMAGE Consortium	AA000495
7297 B	2 021A1ET	Hyaluronan mediated motility receptor (RHAMM) {IMAGE:424197}	424197 IMAGE Consortium	W98230
7298 B	4 021A1ET	Cytochrome P450, 2e1, ethanol inducible {IMAGE:426084}	426084 IMAGE Consortium	AA002850



7299 B	6 0212A1ET	IMAGE EST {IMAGE:424778}	424778 IMAGE Consortium	W98074
7300 B	8 0212A1ET	ESTs {IMAGE:426230}	426230 IMAGE Consortium	AA002751
7301 B	10 0212A1ET	Untitled {IMAGE:425332}	425332 IMAGE Consortium	W99907
7302 B	12 0212A1ET	ESTs, Weakly similar to glucose transport protein, hepatic [M.musculus] {IMAGE:426758}	426758 IMAGE Consortium	AA002666
7303 D	2 0212A1ET	Cyclin D2 {IMAGE:424433}	424433 IMAGE Consortium	W98440
7304 D	4 0212A1ET	ESTs {IMAGE:424851}	424851 IMAGE Consortium	W98123
7305 D	6 0212A1ET	ESTs {IMAGE:424617}	424617 IMAGE Consortium	AA000564
7306 D	8 0212A1ET	IMAGE EST {IMAGE:424839}	424839 IMAGE Consortium	W98190
7307 D	10 0212A1ET	ESTs {IMAGE:425021}	425021 IMAGE Consortium	W99925
7308 D	12 0212A1ET	Mus musculus intermediate conductance potassium channel mlk1 mRNA, complete cds {IMAGE:425427}	425427 IMAGE Consortium	W99968
7309 F	2 0212A1ET	ESTs {IMAGE:424696}	424696 IMAGE Consortium	AA000515
7310 F	4 0212A1ET	ESTs {IMAGE:426529}	426529 IMAGE Consortium	AA002393
7311 F	6 0212A1ET	IMAGE EST {IMAGE:425348}	425348 IMAGE Consortium	W99858
7312 F	8 0212A1ET	IMAGE EST {IMAGE:425523}	425523 IMAGE Consortium	AA000223
7313 F	10 0212A1ET	Mus musculus guanine nucleotide exchange factor delta subunit (JGR1A) mRNA, complete cds {IMAGE:425503}	425503 IMAGE Consortium	AA000219
7314 F	12 0212A1ET	IMAGE EST {IMAGE:425482}	425482 IMAGE Consortium	AA000044
7315 H	2 0212A1ET	ESTs {IMAGE:424702}	424702 IMAGE Consortium	AA000517
7316 H	4 0212A1ET	IMAGE EST {IMAGE:426601}	426601 IMAGE Consortium	AA002567
7317 H	6 0212A1ET	IMAGE EST {IMAGE:426713}	426713 IMAGE Consortium	AA002760
7318 H	8 0212A1ET	ESTs {IMAGE:426551}	426551 IMAGE Consortium	AA002444
7319 H	10 0212A1ET	GLIAL CELL LINE-DERIVED NEUROTROPHIC FACTOR PRECURSOR {IMAGE:425671}	425671 IMAGE Consortium	AA000286
7320 H	12 0212A1ET	IMAGE EST {IMAGE:427005}	427005 IMAGE Consortium	AA003280
7321 B	2 0219A1EU	Mus musculus thrombospondin-4 mRNA, complete cds {IMAGE:437213}	437213 IMAGE Consortium	AA003452
7322 B	4 0219A1EU	IMAGE EST {IMAGE:437035}	437035 IMAGE Consortium	AA003165
7323 B	6 0219A1EU	IMAGE EST {IMAGE:438906}	438906 IMAGE Consortium	AA007974
7324 B	8 0219A1EU	IMAGE EST {IMAGE:437485}	437485 IMAGE Consortium	AA003797
7325 B	10 0219A1EU	Receptor-like tyrosine kinase {IMAGE:439496}	439496 IMAGE Consortium	AA004183
7326 B	12 0219A1EU	IMAGE EST {IMAGE:439213}	439213 IMAGE Consortium	AA003899
7327 D	2 0219A1EU	ESTs, Weakly similar to HYPOTHETICAL 433.2 KD PROTEIN IN HXT5-NRK1 INTERGENIC REGION [S.cerevisiae] {IMAGE:438842}	438842 IMAGE Consortium	AA008375
7328 D	4 0219A1EU	IMAGE EST {IMAGE:436988}	436988 IMAGE Consortium	AA003359
7329 D	6 0219A1EU	Thrombospondin 2 {IMAGE:439102}	439102 IMAGE Consortium	AA003904
7330 D	8 0219A1EU	ESTs {IMAGE:437278}	437278 IMAGE Consortium	AA003672
7331 D	10 0219A1EU	IMAGE EST {IMAGE:437674}	437674 IMAGE Consortium	AA007769
7332 D	12 0219A1EU	IMAGE EST {IMAGE:439555}	439555 IMAGE Consortium	AA008549
7333 F	2 0219A1EU	IMAGE EST {IMAGE:437036}	437036 IMAGE Consortium	AA003367
7334 F	4 0219A1EU	Tumor necrosis factor receptor superfamily, member 1b	437512 IMAGE Consortium	AA003594

		{IMAGE:437512}		
7335 F	6 0219A1EU	ESTs, Highly similar to TFG [H.sapiens] {IMAGE:439282}	439282 IMAGE Consortium	AA003979
7336 F	8 0219A1EU	Mouse testis abundant mRNA sequence {IMAGE:437994}	437994 IMAGE Consortium	AA007791
7337 F	10 0219A1EU	ESTs, Highly similar to D123 [R.norvegicus] {IMAGE:438159}	438159 IMAGE Consortium	AA008062
7338 F	12 0219A1EU	ESTs {IMAGE:438401}	438401 IMAGE Consortium	AA008029
7339 H	2 0219A1EU	IMAGE EST {IMAGE:439364}	439364 IMAGE Consortium	AA004121
7340 H	4 0219A1EU	ESTs {IMAGE:439614}	439614 IMAGE Consortium	AA008684
7341 H	6 0219A1EU	ESTs {IMAGE:439928}	439928 IMAGE Consortium	AA008515
7342 H	8 0219A1EU	ESTs, Highly similar to NADH-UBIQUINONE OXIDOREDUCTASE 15 KD SUBUNIT [Bos taurus] {IMAGE:439668}	439668 IMAGE Consortium	AA014507
7343 H	10 0219A1EU	Cartilage derived retinoic acid sensitive protein {IMAGE:427314}	427314 IMAGE Consortium	AA003262
7344 H	12 0219A1EU	ESTs {IMAGE:436969}	436969 IMAGE Consortium	AA002582
7345 B	2 021GA1EV	Solute carrier family 16 (monocarboxylic acid transporters), member 2 {IMAGE:427197}	427197 IMAGE Consortium	AA003078
7346 B	4 021GA1EV	Cytochrome P450, 2e1, ethanol inducible {IMAGE:424084}	424084 IMAGE Consortium	W91348
7347 B	6 021GA1EV	IMAGE EST {IMAGE:424298}	424298 IMAGE Consortium	W98611
7348 B	8 021GA1EV	IMAGE EST {IMAGE:424463}	424463 IMAGE Consortium	W98571
7349 B	10 021GA1EV	Mus musculus HIC-5 mRNA, complete cds {IMAGE:426146}	426146 IMAGE Consortium	AA002985
7350 B	12 021GA1EV	IMAGE EST {IMAGE:424870}	424870 IMAGE Consortium	AA000598
7351 D	2 021GA1EV	ESTs {IMAGE:426170}	426170 IMAGE Consortium	AA002993
7352 D	4 021GA1EV	7-dehydrocholesterol reductase {IMAGE:426304}	426304 IMAGE Consortium	AA003001
7353 D	6 021GA1EV	IMAGE EST {IMAGE:426166}	426166 IMAGE Consortium	AA003083
7354 D	8 021GA1EV	ESTs {IMAGE:426510}	426510 IMAGE Consortium	AA002379
7355 D	10 021GA1EV	IMAGE EST {IMAGE:426523}	426523 IMAGE Consortium	AA002293
7356 D	12 021GA1EV	ESTs {IMAGE:427248}	427248 IMAGE Consortium	AA002937
7357 F	2 021GA1EV	ESTs {IMAGE:424979}	424979 IMAGE Consortium	W99986
7358 F	4 021GA1EV	ESTs, Highly similar to ER LUMEN PROTEIN RETAINING RECEPTOR 1 [Homo sapiens] {IMAGE:425400}	425400 IMAGE Consortium	AA000093
7359 F	6 021GA1EV	Mouse mRNA for KIFC1, complete cds {IMAGE:426992}	426992 IMAGE Consortium	AA003218
7360 F	8 021GA1EV	EST {IMAGE:427350}	427350 IMAGE Consortium	AA003151
7361 F	10 021GA1EV	RNA polymerase II 1 {IMAGE:425615}	425615 IMAGE Consortium	AA000244
7362 F	12 021GA1EV	IMAGE EST {IMAGE:426076}	426076 IMAGE Consortium	AA000959
7363 H	2 021GA1EV	ESTs {IMAGE:427016}	427016 IMAGE Consortium	AA003064
7364 H	4 021GA1EV	Branched chain aminotransferase 1, cytosolic {IMAGE:426976}	426976 IMAGE Consortium	AA003372
7365 H	6 021GA1EV	ESTs {IMAGE:438415}	438415 IMAGE Consortium	AA008301
7366 H	8 021GA1EV	Seven in absentia 2 {IMAGE:438621}	438621 IMAGE Consortium	AA051014
7367 H	10 021GA1EV	IMAGE EST {IMAGE:437005}	437005 IMAGE Consortium	AA002786
7368 H	12 021GA1EV	ESTs {IMAGE:436895}	436895 IMAGE Consortium	AA002517
7369 B	2 021NA1EW	IMAGE EST {IMAGE:439436}	439436 IMAGE Consortium	AA004142
7370 B	4 021NA1EW	Myosin heavy chain, skeletal muscle, embryonic	439638 IMAGE Consortium	AA008698

		{IMAGE:439638}		
7371 B	6	021NA1EW ESTs {IMAGE:439617}	439617	IMAGE Consortium AA008691
7372 B	8	021NA1EW ESTs, Highly similar to DIHYDROPRYRIDINE-SENSITIVE L-TYPE, CALCIUM CHANNEL BETA-1M SUBUNIT [Homo sapiens] {IMAGE:439812}	439812	IMAGE Consortium AA008590
7373 B	10	021NA1EW IMAGE EST {IMAGE:438665}	438665	IMAGE Consortium AA051089
7374 B	12	021NA1EW IMAGE EST {IMAGE:437194}	437194	IMAGE Consortium AA003532
7375 D	2	021NA1EW IMAGE EST {IMAGE:439920}	439920	IMAGE Consortium AA008730
7376 D	4	021NA1EW Homeo box B6 {IMAGE:439758}	439758	IMAGE Consortium AA008326
7377 D	6	021NA1EW IMAGE EST {IMAGE:438580}	438580	IMAGE Consortium AA050964
7378 D	8	021NA1EW IMAGE EST {IMAGE:438706}	438706	IMAGE Consortium AA051104
7379 D	10	021NA1EW IMAGE EST {IMAGE:436955}	436955	IMAGE Consortium AA002636
7380 D	12	021NA1EW Mus musculus brain fatty acid-binding protein (B-FABP) gene, complete cds {IMAGE:436894}	436894	IMAGE Consortium AA002514
7381 F	2	021NA1EW IMAGE EST {IMAGE:439012}	439012	IMAGE Consortium AA008231
7382 F	4	021NA1EW Proteinase 3 {IMAGE:437587}	437587	IMAGE Consortium AA003779
7383 F	6	021NA1EW IMAGE EST {IMAGE:438001}	438001	IMAGE Consortium AA007970
7384 F	8	021NA1EW IMAGE EST {IMAGE:439311}	439311	IMAGE Consortium AA004002
7385 F	10	021NA1EW IMAGE EST {IMAGE:439706}	439706	IMAGE Consortium AA014288
7386 F	12	021NA1EW ESTs, Highly similar to ZINC FINGER PROTEIN HRX [Homo sapiens] {IMAGE:439854}	439854	IMAGE Consortium AA008671
7387 H	2	021NA1EW IMAGE EST {IMAGE:439351}	439351	IMAGE Consortium AA004037
7388 H	4	021NA1EW IMAGE EST {IMAGE:439383}	439383	IMAGE Consortium AA004056
7389 H	6	021NA1EW IMAGE EST {IMAGE:439778}	439778	IMAGE Consortium AA008332
7390 H	8	021NA1EW ESTs {IMAGE:439902}	439902	IMAGE Consortium AA008727
7391 H	10	021NA1EW IMAGE EST {IMAGE:439594}	439594	IMAGE Consortium AA008663
7392 H	12	021NA1EW ESTs, Weakly similar to MOLYBDENUM COFACTOR BIOSYNTHESIS PROTEIN A [Haemophilus influenzae] {IMAGE:427580}	427580	IMAGE Consortium AA002684
7393 B	2	021UA1EX ESTs {IMAGE:439639}	439639	IMAGE Consortium AA008705
7394 B	4	021UA1EX ESTs, Highly similar to vacuolar proton-ATPase subunit M9.2 [H.sapiens] {IMAGE:439715}	439715	IMAGE Consortium AA014456
7395 B	6	021UA1EX ESTs, Moderately similar to THYMIDINE PHOSPHORYLASE [Homo sapiens] {IMAGE:439622}	439622	IMAGE Consortium AA008687
7396 B	8	021UA1EX ESTs {IMAGE:438516}	438516	IMAGE Consortium AA050783
7397 B	10	021UA1EX Unc-51 like kinase 1 {IMAGE:438917}	438917	IMAGE Consortium AA007768
7398 B	12	021UA1EX IMAGE EST {IMAGE:439395}	439395	IMAGE Consortium AA004065
7399 D	2	021UA1EX ESTs {IMAGE:439914}	439914	IMAGE Consortium AA008563
7400 D	4	021UA1EX ESTs {IMAGE:440194}	440194	IMAGE Consortium AA016584
7401 D	6	021UA1EX ESTs, Moderately similar to CPE-receptor [M.musculus] {IMAGE:441695}	441695	IMAGE Consortium AA013960
7402 D	8	021UA1EX ESTs, Weakly similar to HYPOTHETICAL 47.8 KD PROTEIN B0280.9 IN CHROMOSOME III [C.elegans] {IMAGE:441968}	441968	IMAGE Consortium AA014723

7403 D	10 021UA1EX	ESTs, Weakly similar to ORF YOR173w [S.cerevisiae] {IMAGE:441983}	441983 IMAGE Consortium	AA015421
7404 D	12 021UA1EX	IMAGE EST {IMAGE:439699}	439699 IMAGE Consortium	AA014375
7405 F	2 021UA1EX	IMAGE EST {IMAGE:441864}	441864 IMAGE Consortium	AA013506
7406 F	4 021UA1EX	IMAGE EST {IMAGE:440843}	440843 IMAGE Consortium	AA008751
7407 F	6 021UA1EX	IMAGE EST {IMAGE:441043}	441043 IMAGE Consortium	AA008873
7408 F	8 021UA1EX	IMAGE EST {IMAGE:441264}	441264 IMAGE Consortium	AA011737
7409 F	10 021UA1EX	ESTs {IMAGE:441108}	441108 IMAGE Consortium	AA009039
7410 F	12 021UA1EX	IMAGE EST {IMAGE:439959}	439959 IMAGE Consortium	AA008655
7411 H	2 021UA1EX	IMAGE EST {IMAGE:441235}	441235 IMAGE Consortium	AA009159
7412 H	4 021UA1EX	ESTs {IMAGE:440292}	440292 IMAGE Consortium	AA016694
7413 H	6 021UA1EX	IMAGE EST {IMAGE:441645}	441645 IMAGE Consortium	AA014014
7414 H	8 021UA1EX	IMAGE EST {IMAGE:442080}	442080 IMAGE Consortium	AA013559
7415 H	10 021UA1EX	ESTs {IMAGE:442243}	442243 IMAGE Consortium	AA013567
7416 H	12 021UA1EX	Mus musculus FAN protein mRNA, complete cds {IMAGE:442332}	442332 IMAGE Consortium	AA013700
7417 B	2 0211A1EY	ESTs {IMAGE:442835}	442835 IMAGE Consortium	AA014746
7418 B	4 0211A1EY	ESTs, Moderately similar to motor protein [H.sapiens] {IMAGE:445230}	445230 IMAGE Consortium	AA016847
7419 B	6 0211A1EY	Cholecystokinin {IMAGE:445196}	445196 IMAGE Consortium	AA016824
7420 B	8 0211A1EY	Heat shock protein, 74 kDa {IMAGE:444027}	444027 IMAGE Consortium	AA014915
7421 B	10 0211A1EY	ESTs {IMAGE:443894}	443894 IMAGE Consortium	AA014139
7422 B	12 0211A1EY	ESTs {IMAGE:443852}	443852 IMAGE Consortium	AA014858
7423 D	2 0211A1EY	IMAGE EST {IMAGE:445779}	445779 IMAGE Consortium	AA017911
7424 D	4 0211A1EY	IMAGE EST {IMAGE:445961}	445961 IMAGE Consortium	AA020046
7425 D	6 0211A1EY	IMAGE EST {IMAGE:442692}	442692 IMAGE Consortium	AA015451
7426 D	8 0211A1EY	IMAGE EST {IMAGE:443341}	443341 IMAGE Consortium	AA014964
7427 D	10 0211A1EY	Endothelial-specific receptor tyrosine kinase {IMAGE:445565}	445565 IMAGE Consortium	AA017847
7428 D	12 0211A1EY	IMAGE EST {IMAGE:445435}	445435 IMAGE Consortium	AA017784
7429 F	2 0211A1EY	IMAGE EST {IMAGE:445127}	445127 IMAGE Consortium	AA023461
7430 F	4 0211A1EY	IMAGE EST {IMAGE:443865}	443865 IMAGE Consortium	AA014131
7431 F	6 0211A1EY	IMAGE EST {IMAGE:445900}	445900 IMAGE Consortium	AA020287
7432 F	8 0211A1EY	IMAGE EST {IMAGE:442844}	442844 IMAGE Consortium	AA014195
7433 F	10 0211A1EY	ESTs, Moderately similar to GLIOMA PATHOGENESIS-RELATED PROTEIN [Homo sapiens] {IMAGE:445439}	445439 IMAGE Consortium	AA017785
7434 F	12 0211A1EY	Mus musculus tissue factor pathway inhibitor (TFPI) mRNA, complete cds {IMAGE:445829}	445829 IMAGE Consortium	AA020100
7435 H	2 0211A1EY	ESTs {IMAGE:455497}	455497 IMAGE Consortium	AA020515
7436 H	4 0211A1EY	IMAGE EST {IMAGE:457569}	457569 IMAGE Consortium	AA021708
7437 H	6 0211A1EY	IMAGE EST {IMAGE:455955}	455955 IMAGE Consortium	AA023085
7438 H	8 0211A1EY	M.musculus mRNA for 11beta-hydroxysteroid dehydrogenase/carbonyl reductase {IMAGE:455945}	455945 IMAGE Consortium	AA023077
7439 H	10 0211A1EY	ESTs, Weakly similar to similar to alpha-actinin [C.elegans]	457000 IMAGE Consortium	AA023523

7440 H	12 0211A1EY	{IMAGE:457000} ESTs, Highly similar to sorting nexin 1 [H.sapiens] {IMAGE:456862}	456862 IMAGE Consortium	AA023638
7441 B	2 0218A1EZ	IMAGE EST {IMAGE:440985}	440985 IMAGE Consortium	AA008960
7442 B	4 0218A1EZ	Mus musculus junctional adhesion molecule (Jam) mRNA, complete cds {IMAGE:440944}	440944 IMAGE Consortium	AA009015
7443 B	6 0218A1EZ	ESTs, Weakly similar to transcription factor C1 [M.musculus] {IMAGE:441526}	441526 IMAGE Consortium	AA013895
7444 B	8 0218A1EZ	ESTs, Moderately similar to ANTIGEN WC1.1 [Bos taurus] {IMAGE:441064}	441064 IMAGE Consortium	AA008876
7445 B	10 0218A1EZ	Myelocytomatosis oncogene {IMAGE:441346}	441346 IMAGE Consortium	AA009268
7446 B	12 0218A1EZ	IMAGE EST {IMAGE:440264}	440264 IMAGE Consortium	AA016661
7447 D	2 0218A1EZ	ESTs {IMAGE:440490}	440490 IMAGE Consortium	AA009106
7448 D	4 0218A1EZ	IMAGE EST {IMAGE:442135}	442135 IMAGE Consortium	AA013528
7449 D	6 0218A1EZ	IMAGE EST {IMAGE:442396}	442396 IMAGE Consortium	AA013792
7450 D	8 0218A1EZ	IMAGE EST {IMAGE:441327}	441327 IMAGE Consortium	AA009266
7451 D	10 0218A1EZ	IMAGE EST {IMAGE:441519}	441519 IMAGE Consortium	AA013802
7452 D	12 0218A1EZ	IMAGE EST {IMAGE:440911}	440911 IMAGE Consortium	AA008899
7453 F	2 0218A1EZ	Glucosaminyl (N-acetyl) transferase 1, core 2 {IMAGE:444411}	444411 IMAGE Consortium	AA016940
7454 F	4 0218A1EZ	IMAGE EST {IMAGE:442966}	442966 IMAGE Consortium	AA014142
7455 F	6 0218A1EZ	ESTs, Weakly similar to CYTOSKELETAL P17 PROTEIN [Dictyostelium discoideum] {IMAGE:445352}	445352 IMAGE Consortium	AA017875
7456 F	8 0218A1EZ	IMAGE EST {IMAGE:444106}	444106 IMAGE Consortium	AA014942
7457 F	10 0218A1EZ	IMAGE EST {IMAGE:443965}	443965 IMAGE Consortium	AA014971
7458 F	12 0218A1EZ	ESTs {IMAGE:444162}	444162 IMAGE Consortium	AA015176
7459 H	2 0218A1EZ	IMAGE EST {IMAGE:444412}	444412 IMAGE Consortium	AA016920
7460 H	4 0218A1EZ	IMAGE EST {IMAGE:444817}	444817 IMAGE Consortium	AA016430
7461 H	6 0218A1EZ	ESTs, Weakly similar to K04G7.11 [C.elegans] {IMAGE:445520}	445520 IMAGE Consortium	AA017918
7462 H	8 0218A1EZ	EST {IMAGE:443843}	443843 IMAGE Consortium	AA014121
7463 H	10 0218A1EZ	SECRETORY GRANULE PROTEOGLYCAN CORE PROTEIN PRECURSOR {IMAGE:445929}	445929 IMAGE Consortium	AA020057
7464 H	12 0218A1EZ	ESTs {IMAGE:444436}	444436 IMAGE Consortium	AA016942
7465 B	2 0218A1F0	Mus musculus mRNA for glycoprotein-associated amino acid transporter y+LAT1b {IMAGE:457955}	457955 IMAGE Consortium	AA030193
7466 B	4 0218A1F0	EST {IMAGE:458068}	458068 IMAGE Consortium	AA030667
7467 B	6 0218A1F0	Cut (Drosophila)-like 1 {IMAGE:458712}	458712 IMAGE Consortium	AA023906
7468 B	8 0218A1F0	IMAGE EST {IMAGE:458911}	458911 IMAGE Consortium	AA024141
7469 B	10 0218A1F0	IMAGE EST {IMAGE:457289}	457289 IMAGE Consortium	AA023664
7470 B	12 0218A1F0	Granzyme F {IMAGE:456459}	456459 IMAGE Consortium	AA023418
7471 D	2 0218A1F0	IMAGE EST {IMAGE:457832}	457832 IMAGE Consortium	AA021966
7472 D	4 0218A1F0	Mus musculus semaF cytoplasmic domain associated protein	457038 IMAGE Consortium	AA023463

		2 (Semcap2) mRNA, complete cds {IMAGE:457038}		
7473 D	6 0218A1F0	Dynamin 2 {IMAGE:457445}	457445 IMAGE Consortium	AA021681
7474 D	8 0218A1F0	IMAGE EST {IMAGE:457704}	457704 IMAGE Consortium	AA021760
7475 D	10 0218A1F0	ESTs, Weakly similar to ACYL CARRIER PROTEIN, MITOCHONDRIAL PRECURSOR [Neurospora crassa] {IMAGE:456543}	456543 IMAGE Consortium	AA023445
7476 D	12 0218A1F0	ESTs {IMAGE:446015}	446015 IMAGE Consortium	AA020272
7477 F	2 0218A1F0	ESTs {IMAGE:458684}	458684 IMAGE Consortium	AA023914
7478 F	4 0218A1F0	IMAGE EST {IMAGE:455438}	455438 IMAGE Consortium	AA020480
7479 F	6 0218A1F0	Signal transducer and activator of transcription 1 {IMAGE:455610}	455610 IMAGE Consortium	AA023159
7480 F	8 0218A1F0	IMAGE EST {IMAGE:457827}	457827 IMAGE Consortium	AA021858
7481 F	10 0218A1F0	IMAGE EST {IMAGE:457769}	457769 IMAGE Consortium	AA021871
7482 F	12 0218A1F0	Mus musculus Xin mRNA, complete cds {IMAGE:456666}	456666 IMAGE Consortium	AA023480
7483 H	2 0218A1F0	IMAGE EST {IMAGE:458349}	458349 IMAGE Consortium	AA024360
7484 H	4 0218A1F0	Mus musculus mRNA for transcription factor TFIIH, 62 kD subunit {IMAGE:458432}	458432 IMAGE Consortium	AA024318
7485 H	6 0218A1F0	Transcription factor AP-2, alpha {IMAGE:458629}	458629 IMAGE Consortium	AA023994
7486 H	8 0218A1F0	IMAGE EST {IMAGE:457442}	457442 IMAGE Consortium	AA021678
7487 H	10 0218A1F0	Mus musculus LIM protein 3 (mSLIM3) mRNA, complete cds {IMAGE:457264}	457264 IMAGE Consortium	AA023645
7488 H	12 0218A1F0	IMAGE EST {IMAGE:455825}	455825 IMAGE Consortium	AA023805
7489 B	2 021FA1F1	Mus musculus mRNA for 3'UTR of Clc1 gene {IMAGE:464457}	464457 IMAGE Consortium	AA033085
7490 B	4 021FA1F1	IMAGE EST {IMAGE:463064}	463064 IMAGE Consortium	AA027673
7491 B	6 021FA1F1	Defender against cell death 1 {IMAGE:464622}	464622 IMAGE Consortium	AA033006
7492 B	8 021FA1F1	ESTs, Moderately similar to 60S RIBOSOMAL PROTEIN L7 [Mus musculus] {IMAGE:464556}	464556 IMAGE Consortium	AA030122
7493 B	10 021FA1F1	Hydroxysteroid dehydrogenase-1, delta<5>-3-beta {IMAGE:463566}	463566 IMAGE Consortium	AA028760
7494 B	12 021FA1F1	ESTs, Weakly similar to HYPOTHETICAL 54.9 KD PROTEIN C02F5.7 IN CHROMOSOME III [Caenorhabditis elegans] {IMAGE:464118}	464118 IMAGE Consortium	AA028564
7495 D	2 021FA1F1	Mus musculus unknown protein mRNA, complete cds {IMAGE:463590}	463590 IMAGE Consortium	AA028678
7496 D	4 021FA1F1	ESTs, Moderately similar to ACYL-COA-BINDING PROTEIN [Bos taurus] {IMAGE:463575}	463575 IMAGE Consortium	AA030780
7497 D	6 021FA1F1	Mus musculus RaiBP1-associated EH domain protein Repl1 (reps1) mRNA, complete cds {IMAGE:458966}	458966 IMAGE Consortium	AA024100
7498 D	8 021FA1F1	ESTs {IMAGE:463927}	463927 IMAGE Consortium	AA028495
7499 D	10 021FA1F1	ESTs {IMAGE:459390}	459390 IMAGE Consortium	AA027412
7500 D	12 021FA1F1	Mus musculus sex-determination protein homolog Fem1b mRNA, complete cds {IMAGE:459309}	459309 IMAGE Consortium	AA030303
7501 F	2 021FA1F1	IMAGE EST {IMAGE:463599}	463599 IMAGE Consortium	AA030786

7502 F	4 021FA1F1	IMAGE EST {IMAGE:459090}	459090 IMAGE Consortium	AA024025
7503 F	6 021FA1F1	Enhancer trap locus 1 {IMAGE:459319}	459319 IMAGE Consortium	AA024295
7504 F	8 021FA1F1	IMAGE EST {IMAGE:464134}	464134 IMAGE Consortium	AA028359
7505 F	10 021FA1F1	IMAGE EST {IMAGE:464434}	464434 IMAGE Consortium	AA033058
7506 F	12 021FA1F1	ESTs {IMAGE:464744}	464744 IMAGE Consortium	AA030822
7507 H	2 021FA1F1	IMAGE EST {IMAGE:463038}	463038 IMAGE Consortium	AA027649
7508 H	4 021FA1F1	Natriuretic peptide precursor type B {IMAGE:463618}	463618 IMAGE Consortium	AA030805
7509 H	6 021FA1F1	IMAGE EST {IMAGE:458871}	458871 IMAGE Consortium	AA024191
7510 H	8 021FA1F1	ESTs {IMAGE:459488}	459488 IMAGE Consortium	AA027371
7511 H	10 021FA1F1	IMAGE EST {IMAGE:464263}	464263 IMAGE Consortium	AA030682
7512 H	12 021FA1F1	IMAGE EST {IMAGE:463273}	463273 IMAGE Consortium	AA027683
7513 B	2 021MA1F2	ESTs {IMAGE:463342}	463342 IMAGE Consortium	AA028696
7514 B	4 021MA1F2	IMAGE EST {IMAGE:464828}	464828 IMAGE Consortium	AA032419
7515 B	6 021MA1F2	ESTs, Highly similar to ZINC FINGER PROTEIN 7 [Homo sapiens] {IMAGE:466781}	466781 IMAGE Consortium	AA031146
7516 B	8 021MA1F2	Seizure related gene 6 {IMAGE:465285}	465285 IMAGE Consortium	AA030773
7517 B	10 021MA1F2	IMAGE EST {IMAGE:465403}	465403 IMAGE Consortium	AA032691
7518 B	12 021MA1F2	ESTs {IMAGE:465804}	465804 IMAGE Consortium	AA032828
7519 D	2 021MA1F2	IMAGE EST {IMAGE:465475}	465475 IMAGE Consortium	AA032702
7520 D	4 021MA1F2	IMAGE EST {IMAGE:467182}	467182 IMAGE Consortium	AA034840
7521 D	6 021MA1F2	IMAGE EST {IMAGE:467511}	467511 IMAGE Consortium	AA036444
7522 D	8 021MA1F2	IMAGE EST {IMAGE:466057}	466057 IMAGE Consortium	AA034677
7523 D	10 021MA1F2	IMAGE EST {IMAGE:468218}	468218 IMAGE Consortium	AA015094
7524 D	12 021MA1F2	Mus musculus myosin light chain 2 mRNA, complete cds {IMAGE:466382}	466382 IMAGE Consortium	AA032362
7525 F	2 021MA1F2	IMAGE EST {IMAGE:466252}	466252 IMAGE Consortium	AA033305
7526 F	4 021MA1F2	IMAGE EST {IMAGE:468175}	468175 IMAGE Consortium	AA015088
7527 F	6 021MA1F2	Mus musculus shd mRNA, complete cds {IMAGE:466339}	466339 IMAGE Consortium	AA033417
7528 F	8 021MA1F2	ESTs {IMAGE:465060}	465060 IMAGE Consortium	AA032514
7529 F	10 021MA1F2	IMAGE EST {IMAGE:465043}	465043 IMAGE Consortium	AA032621
7530 F	12 021MA1F2	ESTs {IMAGE:466822}	466822 IMAGE Consortium	AA031177
7531 H	2 021MA1F2	ESTs, Moderately similar to APC-binding protein EB1 homolog [M.musculus] {IMAGE:466759}	466759 IMAGE Consortium	AA031126
7532 H	4 021MA1F2	IMAGE EST {IMAGE:466715}	466715 IMAGE Consortium	AA031035
7533 H	6 021MA1F2	IMAGE EST {IMAGE:465596}	465596 IMAGE Consortium	AA032786
7534 H	8 021MA1F2	IMAGE EST {IMAGE:467472}	467472 IMAGE Consortium	AA034711
7535 H	10 021MA1F2	IMAGE EST {IMAGE:467560}	467560 IMAGE Consortium	AA034730
7536 H	12 021MA1F2	ESTs {IMAGE:467979}	467979 IMAGE Consortium	AA015192
7537 B	2 021TA1F3	CAMP responsive element modulator {IMAGE:459249}	459249 IMAGE Consortium	AA030254
7538 B	4 021TA1F3	ESTs, Highly similar to L-SERINE DEHYDRATASE [Rattus norvegicus] {IMAGE:463345}	463345 IMAGE Consortium	AA027702
7539 B	6 021TA1F3	ESTs {IMAGE:464396}	464396 IMAGE Consortium	AA030659
7540 B	8 021TA1F3	IMAGE EST {IMAGE:464578}	464578 IMAGE Consortium	AA030144

7541 B	10 021TA1F3	ESTs, Highly similar to 80 KD NUCLEAR CAP BINDING PROTEIN [H.sapiens] {IMAGE:463602}	463602 IMAGE Consortium	AA027752
7542 B	12 021TA1F3	IMAGE EST {IMAGE:458776}	458776 IMAGE Consortium	AA024086
7543 D	2 021TA1F3	ESTs {IMAGE:458996}	458996 IMAGE Consortium	AA024160
7544 D	4 021TA1F3	IMAGE EST {IMAGE:459491}	459491 IMAGE Consortium	AA027443
7545 D	6 021TA1F3	ESTs {IMAGE:464280}	464280 IMAGE Consortium	AA030683
7546 D	8 021TA1F3	IMAGE EST {IMAGE:459571}	459571 IMAGE Consortium	AA027397
7547 D	10 021TA1F3	3-hydroxy-3-methylglutaryl coenzyme A synthase 2 {IMAGE:464392}	464392 IMAGE Consortium	AA030116
7548 D	12 021TA1F3	ESTs, Weakly similar to gene mCBP protein [M.musculus] {IMAGE:464397}	464397 IMAGE Consortium	AA030663
7549 F	2 021TA1F3	ESTs, Highly similar to DOWN REGULATORY PROTEIN OF INTERLEUKIN 2 RECEPTOR [Mus musculus] {IMAGE:463322}	463322 IMAGE Consortium	AA027694
7550 F	4 021TA1F3	IMAGE EST {IMAGE:463207}	463207 IMAGE Consortium	AA030366
7551 F	6 021TA1F3	ESTs {IMAGE:463044}	463044 IMAGE Consortium	AA027652
7552 F	8 021TA1F3	IMAGE EST {IMAGE:464866}	464866 IMAGE Consortium	AA030521
7553 F	10 021TA1F3	ESTs {IMAGE:464535}	464535 IMAGE Consortium	AA030119
7554 F	12 021TA1F3	ESTs, Highly similar to PROCHYMO SIN PRECURSOR, B-FORM [Ovis aries] {IMAGE:463502}	463502 IMAGE Consortium	AA028632
7555 H	2 021TA1F3	IMAGE EST {IMAGE:463092}	463092 IMAGE Consortium	AA030074
7556 H	4 021TA1F3	Adenine phosphoribosyl transferase {IMAGE:463241}	463241 IMAGE Consortium	AA030540
7557 H	6 021TA1F3	ESTs {IMAGE:463270}	463270 IMAGE Consortium	AA030634
7558 H	8 021TA1F3	Mus domesticus nuclear binding factor NF2d9 mRNA, complete cds {IMAGE:463518}	463518 IMAGE Consortium	AA027734
7559 H	10 021TA1F3	IMAGE EST {IMAGE:463646}	463646 IMAGE Consortium	AA030814
7560 H	12 021TA1F3	IMAGE EST {IMAGE:464915}	464915 IMAGE Consortium	AA031238
7561 B	2 0210A1F4	Homeo box A5 {IMAGE:465937}	465937 IMAGE Consortium	AA033195
7562 B	4 0210A1F4	Stromal interaction molecule 1 {IMAGE:465824}	465824 IMAGE Consortium	AA032834
7563 B	6 0210A1F4	IMAGE EST {IMAGE:465758}	465758 IMAGE Consortium	AA033125
7564 B	8 0210A1F4	Mus musculus mRNA for actin-related protein 1 alpha-isoform, complete cds {IMAGE:466134}	466134 IMAGE Consortium	AA034697
7565 B	10 0210A1F4	IMAGE EST {IMAGE:468272}	468272 IMAGE Consortium	AA014385
7566 B	12 0210A1F4	ESTs {IMAGE:608410}	608410 IMAGE Consortium	AA168066
7567 D	2 0210A1F4	Yamaguchi sarcoma viral (v-yes-1) oncogene homolog {IMAGE:465297}	465297 IMAGE Consortium	AA032394
7568 D	4 0210A1F4	ESTs, Highly similar to PROTEASOME COMPONENT C2 [Homo sapiens] {IMAGE:466041}	466041 IMAGE Consortium	AA031120
7569 D	6 0210A1F4	IMAGE EST {IMAGE:467394}	467394 IMAGE Consortium	AA036434
7570 D	8 0210A1F4	IMAGE EST {IMAGE:466178}	466178 IMAGE Consortium	AA033355
7571 D	10 0210A1F4	IMAGE EST {IMAGE:466428}	466428 IMAGE Consortium	AA032386
7572 D	12 0210A1F4	IMAGE EST {IMAGE:466150}	466150 IMAGE Consortium	AA033320
7573 F	2 0210A1F4	IMAGE EST {IMAGE:467830}	467830 IMAGE Consortium	AA036537



7574 F	4 0210A1F4	ESTs, Highly similar to ZINC FINGER PROTEIN ZFP-37 [Mus musculus] {IMAGE:466313}	466313 IMAGE Consortium	AA033410
7575 F	6 0210A1F4	IMAGE EST {IMAGE:468536}	468536 IMAGE Consortium	AA014324
7576 F	8 0210A1F4	IMAGE EST {IMAGE:466645}	466645 IMAGE Consortium	AA031004
7577 F	10 0210A1F4	ESTs {IMAGE:466626}	466626 IMAGE Consortium	AA030988
7578 F	12 0210A1F4	IMAGE EST {IMAGE:465724}	465724 IMAGE Consortium	AA032893
7579 H	2 0210A1F4	ESTs {IMAGE:465292}	465292 IMAGE Consortium	AA032709
7580 H	4 0210A1F4	Adenine nucleotide translocator 2, fibroblast {IMAGE:465520}	465520 IMAGE Consortium	AA033138
7581 H	6 0210A1F4	ESTs {IMAGE:465940}	465940 IMAGE Consortium	AA033193
7582 H	8 0210A1F4	IMAGE EST {IMAGE:466043}	466043 IMAGE Consortium	AA031055
7583 H	10 0210A1F4	ESTs, Highly similar to NADH-UBIQUINONE OXIDOREDUCTASE SUBUNIT B14.5A [Bos taurus] {IMAGE:465903}	465903 IMAGE Consortium	AA033262
7584 H	12 0210A1F4	IMAGE EST {IMAGE:466423}	466423 IMAGE Consortium	AA032330
7585 B	2 0217A1F5	ESTs, Weakly similar to contains similarity to S. cerevisiae mitochondrial outer membrane protein 22kDa protein [C.elegans] {IMAGE:467790}	467790 IMAGE Consortium	AA036494
7586 B	4 0217A1F5	Retinoid X receptor gamma {IMAGE:466208}	466208 IMAGE Consortium	AA033489
7587 B	6 0217A1F5	FMS-like tyrosine kinase 1 {IMAGE:468210}	468210 IMAGE Consortium	AA014250
7588 B	8 0217A1F5	Dlk1-like homolog (Drosophila) {IMAGE:466247}	466247 IMAGE Consortium	AA032368
7589 B	10 0217A1F5	Cytochrome P450, 2b19 {IMAGE:473877}	473877 IMAGE Consortium	AA038397
7590 B	12 0217A1F5	IMAGE EST {IMAGE:468966}	468966 IMAGE Consortium	AA036282
7591 D	2 0217A1F5	IMAGE EST {IMAGE:468809}	468809 IMAGE Consortium	AA035924
7592 D	4 0217A1F5	ESTs {IMAGE:472376}	472376 IMAGE Consortium	AA036357
7593 D	6 0217A1F5	ESTs, Weakly similar to C05D11.3 [C.elegans] {IMAGE:472623}	472623 IMAGE Consortium	AA037970
7594 D	8 0217A1F5	IMAGE EST {IMAGE:475345}	475345 IMAGE Consortium	AA041924
7595 D	10 0217A1F5	IMAGE EST {IMAGE:473759}	473759 IMAGE Consortium	AA038253
7596 D	12 0217A1F5	IMAGE EST {IMAGE:468857}	468857 IMAGE Consortium	AA036169
7597 F	2 0217A1F5	IMAGE EST {IMAGE:469077}	469077 IMAGE Consortium	AA036270
7598 F	4 0217A1F5	DiGeorge syndrome gene c {IMAGE:472948}	472948 IMAGE Consortium	AA038087
7599 F	6 0217A1F5	IMAGE EST {IMAGE:475562}	475562 IMAGE Consortium	AA050185
7600 F	8 0217A1F5	ESTs, Weakly similar to Ydr472wp [S.cerevisiae] {IMAGE:474336}	474336 IMAGE Consortium	AA038812
7601 F	10 0217A1F5	IMAGE EST {IMAGE:469063}	469063 IMAGE Consortium	AA036379
7602 F	12 0217A1F5	IMAGE EST {IMAGE:469125}	469125 IMAGE Consortium	AA036322
7603 H	2 0217A1F5	IMAGE EST {IMAGE:474089}	474089 IMAGE Consortium	AA038718
7604 H	4 0217A1F5	IMAGE EST {IMAGE:474184}	474184 IMAGE Consortium	AA038737
7605 H	6 0217A1F5	IMAGE EST {IMAGE:472450}	472450 IMAGE Consortium	AA036411
7606 H	8 0217A1F5	ESTs, Moderately similar to COATOMER DELTA SUBUNIT [Homo sapiens] {IMAGE:472626}	472626 IMAGE Consortium	AA038054
7607 H	10 0217A1F5	Pyruvate decarboxylase {IMAGE:473778}	473778 IMAGE Consortium	AA038254
7608 H	12 0217A1F5	Cell division cycle control protein 2a {IMAGE:468792}	468792 IMAGE Consortium	AA035888

7609 B	2 021EA1F6	Catenin alpha 2 {IMAGE:475651}	475651 IMAGE Consortium	AA049813
7610 B	4 021EA1F6	ESTs {IMAGE:475824}	475824 IMAGE Consortium	AA050420
7611 B	6 021EA1F6	Mus musculus secretory carrier membrane protein (SCAMP3) mRNA, complete cds {IMAGE:477267}	477267 IMAGE Consortium	AA048238
7612 B	8 021EA1F6	IMAGE EST {IMAGE:475797}	475797 IMAGE Consortium	AA050139
7613 B	10 021EA1F6	IMAGE EST {IMAGE:476055}	476055 IMAGE Consortium	AA050178
7614 B	12 021EA1F6	ESTs, Highly similar to pescadillo [H.sapiens] {IMAGE:477900}	477900 IMAGE Consortium	AA048619
7615 D	2 021EA1F6	ESTs {IMAGE:476082}	476082 IMAGE Consortium	AA049842
7616 D	4 021EA1F6	IMAGE EST {IMAGE:477882}	477882 IMAGE Consortium	AA049406
7617 D	6 021EA1F6	Mus musculus ADP-ribosylation-like factor homolog ARL6 (Arl6) mRNA, complete cds {IMAGE:476610}	476610 IMAGE Consortium	AA050789
7618 D	8 021EA1F6	Mus musculus Ena-VASP like protein (Evl) mRNA, complete cds {IMAGE:477177}	477177 IMAGE Consortium	AA048116
7619 D	10 021EA1F6	Mus musculus mRNA for p67phox, complete cds {IMAGE:475840}	475840 IMAGE Consortium	AA050149
7620 D	12 021EA1F6	IMAGE EST {IMAGE:476099}	476099 IMAGE Consortium	AA050371
7621 F	2 021EA1F6	ESTs, Highly similar to NECDIN [Mus musculus] {IMAGE:476509}	476509 IMAGE Consortium	AA050644
7622 F	4 021EA1F6	IMAGE EST {IMAGE:478172}	478172 IMAGE Consortium	AA049701
7623 F	6 021EA1F6	Mus musculus ubiquitin conjugating enzyme UBC9 mRNA, complete cds {IMAGE:475503}	475503 IMAGE Consortium	AA050086
7624 F	8 021EA1F6	ESTs {IMAGE:475681}	475681 IMAGE Consortium	AA050058
7625 F	10 021EA1F6	ESTs {IMAGE:477484}	477484 IMAGE Consortium	AA048400
7626 F	12 021EA1F6	ESTs {IMAGE:477559}	477559 IMAGE Consortium	AA048262
7627 H	2 021EA1F6	Transcription factor 1 {IMAGE:475809}	475809 IMAGE Consortium	AA049898
7628 H	4 021EA1F6	IMAGE EST {IMAGE:476268}	476268 IMAGE Consortium	AA050767
7629 H	6 021EA1F6	ESTs {IMAGE:476275}	476275 IMAGE Consortium	AA050575
7630 H	8 021EA1F6	Mus musculus Su(var)3-9 homolog (Suv39h) mRNA, complete cds {IMAGE:476760}	476760 IMAGE Consortium	AA050907
7631 H	10 021EA1F6	ESTs {IMAGE:476653}	476653 IMAGE Consortium	AA050547
7632 H	12 021EA1F6	IMAGE EST {IMAGE:478244}	478244 IMAGE Consortium	AA049736
7633 B	2 021LA1F7	IMAGE EST {IMAGE:472374}	472374 IMAGE Consortium	AA036255
7634 B	4 021LA1F7	ESTs {IMAGE:475467}	475467 IMAGE Consortium	AA042006
7635 B	6 021LA1F7	Keratin complex 2, basic, gene 4 {IMAGE:474387}	474387 IMAGE Consortium	AA038826
7636 B	8 021LA1F7	ESTs {IMAGE:469076}	469076 IMAGE Consortium	AA036267
7637 B	10 021LA1F7	ESTs {IMAGE:474091}	474091 IMAGE Consortium	AA038719
7638 B	12 021LA1F7	Ring finger protein 3 {IMAGE:474292}	474292 IMAGE Consortium	AA038785
7639 D	2 021LA1F7	ESTs {IMAGE:474467}	474467 IMAGE Consortium	AA038926
7640 D	4 021LA1F7	IMAGE EST {IMAGE:472423}	472423 IMAGE Consortium	AA036150
7641 D	6 021LA1F7	ESTs {IMAGE:472770}	472770 IMAGE Consortium	AA038182
7642 D	8 021LA1F7	ESTs {IMAGE:474104}	474104 IMAGE Consortium	AA038634
7643 D	10 021LA1F7	RSP-1 PROTEIN {IMAGE:473805}	473805 IMAGE Consortium	AA038395

7644 D	12 021LA1F7	Phospholipase C, beta 3 {IMAGE:475397}	475397 IMAGE Consortium	AA041709
7645 F	2 021LA1F7	IMAGE EST {IMAGE:476198}	476198 IMAGE Consortium	AA050200
7646 F	4 021LA1F7	ESTs {IMAGE:477665}	477665 IMAGE Consortium	AA048581
7647 F	6 021LA1F7	HISTONE H1.1 {IMAGE:477962}	477962 IMAGE Consortium	AA049416
7648 F	8 021LA1F7	ESTs {IMAGE:476776}	476776 IMAGE Consortium	AA050447
7649 F	10 021LA1F7	EST {IMAGE:476567}	476567 IMAGE Consortium	AA050713
7650 F	12 021LA1F7	IMAGE EST {IMAGE:475285}	475285 IMAGE Consortium	AA042043
7651 H	2 021LA1F7	EST {IMAGE:476615}	476615 IMAGE Consortium	AA050810
7652 H	4 021LA1F7	Casein kinase II, alpha 2, polypeptide {IMAGE:475351}	475351 IMAGE Consortium	AA041970
7653 H	6 021LA1F7	IMAGE EST {IMAGE:475333}	475333 IMAGE Consortium	AA042056
7654 H	8 021LA1F7	ESTs {IMAGE:476966}	476966 IMAGE Consortium	AA048028
7655 H	10 021LA1F7	IMAGE EST {IMAGE:477195}	477195 IMAGE Consortium	AA048217
7656 H	12 021LA1F7	ESTs {IMAGE:476185}	476185 IMAGE Consortium	AA050592
7657 B	2 021SA1F8	IMAGE EST {IMAGE:476694}	476694 IMAGE Consortium	AA050504
7658 B	4 021SA1F8	ESTs, Weakly similar to cyclin G2 [M.musculus] {IMAGE:476702}	476702 IMAGE Consortium	AA050559
7659 B	6 021SA1F8	IMAGE EST {IMAGE:475475}	475475 IMAGE Consortium	AA050078
7660 B	8 021SA1F8	Mus musculus nucleolar protein (MSP58) mRNA, complete cds {IMAGE:477036}	477036 IMAGE Consortium	AA048134
7661 B	10 021SA1F8	Keratin complex 2, basic, gene 1 {IMAGE:476999}	476999 IMAGE Consortium	AA047991
7662 B	12 021SA1F8	ESTs, Highly similar to SERINE/THREONINE PROTEIN PHOSPHATASE PP-X [Homo sapiens] {IMAGE:476407}	476407 IMAGE Consortium	AA050169
7663 D	2 021SA1F8	IMAGE EST {IMAGE:477264}	477264 IMAGE Consortium	AA048234
7664 D	4 021SA1F8	IMAGE EST {IMAGE:477509}	477509 IMAGE Consortium	AA048429
7665 D	6 021SA1F8	ESTs, Weakly similar to LIGHT-MEDIATED DEVELOPMENT PROTEIN DET1 [Arabidopsis thaliana] {IMAGE:476096}	476096 IMAGE Consortium	AA050184
7666 D	8 021SA1F8	Mus musculus RAB/Rip protein mRNA, partial cds {IMAGE:476081}	476081 IMAGE Consortium	AA049862
7667 D	10 021SA1F8	IMAGE EST {IMAGE:476767}	476767 IMAGE Consortium	AA050914
7668 D	12 021SA1F8	EST {IMAGE:476703}	476703 IMAGE Consortium	AA050562
7669 F	2 021SA1F8	IMAGE EST {IMAGE:480205}	480205 IMAGE Consortium	AA058194
7670 F	4 021SA1F8	Mus musculus high mobility group protein homolog HMG4 (Hmg4) mRNA, complete cds {IMAGE:478865}	478865 IMAGE Consortium	AA048831
7671 F	6 021SA1F8	ESTs {IMAGE:479233}	479233 IMAGE Consortium	AA048723
7672 F	8 021SA1F8	EST {IMAGE:480583}	480583 IMAGE Consortium	AA064015
7673 F	10 021SA1F8	ESTs {IMAGE:480419}	480419 IMAGE Consortium	AA058171
7674 F	12 021SA1F8	ESTs {IMAGE:478125}	478125 IMAGE Consortium	AA049592
7675 H	2 021SA1F8	ESTs, Weakly similar to R05G6.4 gene product [C.elegans] {IMAGE:480164}	480164 IMAGE Consortium	AA058020
7676 H	4 021SA1F8	IMAGE EST {IMAGE:480434}	480434 IMAGE Consortium	AA058225
7677 H	6 021SA1F8	ESTs, Highly similar to THREONYL-TRNA SYNTHETASE, CYTOPLASMIC [Homo sapiens] {IMAGE:480488}	480488 IMAGE Consortium	AA064056
7678 H	8 021SA1F8	ESTs, Moderately similar to CYTOCHROME B5, OUTER	479126 IMAGE Consortium	AA049211

## MITOCHONDRIAL MEMBRANE [Rattus norvegicus]

		{IMAGE:479126}		
7679 H	10 021SA1F8	ESTs {IMAGE:478006}	478006 IMAGE Consortium	AA049551
7680 H	12 021SA1F8	IMAGE EST {IMAGE:478426}	478426 IMAGE Consortium	AA049818
7681 B	2 021ZA1F9	ESTs {IMAGE:479811}	479811 IMAGE Consortium	AA051493
7682 B	4 021ZA1F9	IMAGE EST {IMAGE:479726}	479726 IMAGE Consortium	AA051613
7683 B	6 021ZA1F9	Mast cell protease 6 {IMAGE:479561}	479561 IMAGE Consortium	AA049080
7684 B	8 021ZA1F9	IMAGE EST {IMAGE:480193}	480193 IMAGE Consortium	AA058057
7685 B	10 021ZA1F9	IMAGE EST {IMAGE:479962}	479962 IMAGE Consortium	AA051699
7686 B	12 021ZA1F9	Platelet derived growth factor receptor, alpha polypeptide {IMAGE:479895}	479895 IMAGE Consortium	AA051556
7687 D	2 021ZA1F9	ESTs, Weakly similar to cDNA EST EMBL:T00822 comes from this gene [C.elegans] {IMAGE:479978}	479978 IMAGE Consortium	AA051433
7688 D	4 021ZA1F9	ESTs {IMAGE:478904}	478904 IMAGE Consortium	AA051271
7689 D	6 021ZA1F9	ESTs {IMAGE:478963}	478963 IMAGE Consortium	AA049152
7690 D	8 021ZA1F9	IMAGE EST {IMAGE:480612}	480612 IMAGE Consortium	AA060687
7691 D	10 021ZA1F9	ESTs {IMAGE:480430}	480430 IMAGE Consortium	AA058134
7692 D	12 021ZA1F9	IMAGE EST {IMAGE:478686}	478686 IMAGE Consortium	AA051374
7693 F	2 021ZA1F9	IMAGE EST {IMAGE:478512}	478512 IMAGE Consortium	AA049927
7694 F	4 021ZA1F9	IMAGE EST {IMAGE:478710}	478710 IMAGE Consortium	AA051380
7695 F	6 021ZA1F9	ESTs {IMAGE:479846}	479846 IMAGE Consortium	AA051480
7696 F	8 021ZA1F9	ESTs {IMAGE:480050}	480050 IMAGE Consortium	AA051515
7697 F	10 021ZA1F9	Tenascin X {IMAGE:480221}	480221 IMAGE Consortium	AA058085
7698 F	12 021ZA1F9	IMAGE EST {IMAGE:480462}	480462 IMAGE Consortium	AA058269
7699 H	2 021ZA1F9	IMAGE EST {IMAGE:479077}	479077 IMAGE Consortium	AA048878
7700 H	4 021ZA1F9	Mouse mRNA for meiosis-specific nuclear structural protein 1 (MNS1), complete cds {IMAGE:480118}	480118 IMAGE Consortium	AA057998
7701 H	6 021ZA1F9	ESTs {IMAGE:479287}	479287 IMAGE Consortium	AA048738
7702 H	8 021ZA1F9	IMAGE EST {IMAGE:480639}	480639 IMAGE Consortium	AA060697
7703 H	10 021ZA1F9	IMAGE EST {IMAGE:480404}	480404 IMAGE Consortium	AA058302
7704 H	12 021ZA1F9	Protein kinase C substrate 80K-H {IMAGE:479665}	479665 IMAGE Consortium	AA049206
7705 B	2 021JA1FA	IMAGE EST {IMAGE:482943}	482943 IMAGE Consortium	AA062182
7706 B	4 021JA1FA	ESTs, Highly similar to NUCLEAR FACTOR NF-KAPPA-B P100 SUBUNIT [Homo sapiens] {IMAGE:482952}	482952 IMAGE Consortium	AA060802
7707 B	6 021JA1FA	ESTs {IMAGE:483320}	483320 IMAGE Consortium	AA061586
7708 B	8 021JA1FA	Zinc finger protein 147 {IMAGE:483232}	483232 IMAGE Consortium	AA063824
7709 B	10 021JA1FA	IMAGE EST {IMAGE:483309}	483309 IMAGE Consortium	AA061630
7710 B	12 021JA1FA	ESTs {IMAGE:481481}	481481 IMAGE Consortium	AA060306
7711 D	2 021JA1FA	IMAGE EST {IMAGE:482771}	482771 IMAGE Consortium	AA062034
7712 D	4 021JA1FA	Cytochrome P450, 2e1, ethanol inducible {IMAGE:482632}	482632 IMAGE Consortium	AA062043
7713 D	6 021JA1FA	IMAGE EST {IMAGE:483024}	483024 IMAGE Consortium	AA060880
7714 D	8 021JA1FA	IMAGE EST {IMAGE:481521}	481521 IMAGE Consortium	AA060594
7715 D	10 021JA1FA	ESTs {IMAGE:483628}	483628 IMAGE Consortium	AA061286

7716 D	12 021JA1FA	ESTs, Highly similar to nuclear protein Skip [H.sapiens] {IMAGE:481872}	481872 IMAGE Consortium	AA059626
7717 F	2 021JA1FA	IMAGE EST {IMAGE:481234}	481234 IMAGE Consortium	AA060385
7718 F	4 021JA1FA	IMAGE EST {IMAGE:481142}	481142 IMAGE Consortium	AA064311
7719 F	6 021JA1FA	IMAGE EST {IMAGE:483210}	483210 IMAGE Consortium	AA063801
7720 F	8 021JA1FA	Mouse housekeeping DXS253E (P3) and DXS254E (GdX) genes, 3' end, and complete cds {IMAGE:483262}	483262 IMAGE Consortium	AA061673
7721 F	10 021JA1FA	Mus musculus short-chain dehydrogenase CRAD2 mRNA, complete cds {IMAGE:483701}	483701 IMAGE Consortium	AA061397
7722 F	12 021JA1FA	Cytochrome P450, 4a14 {IMAGE:483348}	483348 IMAGE Consortium	AA061737
7723 H	2 021JA1FA	IMAGE EST {IMAGE:481307}	481307 IMAGE Consortium	AA060448
7724 H	4 021JA1FA	ESTs {IMAGE:483002}	483002 IMAGE Consortium	AA060856
7725 H	6 021JA1FA	IMAGE EST {IMAGE:483177}	483177 IMAGE Consortium	AA060995
7726 H	8 021JA1FA	IMAGE EST {IMAGE:481450}	481450 IMAGE Consortium	AA060579
7727 H	10 021JA1FA	IMAGE EST {IMAGE:482136}	482136 IMAGE Consortium	AA059642
7728 H	12 021JA1FA	Mus musculus mRNA for Mad-related protein Smad7 {IMAGE:483550}	483550 IMAGE Consortium	AA061644
7729 B	2 021QA1FB	ESTs {IMAGE:479311}	479311 IMAGE Consortium	AA048769
7730 B	4 021QA1FB	EST {IMAGE:479316}	479316 IMAGE Consortium	AA048764
7731 B	6 021QA1FB	ESTs {IMAGE:480428}	480428 IMAGE Consortium	AA058310
7732 B	8 021QA1FB	IMAGE EST {IMAGE:480431}	480431 IMAGE Consortium	AA058137
7733 B	10 021QA1FB	ESTs {IMAGE:479713}	479713 IMAGE Consortium	AA051532
7734 B	12 021QA1FB	ESTs {IMAGE:478428}	478428 IMAGE Consortium	AA049819
7735 D	2 021QA1FB	ESTs {IMAGE:479364}	479364 IMAGE Consortium	AA048795
7736 D	4 021QA1FB	Cytochrome P450, 2e1, ethanol inducible {IMAGE:479487}	479487 IMAGE Consortium	AA049267
7737 D	6 021QA1FB	Mus musculus branched chain alpha ketoacid dehydrogenase kinase (bckd-kinase) mRNA, nuclear gene encoding mitochondrial protein, complete cds {IMAGE:480575}	480575 IMAGE Consortium	AA059497
7738 D	8 021QA1FB	ESTs {IMAGE:478452}	478452 IMAGE Consortium	AA049979
7739 D	10 021QA1FB	Tumor necrosis factor induced protein 6 {IMAGE:478579}	478579 IMAGE Consortium	AA051341
7740 D	12 021QA1FB	ACTIN, AORTIC SMOOTH MUSCLE {IMAGE:480128}	480128 IMAGE Consortium	AA051790
7741 F	2 021QA1FB	ESTs {IMAGE:478476}	478476 IMAGE Consortium	AA050001
7742 F	4 021QA1FB	ESTs, Highly similar to C-1-TETRAHYDROFOLATE SYNTHASE, CYTOPLASMIC [Rattus norvegicus] {IMAGE:478383}	478383 IMAGE Consortium	AA049796
7743 F	6 021QA1FB	ESTs, Highly similar to Similar to S.cerevisiae hypothetical protein N0330 [H.sapiens] {IMAGE:479759}	479759 IMAGE Consortium	AA051679
7744 F	8 021QA1FB	ESTs, Highly similar to angio-associated migratory cell protein [H.sapiens] {IMAGE:480176}	480176 IMAGE Consortium	AA058182
7745 F	10 021QA1FB	IMAGE EST {IMAGE:480181}	480181 IMAGE Consortium	AA058187
7746 F	12 021QA1FB	IMAGE EST {IMAGE:478744}	478744 IMAGE Consortium	AA051156
7747 H	2 021QA1FB	ESTs {IMAGE:482626}	482626 IMAGE Consortium	AA061950
7748 H	4 021QA1FB	Alpha 1 microglobulin/bikunin {IMAGE:482830}	482830 IMAGE Consortium	AA062198

7749 H	6 021QA1FB	ESTs {IMAGE:481712}	481712 IMAGE Consortium	AA060114
7750 H	8 021QA1FB	IMAGE EST {IMAGE:481548}	481548 IMAGE Consortium	AA060601
7751 H	10 021QA1FB	IMAGE EST {IMAGE:481433}	481433 IMAGE Consortium	AA060293
7752 H	12 021QA1FB	ESTs, Moderately similar to CLATHRIN LIGHT CHAIN B [Bos taurus] {IMAGE:481655}	481655 IMAGE Consortium	AA060277
7753 B	2 021XA1FC	IMAGE EST {IMAGE:482995}	482995 IMAGE Consortium	AA060863
7754 B	4 021XA1FC	IMAGE EST {IMAGE:481765}	481765 IMAGE Consortium	AA060074
7755 B	6 021XA1FC	IMAGE EST {IMAGE:481969}	481969 IMAGE Consortium	AA059959
7756 B	8 021XA1FC	M.musculus mRNA for NfiB3-protein, splice variant (exon 9, 10 and 11 missing) {IMAGE:481927}	481927 IMAGE Consortium	AA059930
7757 B	10 021XA1FC	Mus musculus Keap1 mRNA, complete cds {IMAGE:482028}	482028 IMAGE Consortium	AA059890
7758 B	12 021XA1FC	Fatty acid transport protein {IMAGE:482059}	482059 IMAGE Consortium	AA059983
7759 D	2 021XA1FC	ESTs {IMAGE:483368}	483368 IMAGE Consortium	AA061710
7760 D	4 021XA1FC	ESTs {IMAGE:481898}	481898 IMAGE Consortium	AA059876
7761 D	6 021XA1FC	ESTs {IMAGE:483695}	483695 IMAGE Consortium	AA061216
7762 D	8 021XA1FC	IMAGE EST {IMAGE:483758}	483758 IMAGE Consortium	AA061404
7763 D	10 021XA1FC	ESTs {IMAGE:480893}	480893 IMAGE Consortium	AA064183
7764 D	12 021XA1FC	ESTs, Highly similar to SODIUM/GLUCOSE COTRANSPORTER [Sus scrofa] {IMAGE:480901}	480901 IMAGE Consortium	AA064246
7765 F	2 021XA1FC	Erythrocyte protein band 7.2 {IMAGE:481880}	481880 IMAGE Consortium	AA059683
7766 F	4 021XA1FC	ESTs {IMAGE:483735}	483735 IMAGE Consortium	AA061276
7767 F	6 021XA1FC	ESTs, Moderately similar to ENDOTHELIAL ACTIN-BINDING PROTEIN [Homo sapiens] {IMAGE:484301}	484301 IMAGE Consortium	AA080279
7768 F	8 021XA1FC	ESTs, Weakly similar to similar to nucleotide translocator [C.elegans] {IMAGE:480706}	480706 IMAGE Consortium	AA064156
7769 F	10 021XA1FC	ESTs {IMAGE:482641}	482641 IMAGE Consortium	AA061982
7770 F	12 021XA1FC	IMAGE EST {IMAGE:482817}	482817 IMAGE Consortium	AA062120
7771 H	2 021XA1FC	ESTs, Highly similar to HYPOTHETICAL 70.2 KD PROTEIN IN GSH1-CHS6 INTERGENIC REGION [Saccharomyces cerevisiae] {IMAGE:481947}	481947 IMAGE Consortium	AA059952
7772 H	4 021XA1FC	ESTs, Moderately similar to JAK3 [M.musculus] {IMAGE:481857}	481857 IMAGE Consortium	AA059785
7773 H	6 021XA1FC	Mus musculus signal recognition particle receptor beta subunit mRNA, complete cds {IMAGE:482029}	482029 IMAGE Consortium	AA059893
7774 H	8 021XA1FC	ESTs, Weakly similar to C01A2.4 [C.elegans] {IMAGE:482328}	482328 IMAGE Consortium	AA060898
7775 H	10 021XA1FC	EST {IMAGE:480798}	480798 IMAGE Consortium	AA060715
7776 H	12 021XA1FC	IMAGE EST {IMAGE:480994}	480994 IMAGE Consortium	AA060190
7777 B	2 0214A1FD	ESTs {IMAGE:481701}	481701 IMAGE Consortium	AA060287
7778 B	4 0214A1FD	Cytochrome P450, 2e1, ethanol inducible {IMAGE:483404}	483404 IMAGE Consortium	AA061873
7779 B	6 0214A1FD	IMAGE EST {IMAGE:483750}	483750 IMAGE Consortium	AA061278
7780 B	8 0214A1FD	IMAGE EST {IMAGE:480731}	480731 IMAGE Consortium	AA064166
7781 B	10 0214A1FD	ESTs {IMAGE:481227}	481227 IMAGE Consortium	AA060389

7782 B	12 0214A1FD	N-acetyl galactosaminidase, alpha {IMAGE:481351}	481351 IMAGE Consortium	AA060459
7783 D	2 0214A1FD	ESTs {IMAGE:484254}	484254 IMAGE Consortium	AA080143
7784 D	4 0214A1FD	TALIN {IMAGE:480684}	480684 IMAGE Consortium	AA064134
7785 D	6 0214A1FD	ESTs {IMAGE:482882}	482882 IMAGE Consortium	AA062142
7786 D	8 0214A1FD	Tubulin, beta 2 {IMAGE:482602}	482602 IMAGE Consortium	AA061838
7787 D	10 0214A1FD	Paired-like homeodomain transcription factor 3 {IMAGE:482871}	482871 IMAGE Consortium	AA062140
7788 D	12 0214A1FD	ESTs {IMAGE:481362}	481362 IMAGE Consortium	AA060545
7789 F	2 0214A1FD	ESTs {IMAGE:571419}	571419 IMAGE Consortium	AA109017
7790 F	4 0214A1FD	ESTs {IMAGE:571643}	571643 IMAGE Consortium	AA108880
7791 F	6 0214A1FD	Secreted phosphoprotein 1 {IMAGE:571759}	571759 IMAGE Consortium	AA108928
7792 F	8 0214A1FD	ESTs {IMAGE:571816}	571816 IMAGE Consortium	AA109854
7793 F	10 0214A1FD	IMAGE EST {IMAGE:572386}	572386 IMAGE Consortium	AA108501
7794 F	12 0214A1FD	Microtubule-associated protein tau {IMAGE:493423}	493423 IMAGE Consortium	AA086907
7795 H	2 0214A1FD	ESTs, Highly similar to SUCCINATE DEHYDROGENASE [Homo sapiens] {IMAGE:572339}	572339 IMAGE Consortium	AA108475
7796 H	4 0214A1FD	IMAGE EST {IMAGE:573075}	573075 IMAGE Consortium	AA118709
7797 H	6 0214A1FD	IMAGE EST {IMAGE:572986}	572986 IMAGE Consortium	AA111560
7798 H	8 0214A1FD	ESTs, Moderately similar to ACTIVATOR 1 36 KD SUBUNIT [Homo sapiens] {IMAGE:572970}	572970 IMAGE Consortium	AA111689
7799 H	10 0214A1FD	Annexin III {IMAGE:573265}	573265 IMAGE Consortium	AA119121
7800 H	12 0214A1FD	ESTs, Highly similar to ATP SYNTHASE EPSILON CHAIN, MITOCHONDRIAL PRECURSOR [Bos taurus] {IMAGE:571214}	571214 IMAGE Consortium	AA108733
7801 B	2 021BA1FE	IMAGE EST {IMAGE:492439}	492439 IMAGE Consortium	AA097213
7802 B	4 021BA1FE	IMAGE EST {IMAGE:571955}	571955 IMAGE Consortium	AA108342
7803 B	6 021BA1FE	ESTs {IMAGE:493205}	493205 IMAGE Consortium	AA097495
7804 B	8 021BA1FE	ESTs, Weakly similar to unknown [S.cerevisiae] {IMAGE:572718}	572718 IMAGE Consortium	AA111661
7805 B	10 021BA1FE	Mus musculus Oct binding factor 1 (OBF-1) mRNA, complete cds {IMAGE:572880}	572880 IMAGE Consortium	AA110709
7806 B	12 021BA1FE	ESTs {IMAGE:493434}	493434 IMAGE Consortium	AA086952
7807 D	2 021BA1FE	Neurofibromatosis 2 {IMAGE:572275}	572275 IMAGE Consortium	AA105153
7808 D	4 021BA1FE	ESTs {IMAGE:572601}	572601 IMAGE Consortium	AA111003
7809 D	6 021BA1FE	FMS-like tyrosine kinase 4 {IMAGE:493563}	493563 IMAGE Consortium	AA086840
7810 D	8 021BA1FE	ESTs {IMAGE:493451}	493451 IMAGE Consortium	AA087038
7811 D	10 021BA1FE	Histidine decarboxylase cluster {IMAGE:573039}	573039 IMAGE Consortium	AA118747
7812 D	12 021BA1FE	Adrenergic receptor, beta 2 {IMAGE:573046}	573046 IMAGE Consortium	AA111732
7813 F	2 021BA1FE	ESTs {IMAGE:573063}	573063 IMAGE Consortium	AA118422
7814 F	4 021BA1FE	IMAGE EST {IMAGE:572879}	572879 IMAGE Consortium	AA110346
7815 F	6 021BA1FE	Histocompatibility 2, O region alpha locus {IMAGE:573556}	573556 IMAGE Consortium	AA118886
7816 F	8 021BA1FE	IMAGE EST {IMAGE:573227}	573227 IMAGE Consortium	AA119357
7817 F	10 021BA1FE	IMAGE EST {IMAGE:484289}	484289 IMAGE Consortium	AA073833

7818 F	12 021BA1FE	Mus musculus E1B 19K/Bcl-2-binding protein homolog (Nip3) mRNA, nuclear gene encoding mitochondrial protein, complete cds {IMAGE:571367}	571367 IMAGE Consortium	AA105295
7819 H	2 021BA1FE	ESTs {IMAGE:573395}	573395 IMAGE Consortium	AA118818
7820 H	4 021BA1FE	ESTs {IMAGE:573209}	573209 IMAGE Consortium	AA110934
7821 H	6 021BA1FE	IMAGE EST {IMAGE:484342}	484342 IMAGE Consortium	AA073756
7822 H	8 021BA1FE	Kinesin heavy chain member 1A {IMAGE:492514}	492514 IMAGE Consortium	AA097858
7823 H	10 021BA1FE	ESTs {IMAGE:572310}	572310 IMAGE Consortium	AA110273
7824 H	12 021BA1FE	ESTs, Weakly similar to secretory leukocyte protease inhibitor [M.musculus] {IMAGE:572422}	572422 IMAGE Consortium	AA105830
7825 B	2 021IA1FF	ESTs {IMAGE:573052}	573052 IMAGE Consortium	AA118681
7826 B	4 021IA1FF	ESTs {IMAGE:572819}	572819 IMAGE Consortium	AA110791
7827 B	6 021IA1FF	COPROPORPHYRINOGEN III OXIDASE PRECURSOR {IMAGE:570602}	570602 IMAGE Consortium	AA108600
7828 B	8 021IA1FF	ESTs, Highly similar to NEDD-4 PROTEIN [Homo sapiens] {IMAGE:573510}	573510 IMAGE Consortium	AA118878
7829 B	10 021IA1FF	ESTs, Highly similar to LEUCOCYTE ANTIGEN CD97 PRECURSOR [Homo sapiens] {IMAGE:573301}	573301 IMAGE Consortium	AA118715
7830 B	12 021IA1FF	Mus musculus gene for DJ-1, complete cds {IMAGE:571278}	571278 IMAGE Consortium	AA108826
7831 D	2 021IA1FF	IMAGE EST {IMAGE:493433}	493433 IMAGE Consortium	AA086829
7832 D	4 021IA1FF	IMAGE EST {IMAGE:572875}	572875 IMAGE Consortium	AA110344
7833 D	6 021IA1FF	ESTs {IMAGE:573338}	573338 IMAGE Consortium	AA118670
7834 D	8 021IA1FF	IMAGE EST {IMAGE:571219}	571219 IMAGE Consortium	AA107718
7835 D	10 021IA1FF	IMAGE EST {IMAGE:571398}	571398 IMAGE Consortium	AA108992
7836 D	12 021IA1FF	EST {IMAGE:492529}	492529 IMAGE Consortium	AA097333
7837 F	2 021IA1FF	ESTs, Weakly similar to HYPOTHETICAL PROTEIN F44G4.1 IN CHROMOSOME II [C.elegans] {IMAGE:573224}	573224 IMAGE Consortium	AA119353
7838 F	4 021IA1FF	ESTs {IMAGE:573302}	573302 IMAGE Consortium	AA118713
7839 F	6 021IA1FF	ESTs {IMAGE:571422}	571422 IMAGE Consortium	AA109015
7840 F	8 021IA1FF	ESTs {IMAGE:571503}	571503 IMAGE Consortium	AA109230
7841 F	10 021IA1FF	ESTs, Weakly similar to GS2 PROTEIN [H.sapiens] {IMAGE:492447}	492447 IMAGE Consortium	AA097194
7842 F	12 021IA1FF	Mus musculus WDR protein mRNA, complete cds {IMAGE:493109}	493109 IMAGE Consortium	AA097808
7843 H	2 021IA1FF	Cytochrome P450, 4a10 {IMAGE:570575}	570575 IMAGE Consortium	AA109684
7844 H	4 021IA1FF	ESTs, Highly similar to VACUOLAR ATP SYNTHASE SUBUNIT D [Bos taurus] {IMAGE:571005}	571005 IMAGE Consortium	AA109695
7845 H	6 021IA1FF	ESTs {IMAGE:484118}	484118 IMAGE Consortium	AA080048
7846 H	8 021IA1FF	IMAGE EST {IMAGE:484315}	484315 IMAGE Consortium	AA073751
7847 H	10 021IA1FF	Glutathione S-transferase, pi 2 {IMAGE:572002}	572002 IMAGE Consortium	AA108370
7848 H	12 021IA1FF	Laminin, alpha 2 {IMAGE:493133}	493133 IMAGE Consortium	AA097476
7849 B	2 021PA1FG	IMAGE EST {IMAGE:575247}	575247 IMAGE Consortium	AA125390
7850 B	4 021PA1FG	IMAGE EST {IMAGE:575776}	575776 IMAGE Consortium	AA120429



7851 B	6 021PA1FG	IMAGE EST {IMAGE:574657}	574657 IMAGE Consortium	AA125257
7852 B	8 021PA1FG	ESTs {IMAGE:574351}	574351 IMAGE Consortium	AA119543
7853 B	10 021PA1FG	IMAGE EST {IMAGE:574500}	574500 IMAGE Consortium	AA125110
7854 B	12 021PA1FG	ESTs {IMAGE:576353}	576353 IMAGE Consortium	AA119478
7855 D	2 021PA1FG	IMAGE EST {IMAGE:574042}	574042 IMAGE Consortium	AA118998
7856 D	4 021PA1FG	IMAGE EST {IMAGE:574245}	574245 IMAGE Consortium	AA111234
7857 D	6 021PA1FG	ESTs, Weakly similar to similar to D. melanogaster trithorax protein [C.elegans] {IMAGE:575899}	575899 IMAGE Consortium	AA117603
7858 D	8 021PA1FG	IMAGE EST {IMAGE:576459}	576459 IMAGE Consortium	AA123781
7859 D	10 021PA1FG	ESTs {IMAGE:574459}	574459 IMAGE Consortium	AA118312
7860 D	12 021PA1FG	Downstream of tyrosine kinase 1 {IMAGE:576287}	576287 IMAGE Consortium	AA119595
7861 F	2 021PA1FG	ESTs, Weakly similar to weak similarity to SP:YAD5_CLOAB [C.elegans] {IMAGE:574234}	574234 IMAGE Consortium	AA119401
7862 F	4 021PA1FG	ESTs, Moderately similar to This sequence is almost identical to HUMRSC433, D13632. [H.sapiens] {IMAGE:576156}	576156 IMAGE Consortium	AA119802
7863 F	6 021PA1FG	IMAGE EST {IMAGE:576068}	576068 IMAGE Consortium	AA119161
7864 F	8 021PA1FG	Mus musculus TNF-response element binding protein mRNA, complete cds {IMAGE:576531}	576531 IMAGE Consortium	AA123821
7865 F	10 021PA1FG	Mus musculus major histocompatibility locus class III region: complement C4 (C4) and cytochrome P450 hydroxylase A (CYP21OH-A) genes, complete cds; slp pseudogene, complete sequence; NG6, SKI, and complement factor B (Bf) genes, c {IMAGE:576545}	576545 IMAGE Consortium	AA117411
7866 F	12 021PA1FG	PCTAIRE-motif protein kinase 1 {IMAGE:576407}	576407 IMAGE Consortium	AA119679
7867 H	2 021PA1FG	IMAGE EST {IMAGE:574586}	574586 IMAGE Consortium	AA124750
7868 H	4 021PA1FG	IMAGE EST {IMAGE:574495}	574495 IMAGE Consortium	AA117628
7869 H	6 021PA1FG	Mus musculus Hsp70-related NST-1 (hsr.1) mRNA, complete cds {IMAGE:574381}	574381 IMAGE Consortium	AA119654
7870 H	8 021PA1FG	IMAGE EST {IMAGE:574725}	574725 IMAGE Consortium	AA120657
7871 H	10 021PA1FG	ESTs, Highly similar to HYPOTHETICAL 67.2 KD PROTEIN T16H12.4 IN CHROMOSOME III [Caenorhabditis elegans] {IMAGE:575067}	575067 IMAGE Consortium	AA125364
7872 H	12 021PA1FG	IMAGE EST {IMAGE:574909}	574909 IMAGE Consortium	AA124907
7873 B	2 021WA1FH	IMAGE EST {IMAGE:575998}	575998 IMAGE Consortium	AA119141
7874 B	4 021WA1FH	IMAGE EST {IMAGE:576436}	576436 IMAGE Consortium	AA120358
7875 B	6 021WA1FH	IMAGE EST {IMAGE:574439}	574439 IMAGE Consortium	AA118291
7876 B	8 021WA1FH	ESTs, Highly similar to zinc finger protein C2H2-171 [H.sapiens] {IMAGE:574756}	574756 IMAGE Consortium	AA124575
7877 B	10 021WA1FH	IMAGE EST {IMAGE:574957}	574957 IMAGE Consortium	AA125282
7878 B	12 021WA1FH	IMAGE EST {IMAGE:574747}	574747 IMAGE Consortium	AA118337
7879 D	2 021WA1FH	Mus musculus mRNA for SMRP/MRP5, partial cds {IMAGE:576245}	576245 IMAGE Consortium	AA119896
7880 D	4 021WA1FH	ESTs {IMAGE:574395}	574395 IMAGE Consortium	AA119577
7881 D	6 021WA1FH	IMAGE EST {IMAGE:574711}	574711 IMAGE Consortium	AA125274

7882 D	8 021WA1FH	ESTs {IMAGE:574607}	574607 IMAGE Consortium	AA118379
7883 D	10 021WA1FH	IMAGE EST {IMAGE:574948}	574948 IMAGE Consortium	AA124983
7884 D	12 021WA1FH	ESTs, Highly similar to PROBABLE RIBOSOMAL PROTEIN B0303.15 IN CHROMOSOME III [Caenorhabditis elegans] {IMAGE:574981}	574981 IMAGE Consortium	AA125287
7885 F	2 021WA1FH	ESTs {IMAGE:576216}	576216 IMAGE Consortium	AA119887
7886 F	4 021WA1FH	Cyclin B1, related sequence 1 {IMAGE:576166}	576166 IMAGE Consortium	AA124592
7887 F	6 021WA1FH	ESTs {IMAGE:574430}	574430 IMAGE Consortium	AA119664
7888 F	8 021WA1FH	IMAGE EST {IMAGE:575045}	575045 IMAGE Consortium	AA125359
7889 F	10 021WA1FH	IMAGE EST {IMAGE:576823}	576823 IMAGE Consortium	AA124419
7890 F	12 021WA1FH	IMAGE EST {IMAGE:576974}	576974 IMAGE Consortium	AA124340
7891 H	2 021WA1FH	ESTs {IMAGE:574635}	574635 IMAGE Consortium	AA125252
7892 H	4 021WA1FH	IMAGE EST {IMAGE:576558}	576558 IMAGE Consortium	AA124116
7893 H	6 021WA1FH	IMAGE EST {IMAGE:576428}	576428 IMAGE Consortium	AA119544
7894 H	8 021WA1FH	ESTs {IMAGE:577010}	577010 IMAGE Consortium	AA122546
7895 H	10 021WA1FH	IMAGE EST {IMAGE:576943}	576943 IMAGE Consortium	AA122529
7896 H	12 021WA1FH	IMAGE EST {IMAGE:576958}	576958 IMAGE Consortium	AA116447
7897 B	2 0213A1FI	ESTs {IMAGE:577413}	577413 IMAGE Consortium	AA116640
7898 B	4 0213A1FI	IMAGE EST {IMAGE:577786}	577786 IMAGE Consortium	AA139231
7899 B	6 0213A1FI	ESTs {IMAGE:577720}	577720 IMAGE Consortium	AA140537
7900 B	8 0213A1FI	IMAGE EST {IMAGE:581794}	581794 IMAGE Consortium	AA153900
7901 B	10 0213A1FI	Itchy {IMAGE:582175}	582175 IMAGE Consortium	AA153909
7902 B	12 0213A1FI	IMAGE EST {IMAGE:577156}	577156 IMAGE Consortium	AA124525
7903 D	2 0213A1FI	ESTs, Moderately similar to SUPEROXIDE DISMUTASE [Brassica oleracea var. capitata] {IMAGE:578433}	578433 IMAGE Consortium	AA123360
7904 D	4 0213A1FI	ESTs {IMAGE:579509}	579509 IMAGE Consortium	AA124332
7905 D	6 0213A1FI	Dlk1-like homolog (Drosophila) {IMAGE:579523}	579523 IMAGE Consortium	AA124612
7906 D	8 0213A1FI	Proliferating cell nuclear antigen {IMAGE:579735}	579735 IMAGE Consortium	AA116947
7907 D	10 0213A1FI	IMAGE EST {IMAGE:577180}	577180 IMAGE Consortium	AA124531
7908 D	12 0213A1FI	Lymphocyte antigen 6 complex {IMAGE:580715}	580715 IMAGE Consortium	AA145865
7909 F	2 0213A1FI	ESTs {IMAGE:581115}	581115 IMAGE Consortium	AA140001
7910 F	4 0213A1FI	IMAGE EST {IMAGE:581124}	581124 IMAGE Consortium	AA140118
7911 F	6 0213A1FI	ESTs {IMAGE:577555}	577555 IMAGE Consortium	AA138025
7912 F	8 0213A1FI	IMAGE EST {IMAGE:577808}	577808 IMAGE Consortium	AA122957
7913 F	10 0213A1FI	ESTs, Weakly similar to 6-PHOSPHOGLUCONATE DEHYDROGENASE, DECARBOXYLATING [Ovis aries] {IMAGE:578436}	578436 IMAGE Consortium	AA122658
7914 F	12 0213A1FI	IMAGE EST {IMAGE:578299}	578299 IMAGE Consortium	AA120312
7915 H	2 0213A1FI	IMAGE EST {IMAGE:581956}	581956 IMAGE Consortium	AA152987
7916 H	4 0213A1FI	ESTs {IMAGE:582063}	582063 IMAGE Consortium	AA154227
7917 H	6 0213A1FI	ESTs, Moderately similar to tetratricopeptide repeat protein [H.sapiens] {IMAGE:582743}	582743 IMAGE Consortium	AA142490
7918 H	8 0213A1FI	ESTs {IMAGE:577409}	577409 IMAGE Consortium	AA116618

7919 H	10 0213A1FI	IMAGE EST {IMAGE:580924}	580924 IMAGE Consortium	AA139817
7920 H	12 0213A1FI	IMAGE EST {IMAGE:577598}	577598 IMAGE Consortium	AA139452
7921 B	2 021AA1FJ	ESTs {IMAGE:574527}	574527 IMAGE Consortium	AA125185
7922 B	4 021AA1FJ	IMAGE EST {IMAGE:576524}	576524 IMAGE Consortium	AA117421
7923 B	6 021AA1FJ	IMAGE EST {IMAGE:576675}	576675 IMAGE Consortium	AA123772
7924 B	8 021AA1FJ	Mus musculus U4/U6 snRNP 90 kDa protein gene, complete cds {IMAGE:576991}	576991 IMAGE Consortium	AA122544
7925 B	10 021AA1FJ	ESTs {IMAGE:576664}	576664 IMAGE Consortium	AA119853
7926 B	12 021AA1FJ	ESTs {IMAGE:576767}	576767 IMAGE Consortium	AA120015
7927 D	2 021AA1FJ	ESTs, Highly similar to SMT3B protein [H.sapiens] {IMAGE:574897}	574897 IMAGE Consortium	AA125200
7928 D	4 021AA1FJ	ESTs {IMAGE:576891}	576891 IMAGE Consortium	AA124157
7929 D	6 021AA1FJ	ESTs {IMAGE:574821}	574821 IMAGE Consortium	AA120697
7930 D	8 021AA1FJ	Lymphotactin {IMAGE:576815}	576815 IMAGE Consortium	AA120030
7931 D	10 021AA1FJ	ESTs {IMAGE:573849}	573849 IMAGE Consortium	AA119049
7932 D	12 021AA1FJ	ESTs, Highly similar to RSP-1 PROTEIN [Mus musculus] {IMAGE:574248}	574248 IMAGE Consortium	AA119463
7933 F	2 021AA1FJ	IMAGE EST {IMAGE:573770}	573770 IMAGE Consortium	AA118841
7934 F	4 021AA1FJ	ESTs {IMAGE:573634}	573634 IMAGE Consortium	AA118853
7935 F	6 021AA1FJ	IMAGE EST {IMAGE:573929}	573929 IMAGE Consortium	AA111044
7936 F	8 021AA1FJ	IMAGE EST {IMAGE:575575}	575575 IMAGE Consortium	AA124959
7937 F	10 021AA1FJ	Ferritin heavy chain {IMAGE:576076}	576076 IMAGE Consortium	AA124593
7938 F	12 021AA1FJ	ESTs {IMAGE:574633}	574633 IMAGE Consortium	AA125251
7939 H	2 021AA1FJ	ESTs {IMAGE:576954}	576954 IMAGE Consortium	AA116445
7940 H	4 021AA1FJ	IMAGE EST {IMAGE:577058}	577058 IMAGE Consortium	AA124459
7941 H	6 021AA1FJ	ESTs {IMAGE:577043}	577043 IMAGE Consortium	AA124425
7942 H	8 021AA1FJ	IMAGE EST {IMAGE:577762}	577762 IMAGE Consortium	AA139225
7943 H	10 021AA1FJ	IMAGE EST {IMAGE:577528}	577528 IMAGE Consortium	AA138012
7944 H	12 021AA1FJ	IMAGE EST {IMAGE:578425}	578425 IMAGE Consortium	AA124104
7945 B	2 021HA1FK	ESTs {IMAGE:582936}	582936 IMAGE Consortium	AA153601
7946 B	4 021HA1FK	IMAGE EST {IMAGE:580302}	580302 IMAGE Consortium	AA137523
7947 B	6 021HA1FK	Ferritin heavy chain {IMAGE:580197}	580197 IMAGE Consortium	AA124396
7948 B	8 021HA1FK	Mus musculus orphan transporter isoform B11 (Xtrp2) mRNA, alternatively spliced, complete cds {IMAGE:581020}	581020 IMAGE Consortium	AA139706
7949 B	10 021HA1FK	Mus musculus Ets-family transcription factor Elf-1 mRNA, complete cds {IMAGE:581665}	581665 IMAGE Consortium	AA152626
7950 B	12 021HA1FK	ESTs {IMAGE:581813}	581813 IMAGE Consortium	AA153861
7951 D	2 021HA1FK	IMAGE EST {IMAGE:577766}	577766 IMAGE Consortium	AA139227
7952 D	4 021HA1FK	ESTs {IMAGE:577654}	577654 IMAGE Consortium	AA140517
7953 D	6 021HA1FK	IMAGE EST {IMAGE:577978}	577978 IMAGE Consortium	AA124168
7954 D	8 021HA1FK	IMAGE EST {IMAGE:582169}	582169 IMAGE Consortium	AA153907
7955 D	10 021HA1FK	IMAGE EST {IMAGE:582261}	582261 IMAGE Consortium	AA146090
7956 D	12 021HA1FK	ESTs {IMAGE:582465}	582465 IMAGE Consortium	AA153039

7957 F	2 021HA1FK	ESTs {IMAGE:581834}	581834 IMAGE Consortium	AA154423
7958 F	4 021HA1FK	Mus musculus golgin-245 (olp-1) mRNA, complete cds {IMAGE:581671}	581671 IMAGE Consortium	AA152629
7959 F	6 021HA1FK	IMAGE EST {IMAGE:582074}	582074 IMAGE Consortium	AA154187
7960 F	8 021HA1FK	ESTs, Weakly similar to p76 [H.sapiens] {IMAGE:582571}	582571 IMAGE Consortium	AA144167
7961 F	10 021HA1FK	IMAGE EST {IMAGE:580247}	580247 IMAGE Consortium	AA124362
7962 F	12 021HA1FK	IMAGE EST {IMAGE:581094}	581094 IMAGE Consortium	AA139949
7963 H	2 021HA1FK	ESTs, Highly similar to PROTEIN TRANSPORT PROTEIN SEC13 [Saccharomyces cerevisiae] {IMAGE:580259}	580259 IMAGE Consortium	AA137370
7964 H	4 021HA1FK	Tissue inhibitor of metalloproteinase 3 {IMAGE:580753}	580753 IMAGE Consortium	AA145784
7965 H	6 021HA1FK	ESTs, Highly similar to CYTOCHROME C OXIDASE POLYPEPTIDE VIB [Bos taurus] {IMAGE:581175}	581175 IMAGE Consortium	AA139624
7966 H	8 021HA1FK	IMAGE EST {IMAGE:577852}	577852 IMAGE Consortium	AA124777
7967 H	10 021HA1FK	ESTs {IMAGE:578496}	578496 IMAGE Consortium	AA162895
7968 H	12 021HA1FK	ESTs, Weakly similar to VITAMIN K-DEPENDENT GAMMA-CARBOXYLASE [Homo sapiens] {IMAGE:582706}	582706 IMAGE Consortium	AA154146
7969 B	2 021OA1FL	ESTs {IMAGE:583943}	583943 IMAGE Consortium	AA145721
7970 B	4 021OA1FL	ESTs {IMAGE:596213}	596213 IMAGE Consortium	AA137829
7971 B	6 021OA1FL	ESTs {IMAGE:598305}	598305 IMAGE Consortium	AA164095
7972 B	8 021OA1FL	IMAGE EST {IMAGE:598289}	598289 IMAGE Consortium	AA155356
7973 B	10 021OA1FL	IMAGE EST {IMAGE:598826}	598826 IMAGE Consortium	AA168417
7974 B	12 021OA1FL	ESTs {IMAGE:598739}	598739 IMAGE Consortium	AA168435
7975 D	2 021OA1FL	Cytochrome P450, 2e1, ethanol inducible {IMAGE:597574}	597574 IMAGE Consortium	AA144393
7976 D	4 021OA1FL	IMAGE EST {IMAGE:598085}	598085 IMAGE Consortium	AA265372
7977 D	6 021OA1FL	IMAGE EST {IMAGE:597883}	597883 IMAGE Consortium	AA163649
7978 D	8 021OA1FL	Mus musculus mRNA, complete cds {IMAGE:598204}	598204 IMAGE Consortium	AA161816
7979 D	10 021OA1FL	IMAGE EST {IMAGE:598426}	598426 IMAGE Consortium	AA163966
7980 D	12 021OA1FL	Mus musculus ZW10 homolog (MmZW10) mRNA, partial cds {IMAGE:598218}	598218 IMAGE Consortium	AA163727
7981 F	2 021OA1FL	ESTs {IMAGE:597688}	597688 IMAGE Consortium	AA144356
7982 F	4 021OA1FL	EST {IMAGE:597742}	597742 IMAGE Consortium	AA144537
7983 F	6 021OA1FL	ESTs, Weakly similar to HYPOTHETICAL 67.0 KD PROTEIN IN PRE3-SAG1 INTERGENIC REGION [Saccharomyces cerevisiae] {IMAGE:597896}	597896 IMAGE Consortium	AA162148
7984 F	8 021OA1FL	IMAGE EST {IMAGE:598123}	598123 IMAGE Consortium	AA265255
7985 F	10 021OA1FL	Ryanodine receptor 1, skeletal muscle {IMAGE:598468}	598468 IMAGE Consortium	AA162446
7986 F	12 021OA1FL	IMAGE EST {IMAGE:596473}	596473 IMAGE Consortium	AA153794
7987 H	2 021OA1FL	IMAGE EST {IMAGE:583922}	583922 IMAGE Consortium	AA154540
7988 H	4 021OA1FL	IMAGE EST {IMAGE:597736}	597736 IMAGE Consortium	AA144535
7989 H	6 021OA1FL	ESTs {IMAGE:583652}	583652 IMAGE Consortium	AA146200
7990 H	8 021OA1FL	Mus musculus double-stranded RNA-specific adenosine deaminase mRNA, complete cds {IMAGE:595849}	595849 IMAGE Consortium	AA137946
7991 H	10 021OA1FL	IMAGE EST {IMAGE:596201}	596201 IMAGE Consortium	AA122763

7992 H	12 0210A1FL	Zinc finger protein 62 {IMAGE:596330}	596330 IMAGE Consortium	AA138529
7993 B	2 021VA1FM	ESTs {IMAGE:583566}	583566 IMAGE Consortium	AA143986
7994 B	4 021VA1FM	IMAGE EST {IMAGE:583809}	583809 IMAGE Consortium	AA153024
7995 B	6 021VA1FM	ESTs {IMAGE:583888}	583888 IMAGE Consortium	AA145822
7996 B	8 021VA1FM	IMAGE EST {IMAGE:583678}	583678 IMAGE Consortium	AA146275
7997 B	10 021VA1FM	ESTs {IMAGE:598464}	598464 IMAGE Consortium	AA162444
7998 B	12 021VA1FM	IMAGE EST {IMAGE:598326}	598326 IMAGE Consortium	AA162667
7999 D	2 021VA1FM	ESTs, Highly similar to PROTEIN 4.1 [Homo sapiens] {IMAGE:597748}	597748 IMAGE Consortium	AA153175
8000 D	4 021VA1FM	SIGNAL RECOGNITION PARTICLE 54 KD PROTEIN {IMAGE:597566}	597566 IMAGE Consortium	AA154373
8001 D	6 021VA1FM	Apoptosis inhibitor 2 {IMAGE:597715}	597715 IMAGE Consortium	AA144490
8002 D	8 021VA1FM	ESTs {IMAGE:598059}	598059 IMAGE Consortium	AA265364
8003 D	10 021VA1FM	ESTs, Highly similar to nucleoporin p54 [R.norvegicus] {IMAGE:595972}	595972 IMAGE Consortium	AA137295
8004 D	12 021VA1FM	ESTs {IMAGE:596027}	596027 IMAGE Consortium	AA137485
8005 F	2 021VA1FM	ESTs {IMAGE:597614}	597614 IMAGE Consortium	AA152944
8006 F	4 021VA1FM	ESTs, Highly similar to cyclic nucleotide phosphodiesterase [M.musculus] {IMAGE:597524}	597524 IMAGE Consortium	AA144377
8007 F	6 021VA1FM	Transcription factor CP2 {IMAGE:597820}	597820 IMAGE Consortium	AA164013
8008 F	8 021VA1FM	ESTs {IMAGE:597832}	597832 IMAGE Consortium	AA163421
8009 F	10 021VA1FM	ESTs {IMAGE:598274}	598274 IMAGE Consortium	AA162652
8010 F	12 021VA1FM	ESTs {IMAGE:598432}	598432 IMAGE Consortium	AA155074
8011 H	2 021VA1FM	IMAGE EST {IMAGE:597399}	597399 IMAGE Consortium	AA153205
8012 H	4 021VA1FM	Hemopoietic cell phosphatase {IMAGE:597644}	597644 IMAGE Consortium	AA144286
8013 H	6 021VA1FM	IMAGE EST {IMAGE:583823}	583823 IMAGE Consortium	AA146391
8014 H	8 021VA1FM	IMAGE EST {IMAGE:596141}	596141 IMAGE Consortium	AA137839
8015 H	10 021VA1FM	ESTs {IMAGE:596030}	596030 IMAGE Consortium	AA137447
8016 H	12 021VA1FM	ESTs, Highly similar to BN51 PROTEIN [Homo sapiens] {IMAGE:596063}	596063 IMAGE Consortium	AA122727
8017 B	2 0212A1FN	IMAGE EST {IMAGE:583311}	583311 IMAGE Consortium	AA162935
8018 B	4 0212A1FN	ESTs {IMAGE:583659}	583659 IMAGE Consortium	AA144095
8019 B	6 0212A1FN	ESTs, Highly similar to INTERFERON-INDUCED GUANYLATE-BINDING PROTEIN 1 [Mus musculus] {IMAGE:583808}	583808 IMAGE Consortium	AA153021
8020 B	8 0212A1FN	Mus musculus (clone HIC-53) hydrogen peroxide-inducible protein mRNA, complete cds {IMAGE:597479}	597479 IMAGE Consortium	AA144221
8021 B	10 0212A1FN	ESTs {IMAGE:597825}	597825 IMAGE Consortium	AA163820
8022 B	12 0212A1FN	IMAGE EST {IMAGE:596023}	596023 IMAGE Consortium	AA137320
8023 D	2 0212A1FN	ESTs {IMAGE:597280}	597280 IMAGE Consortium	AA138290
8024 D	4 0212A1FN	Mus musculus B lymphocyte chemoattractant BLC mRNA, complete cds {IMAGE:597433}	597433 IMAGE Consortium	AA152885
8025 D	6 0212A1FN	Bruton agammaglobulinemia tyrosine kinase	597683 IMAGE Consortium	AA154035

		{IMAGE:597683}		
8026 D	8 0212A1FN	IMAGE EST {IMAGE:583815}	583815 IMAGE Consortium	AA153026
8027 D	10 0212A1FN	ESTs {IMAGE:583845}	583845 IMAGE Consortium	AA146395
8028 D	12 0212A1FN	IMAGE EST {IMAGE:596066}	596066 IMAGE Consortium	AA137628
8029 F	2 0212A1FN	ESTs {IMAGE:599194}	599194 IMAGE Consortium	AA154932
8030 F	4 0212A1FN	B-cell translocation gene 2, anti-proliferative {IMAGE:583186}	583186 IMAGE Consortium	AA154848
8031 F	6 0212A1FN	ESTs, Highly similar to RAS-RELATED PROTEIN RAB-5A [Homo sapiens] {IMAGE:597697}	597697 IMAGE Consortium	AA153228
8032 F	8 0212A1FN	IMAGE EST {IMAGE:597445}	597445 IMAGE Consortium	AA153212
8033 F	10 0212A1FN	IMAGE EST {IMAGE:583935}	583935 IMAGE Consortium	AA154062
8034 F	12 0212A1FN	ESTs, Weakly similar to coded for by C. elegans cDNA yk13g5.3 [C.elegans] {IMAGE:583869}	583869 IMAGE Consortium	AA145820
8035 H	2 0212A1FN	ESTs {IMAGE:596927}	596927 IMAGE Consortium	AA138887
8036 H	4 0212A1FN	ESTs {IMAGE:616690}	616690 IMAGE Consortium	AA174388
8037 H	6 0212A1FN	IMAGE EST {IMAGE:583509}	583509 IMAGE Consortium	AA146379
8038 H	8 0212A1FN	ESTs, Weakly similar to C15H9.5 [C.elegans] {IMAGE:583925}	583925 IMAGE Consortium	AA154527
8039 H	10 0212A1FN	IMAGE EST {IMAGE:583689}	583689 IMAGE Consortium	AA153420
8040 H	12 0212A1FN	ESTs {IMAGE:597938}	597938 IMAGE Consortium	AA163878
8041 B	2 0219A1FO	ESTs {IMAGE:617258}	617258 IMAGE Consortium	AA174859
8042 B	4 0219A1FO	ESTs {IMAGE:619401}	619401 IMAGE Consortium	AA175583
8043 B	6 0219A1FO	ESTs {IMAGE:617360}	617360 IMAGE Consortium	AA174512
8044 B	8 0219A1FO	Mus musculus mRNA for LOK, complete cds {IMAGE:618001}	618001 IMAGE Consortium	AA175853
8045 B	10 0219A1FO	Mus musculus von Hippel-Lindau binding protein homolog (vbp-1) mRNA, complete cds {IMAGE:617975}	617975 IMAGE Consortium	AA170334
8046 B	12 0219A1FO	ESTs, Highly similar to tex292 [M.musculus] {IMAGE:620360}	620360 IMAGE Consortium	AA178163
8047 D	2 0219A1FO	IMAGE EST {IMAGE:619855}	619855 IMAGE Consortium	AA177930
8048 D	4 0219A1FO	IMAGE EST {IMAGE:618480}	618480 IMAGE Consortium	AA175279
8049 D	6 0219A1FO	Histocompatibility 2, class II antigen A, beta 1 {IMAGE:618271}	618271 IMAGE Consortium	AA175329
8050 D	8 0219A1FO	IMAGE EST {IMAGE:618474}	618474 IMAGE Consortium	AA170710
8051 D	10 0219A1FO	Cytochrome P450, 2e1, ethanol inducible {IMAGE:618870}	618870 IMAGE Consortium	AA175201
8052 D	12 0219A1FO	IMAGE EST {IMAGE:618834}	618834 IMAGE Consortium	AA170417
8053 F	2 0219A1FO	ESTs {IMAGE:618481}	618481 IMAGE Consortium	AA175283
8054 F	4 0219A1FO	ESTs {IMAGE:618375}	618375 IMAGE Consortium	AA175027
8055 F	6 0219A1FO	ESTs {IMAGE:620716}	620716 IMAGE Consortium	AA178203
8056 F	8 0219A1FO	IMAGE EST {IMAGE:620656}	620656 IMAGE Consortium	AA174531
8057 F	10 0219A1FO	ESTs, Highly similar to ERYTHROCYTE ADDUCIN BETA SUBUNIT [Homo sapiens] {IMAGE:620815}	620815 IMAGE Consortium	AA178305
8058 F	12 0219A1FO	IMAGE EST {IMAGE:620942}	620942 IMAGE Consortium	AA177241
8059 H	2 0219A1FO	Mus musculus classII MHC transactivator CIITA mRNA, complete cds {IMAGE:620236}	620236 IMAGE Consortium	AA178010

8060 H	4 0219A1FO	IMAGE EST {IMAGE:620128}	620128 IMAGE Consortium	AA172546
8061 H	6 0219A1FO	ESTs {IMAGE:620453}	620453 IMAGE Consortium	AA178166
8062 H	8 0219A1FO	ESTs {IMAGE:620659}	620659 IMAGE Consortium	AA174533
8063 H	10 0219A1FO	IMAGE EST {IMAGE:620839}	620839 IMAGE Consortium	AA178103
8064 H	12 0219A1FO	ESTs {IMAGE:620990}	620990 IMAGE Consortium	AA178253
8065 B	2 021GA1FP	IMAGE EST {IMAGE:620669}	620669 IMAGE Consortium	AA178181
8066 B	4 021GA1FP	IMAGE EST {IMAGE:620492}	620492 IMAGE Consortium	AA174367
8067 B	6 021GA1FP	ESTs {IMAGE:621040}	621040 IMAGE Consortium	AA178132
8068 B	8 021GA1FP	Small inducible cytokine A4 {IMAGE:621095}	621095 IMAGE Consortium	AA178155
8069 B	10 021GA1FP	Protein tyrosine phosphatase, non-receptor type 12 {IMAGE:619322}	619322 IMAGE Consortium	AA175740
8070 B	12 021GA1FP	IMAGE EST {IMAGE:619596}	619596 IMAGE Consortium	AA175803
8071 D	2 021GA1FP	ESTs {IMAGE:618304}	618304 IMAGE Consortium	AA170598
8072 D	4 021GA1FP	ESTs {IMAGE:618238}	618238 IMAGE Consortium	AA170574
8073 D	6 021GA1FP	IMAGE EST {IMAGE:618585}	618585 IMAGE Consortium	AA175915
8074 D	8 021GA1FP	Interferon activated gene 203 {IMAGE:618572}	618572 IMAGE Consortium	AA170251
8075 D	10 021GA1FP	IMAGE EST {IMAGE:619134}	619134 IMAGE Consortium	AA175597
8076 D	12 021GA1FP	IMAGE EST {IMAGE:616679}	616679 IMAGE Consortium	AA170476
8077 F	2 021GA1FP	IMAGE EST {IMAGE:620148}	620148 IMAGE Consortium	AA177887
8078 F	4 021GA1FP	ESTs {IMAGE:620766}	620766 IMAGE Consortium	AA178217
8079 F	6 021GA1FP	Interferon activated gene 203 {IMAGE:620541}	620541 IMAGE Consortium	AA174447
8080 F	8 021GA1FP	MEK kinase {IMAGE:621032}	621032 IMAGE Consortium	AA178632
8081 F	10 021GA1FP	IMAGE EST {IMAGE:621312}	621312 IMAGE Consortium	AA178547
8082 F	12 021GA1FP	IMAGE EST {IMAGE:618909}	618909 IMAGE Consortium	AA170529
8083 H	2 021GA1FP	ESTs {IMAGE:618137}	618137 IMAGE Consortium	AA170437
8084 H	4 021GA1FP	IMAGE EST {IMAGE:620479}	620479 IMAGE Consortium	AA178173
8085 H	6 021GA1FP	ESTs, Highly similar to MITOTIC MAD2 PROTEIN [Saccharomyces cerevisiae] {IMAGE:620733}	620733 IMAGE Consortium	AA174630
8086 H	8 021GA1FP	ESTs {IMAGE:618992}	618992 IMAGE Consortium	AA175486
8087 H	10 021GA1FP	ESTs, Weakly similar to Ydr282cp [S.cerevisiae] {IMAGE:621480}	621480 IMAGE Consortium	AA178731
8088 H	12 021GA1FP	Topoisomerase (DNA) I {IMAGE:619269}	619269 IMAGE Consortium	AA170792
8089 B	2 021NA1FQ	Interleukin 16 {IMAGE:634936}	634936 IMAGE Consortium	AA155344
8090 B	4 021NA1FQ	IMAGE EST {IMAGE:635152}	635152 IMAGE Consortium	AA162499
8091 B	6 021NA1FQ	IMAGE EST {IMAGE:635859}	635859 IMAGE Consortium	AA185328
8092 B	8 021NA1FQ	ESTs {IMAGE:622742}	622742 IMAGE Consortium	AA184254
8093 B	10 021NA1FQ	ESTs {IMAGE:621555}	621555 IMAGE Consortium	AA178763
8094 B	12 021NA1FQ	ESTs {IMAGE:621202}	621202 IMAGE Consortium	AA178487
8095 D	2 021NA1FQ	IMAGE EST {IMAGE:622693}	622693 IMAGE Consortium	AA184189
8096 D	4 021NA1FQ	ESTs {IMAGE:635507}	635507 IMAGE Consortium	AA166388
8097 D	6 021NA1FQ	IMAGE EST {IMAGE:635917}	635917 IMAGE Consortium	AA183250
8098 D	8 021NA1FQ	ESTs, Weakly similar to The KIAA0127 gene product is novel. [H.sapiens] {IMAGE:636433}	636433 IMAGE Consortium	AA183257

8099 D	10 021NA1FQ	IMAGE EST {IMAGE:636277}	636277 IMAGE Consortium	AA189315
8100 D	12 021NA1FQ	IMAGE EST {IMAGE:634110}	634110 IMAGE Consortium	AA184614
8101 F	2 021NA1FQ	ESTs, Weakly similar to myosin phosphatase target subunit 1 [H.sapiens] {IMAGE:635847}	635847 IMAGE Consortium	AA182309
8102 F	4 021NA1FQ	IMAGE EST {IMAGE:636218}	636218 IMAGE Consortium	AA183194
8103 F	6 021NA1FQ	ESTs {IMAGE:621346}	621346 IMAGE Consortium	AA178613
8104 F	8 021NA1FQ	IMAGE EST {IMAGE:621775}	621775 IMAGE Consortium	AA184105
8105 F	10 021NA1FQ	ESTs {IMAGE:634580}	634580 IMAGE Consortium	AA189624
8106 F	12 021NA1FQ	ESTs {IMAGE:634825}	634825 IMAGE Consortium	AA163483
8107 H	2 021NA1FQ	ESTs {IMAGE:634389}	634389 IMAGE Consortium	AA185231
8108 H	4 021NA1FQ	ESTs, Weakly similar to SSM4 PROTEIN [S.cerevisiae] {IMAGE:634931}	634931 IMAGE Consortium	AA164038
8109 H	6 021NA1FQ	IMAGE EST {IMAGE:634675}	634675 IMAGE Consortium	AA155387
8110 H	8 021NA1FQ	ESTs {IMAGE:635025}	635025 IMAGE Consortium	AA163246
8111 H	10 021NA1FQ	IMAGE EST {IMAGE:622724}	622724 IMAGE Consortium	AA177555
8112 H	12 021NA1FQ	IMAGE EST {IMAGE:636078}	636078 IMAGE Consortium	AA189360
8113 B	2 021UA1FR	IMAGE EST {IMAGE:620086}	620086 IMAGE Consortium	AA172493
8114 B	4 021UA1FR	ESTs {IMAGE:620675}	620675 IMAGE Consortium	AA177221
8115 B	6 021UA1FR	ESTs {IMAGE:618863}	618863 IMAGE Consortium	AA170446
8116 B	8 021UA1FR	ESTs {IMAGE:619232}	619232 IMAGE Consortium	AA175750
8117 B	10 021UA1FR	IMAGE EST {IMAGE:618910}	618910 IMAGE Consortium	AA170527
8118 B	12 021UA1FR	IMAGE EST {IMAGE:619636}	619636 IMAGE Consortium	AA175786
8119 D	2 021UA1FR	ESTs {IMAGE:618611}	618611 IMAGE Consortium	AA174941
8120 D	4 021UA1FR	Mus musculus WSB-1 mRNA, complete cds {IMAGE:619201}	619201 IMAGE Consortium	AA175692
8121 D	6 021UA1FR	ESTs, Highly similar to 60S RIBOSOMAL PROTEIN L22 [Tripneustes gratilla] {IMAGE:620947}	620947 IMAGE Consortium	AA174807
8122 D	8 021UA1FR	ESTs {IMAGE:617161}	617161 IMAGE Consortium	AA174656
8123 D	10 021UA1FR	ESTs {IMAGE:617197}	617197 IMAGE Consortium	AA174776
8124 D	12 021UA1FR	IMAGE EST {IMAGE:619385}	619385 IMAGE Consortium	AA172519
8125 F	2 021UA1FR	ESTs, Moderately similar to CELL SURFACE GLYCOPROTEIN MAC-1 ALPHA SUBUNIT PRECURSOR [Mus musculus] {IMAGE:620795}	620795 IMAGE Consortium	AA178276
8126 F	4 021UA1FR	ESTs {IMAGE:621218}	621218 IMAGE Consortium	AA178773
8127 F	6 021UA1FR	ESTs {IMAGE:617038}	617038 IMAGE Consortium	AA163432
8128 F	8 021UA1FR	ESTs {IMAGE:620018}	620018 IMAGE Consortium	AA178059
8129 F	10 021UA1FR	IMAGE EST {IMAGE:620022}	620022 IMAGE Consortium	AA178061
8130 F	12 021UA1FR	Small inducible cytokine A12 {IMAGE:620034}	620034 IMAGE Consortium	AA172456
8131 H	2 021UA1FR	L-SELECTIN PRECURSOR {IMAGE:621878}	621878 IMAGE Consortium	AA183698
8132 H	4 021UA1FR	ESTs {IMAGE:621883}	621883 IMAGE Consortium	AA178388
8133 H	6 021UA1FR	ESTs {IMAGE:622274}	622274 IMAGE Consortium	AA183371
8134 H	8 021UA1FR	Transcription factor 21 {IMAGE:634816}	634816 IMAGE Consortium	AA154683
8135 H	10 021UA1FR	ESTs {IMAGE:634989}	634989 IMAGE Consortium	AA155490
8136 H	12 021UA1FR	ESTs {IMAGE:635023}	635023 IMAGE Consortium	AA163245



8137 B	2 0211A1FS	IMAGE EST {IMAGE:622041}	622041 IMAGE Consortium	AA183934
8138 B	4 0211A1FS	IMAGE EST {IMAGE:622123}	622123 IMAGE Consortium	AA177600
8139 B	6 0211A1FS	Mus musculus double-stranded RNA-specific adenosine deaminase mRNA, complete cds {IMAGE:622431}	622431 IMAGE Consortium	AA184221
8140 B	8 0211A1FS	ESTs {IMAGE:636115}	636115 IMAGE Consortium	AA183081
8141 B	10 0211A1FS	ESTs {IMAGE:636375}	636375 IMAGE Consortium	AA183022
8142 B	12 0211A1FS	IMAGE EST {IMAGE:634235}	634235 IMAGE Consortium	AA182114
8143 D	2 0211A1FS	IMAGE EST {IMAGE:635027}	635027 IMAGE Consortium	AA163247
8144 D	4 0211A1FS	ESTs, Moderately similar to P21 activated kinase-3 [M.musculus] {IMAGE:635569}	635569 IMAGE Consortium	AA165742
8145 D	6 0211A1FS	ESTs {IMAGE:635607}	635607 IMAGE Consortium	AA166303
8146 D	8 0211A1FS	IMAGE EST {IMAGE:623103}	623103 IMAGE Consortium	AA186126
8147 D	10 0211A1FS	IMAGE EST {IMAGE:622822}	622822 IMAGE Consortium	AA177630
8148 D	12 0211A1FS	ESTs {IMAGE:634467}	634467 IMAGE Consortium	AA184913
8149 F	2 0211A1FS	IMAGE EST {IMAGE:635220}	635220 IMAGE Consortium	AA189711
8150 F	4 0211A1FS	IMAGE EST {IMAGE:635565}	635565 IMAGE Consortium	AA163378
8151 F	6 0211A1FS	Heat shock protein, DNAJ-like 2 {IMAGE:622931}	622931 IMAGE Consortium	AA184380
8152 F	8 0211A1FS	ESTs {IMAGE:622792}	622792 IMAGE Consortium	AA177621
8153 F	10 0211A1FS	B-cell receptor-associated protein 29 {IMAGE:622846}	622846 IMAGE Consortium	AA177687
8154 F	12 0211A1FS	IMAGE EST {IMAGE:621416}	621416 IMAGE Consortium	AA178654
8155 H	2 0211A1FS	ESTs {IMAGE:635197}	635197 IMAGE Consortium	AA178792
8156 H	4 0211A1FS	ESTs, Moderately similar to Toll-like receptor 1 [H.sapiens] {IMAGE:622702}	622702 IMAGE Consortium	AA177549
8157 H	6 0211A1FS	IMAGE EST {IMAGE:636033}	636033 IMAGE Consortium	AA183132
8158 H	8 0211A1FS	IMAGE EST {IMAGE:636275}	636275 IMAGE Consortium	AA189314
8159 H	10 0211A1FS	IMAGE EST {IMAGE:634167}	634167 IMAGE Consortium	AA182068
8160 H	12 0211A1FS	Mus musculus potentially prenylated protein tyrosine phosphatase mPRL-2 (Prl2) mRNA, complete cds {IMAGE:634539}	634539 IMAGE Consortium	AA186138
8161 B	2 0218A1FT	ESTs {IMAGE:622458}	622458 IMAGE Consortium	AA177358
8162 B	4 0218A1FT	IMAGE EST {IMAGE:635810}	635810 IMAGE Consortium	AA185313
8163 B	6 0218A1FT	IMAGE EST {IMAGE:622812}	622812 IMAGE Consortium	AA184236
8164 B	8 0218A1FT	ESTs, Weakly similar to HIGH AFFINITY IMMUNOGLOBULIN GAMMA FC RECEPTOR I PRECURSOR [M.musculus] {IMAGE:622960}	622960 IMAGE Consortium	AA177761
8165 B	10 0218A1FT	IMAGE EST {IMAGE:636185}	636185 IMAGE Consortium	AA183309
8166 B	12 0218A1FT	IMAGE EST {IMAGE:634252}	634252 IMAGE Consortium	AA182981
8167 D	2 0218A1FT	ESTs {IMAGE:635366}	635366 IMAGE Consortium	AA185984
8168 D	4 0218A1FT	IMAGE EST {IMAGE:635650}	635650 IMAGE Consortium	AA166543
8169 D	6 0218A1FT	ESTs {IMAGE:635882}	635882 IMAGE Consortium	AA182357
8170 D	8 0218A1FT	ESTs {IMAGE:636337}	636337 IMAGE Consortium	AA189332
8171 D	10 0218A1FT	T-box 6 {IMAGE:636895}	636895 IMAGE Consortium	AA182995
8172 D	12 0218A1FT	IMAGE EST {IMAGE:636837}	636837 IMAGE Consortium	AA183078

8173 F	2 0218A1FT	Mouse spi2 proteinase inhibitor (spi2/eb1) mRNA, 3' end {IMAGE:640951}	640951 IMAGE Consortium	AA210495
8174 F	4 0218A1FT	IMAGE EST {IMAGE:638829}	638829 IMAGE Consortium	AA198969
8175 F	6 0218A1FT	IMAGE EST {IMAGE:639212}	639212 IMAGE Consortium	AA200984
8176 F	8 0218A1FT	ESTs, Highly similar to RNA splicing-related protein [R.norvegicus] {IMAGE:636766}	636766 IMAGE Consortium	AA183061
8177 F	10 0218A1FT	ESTs {IMAGE:639990}	639990 IMAGE Consortium	AA197474
8178 F	12 0218A1FT	Interleukin 4 {IMAGE:640300}	640300 IMAGE Consortium	AA199406
8179 H	2 0218A1FT	IMAGE EST {IMAGE:639431}	639431 IMAGE Consortium	AA200695
8180 H	4 0218A1FT	ESTs {IMAGE:637291}	637291 IMAGE Consortium	AA185873
8181 H	6 0218A1FT	ESTs {IMAGE:639863}	639863 IMAGE Consortium	AA197405
8182 H	8 0218A1FT	ESTs {IMAGE:638311}	638311 IMAGE Consortium	AA200775
8183 H	10 0218A1FT	ESTs {IMAGE:640579}	640579 IMAGE Consortium	AA204257
8184 H	12 0218A1FT	IMAGE EST {IMAGE:640952}	640952 IMAGE Consortium	AA210474
8185 B	2 021FA1FU	IMAGE EST {IMAGE:638823}	638823 IMAGE Consortium	AA200071
8186 B	4 021FA1FU	IMAGE EST {IMAGE:639147}	639147 IMAGE Consortium	AA201092
8187 B	6 021FA1FU	IMAGE EST {IMAGE:639206}	639206 IMAGE Consortium	AA200606
8188 B	8 021FA1FU	IMAGE EST {IMAGE:636827}	636827 IMAGE Consortium	AA183302
8189 B	10 021FA1FU	Caspase 2 {IMAGE:639403}	639403 IMAGE Consortium	AA200808
8190 B	12 021FA1FU	ESTs {IMAGE:639994}	639994 IMAGE Consortium	AA197601
8191 D	2 021FA1FU	ESTs {IMAGE:639451}	639451 IMAGE Consortium	AA200850
8192 D	4 021FA1FU	ESTs {IMAGE:639707}	639707 IMAGE Consortium	AA197398
8193 D	6 021FA1FU	IMAGE EST {IMAGE:639666}	639666 IMAGE Consortium	AA197468
8194 D	8 021FA1FU	IMAGE EST {IMAGE:640106}	640106 IMAGE Consortium	AA197507
8195 D	10 021FA1FU	Palmitoyl-protein thioesterase {IMAGE:637934}	637934 IMAGE Consortium	AA189995
8196 D	12 021FA1FU	IMAGE EST {IMAGE:640757}	640757 IMAGE Consortium	AA203889
8197 F	2 021FA1FU	IMAGE EST {IMAGE:639971}	639971 IMAGE Consortium	AA197597
8198 F	4 021FA1FU	ESTs {IMAGE:639834}	639834 IMAGE Consortium	AA197401
8199 F	6 021FA1FU	ESTs, Moderately similar to cDNA EST EMBL:D74283 comes from this gene [C.elegans] {IMAGE:637463}	637463 IMAGE Consortium	AA185058
8200 F	8 021FA1FU	ESTs {IMAGE:637815}	637815 IMAGE Consortium	AA189751
8201 F	10 021FA1FU	IMAGE EST {IMAGE:638453}	638453 IMAGE Consortium	AA216990
8202 F	12 021FA1FU	ESTs, Moderately similar to The KIAA0191 gene is expressed ubiquitously. [H.sapiens] {IMAGE:638204}	638204 IMAGE Consortium	AA200734
8203 H	2 021FA1FU	Mus musculus non-receptor protein tyrosine kinase Ack mRNA, complete cds {IMAGE:640099}	640099 IMAGE Consortium	AA197459
8204 H	4 021FA1FU	ESTs {IMAGE:640710}	640710 IMAGE Consortium	AA203853
8205 H	6 021FA1FU	ESTs, Highly similar to transcription factor NF-AT 45K chain [H.sapiens] {IMAGE:638252}	638252 IMAGE Consortium	AA200737
8206 H	8 021FA1FU	IMAGE EST {IMAGE:638745}	638745 IMAGE Consortium	AA200945
8207 H	10 021FA1FU	ESTs {IMAGE:638805}	638805 IMAGE Consortium	AA198961
8208 H	12 021FA1FU	IMAGE EST {IMAGE:641910}	641910 IMAGE Consortium	AA210215
8209 B	2 021MA1FV	ESTs {IMAGE:639911}	639911 IMAGE Consortium	AA197440

8210 B	4	021MA1FV	IMAGE EST {IMAGE:640085}	640085 IMAGE Consortium	AA198542
8211 B	6	021MA1FV	Fragile X mental retardation syndrome 1 homolog {IMAGE:638257}	638257 IMAGE Consortium	AA200033
8212 B	8	021MA1FV	Hemopoietic cell kinase {IMAGE:638455}	638455 IMAGE Consortium	AA216991
8213 B	10	021MA1FV	ESTs, Highly similar to DESMOPLAKIN I AND II [Homo sapiens] {IMAGE:640556}	640556 IMAGE Consortium	AA204246
8214 B	12	021MA1FV	ESTs {IMAGE:640919}	640919 IMAGE Consortium	AA204360
8215 D	2	021MA1FV	IMAGE EST {IMAGE:639902}	639902 IMAGE Consortium	AA198514
8216 D	4	021MA1FV	IMAGE EST {IMAGE:637903}	637903 IMAGE Consortium	AA189551
8217 D	6	021MA1FV	ESTs, Highly similar to A-RAF PROTO-ONCOGENE SERINE/THREONINE-PROTEIN KINASE [Rattus norvegicus] {IMAGE:640705}	640705 IMAGE Consortium	AA203871
8218 D	8	021MA1FV	ESTs {IMAGE:638416}	638416 IMAGE Consortium	AA200289
8219 D	10	021MA1FV	Cell division cycle 25A {IMAGE:641051}	641051 IMAGE Consortium	AA204385
8220 D	12	021MA1FV	IMAGE EST {IMAGE:641462}	641462 IMAGE Consortium	AA200457
8221 F	2	021MA1FV	IMAGE EST {IMAGE:637808}	637808 IMAGE Consortium	AA189641
8222 F	4	021MA1FV	IMAGE EST {IMAGE:638234}	638234 IMAGE Consortium	AA200025
8223 F	6	021MA1FV	ESTs {IMAGE:638217}	638217 IMAGE Consortium	AA200330
8224 F	8	021MA1FV	ESTs {IMAGE:638522}	638522 IMAGE Consortium	AA200091
8225 F	10	021MA1FV	IMAGE EST {IMAGE:641234}	641234 IMAGE Consortium	AA200358
8226 F	12	021MA1FV	ESTs {IMAGE:641906}	641906 IMAGE Consortium	AA210213
8227 H	2	021MA1FV	ESTs, Weakly similar to C50F4.12 [C.elegans] {IMAGE:638178}	638178 IMAGE Consortium	AA200003
8228 H	4	021MA1FV	IMAGE EST {IMAGE:641080}	641080 IMAGE Consortium	AA204459
8229 H	6	021MA1FV	ESTs, Highly similar to GTP-binding protein-associated protein [M.musculus] {IMAGE:641451}	641451 IMAGE Consortium	AA200361
8230 H	8	021MA1FV	ESTs {IMAGE:641368}	641368 IMAGE Consortium	AA200482
8231 H	10	021MA1FV	IMAGE EST {IMAGE:641668}	641668 IMAGE Consortium	AA209919
8232 H	12	021MA1FV	ESTs {IMAGE:639379}	639379 IMAGE Consortium	AA200801
8233 B	2	021TA1FW	Mus musculus putative purine nucleotide binding protein mRNA, complete cds {IMAGE:653367}	653367 IMAGE Consortium	AA240404
8234 B	4	021TA1FW	Mouse mRNA for NDPP-1 protein, complete cds {IMAGE:653622}	653622 IMAGE Consortium	AA217593
8235 B	6	021TA1FW	ESTs {IMAGE:653589}	653589 IMAGE Consortium	AA217806
8236 B	8	021TA1FW	ESTs {IMAGE:643257}	643257 IMAGE Consortium	AA186075
8237 B	10	021TA1FW	ESTs {IMAGE:643810}	643810 IMAGE Consortium	AA203910
8238 B	12	021TA1FW	IMAGE EST {IMAGE:641911}	641911 IMAGE Consortium	AA210220
8239 D	2	021TA1FW	ESTs {IMAGE:653368}	653368 IMAGE Consortium	AA240506
8240 D	4	021TA1FW	Mus musculus adaptor protein complex-3 sigma3A subunit isoform mRNA, complete cds {IMAGE:653600}	653600 IMAGE Consortium	AA217687
8241 D	6	021TA1FW	Kinesin heavy chain member 2 {IMAGE:657917}	657917 IMAGE Consortium	AA199274
8242 D	8	021TA1FW	Zinc transporter 4 {IMAGE:643619}	643619 IMAGE Consortium	AA203866
8243 D	10	021TA1FW	ESTs, Highly similar to ALPHA ENOLASE [Mus musculus]	643854 IMAGE Consortium	AA204262

		{IMAGE:643854}		
8244 D	12 021TA1FW	Apolipoprotein A-I {IMAGE:641720}	641720 IMAGE Consortium	AA209970
8245 F	2 021TA1FW	ESTs {IMAGE:643075}	643075 IMAGE Consortium	AA189255
8246 F	4 021TA1FW	ESTs {IMAGE:643298}	643298 IMAGE Consortium	AA189595
8247 F	6 021TA1FW	ESTs {IMAGE:643644}	643644 IMAGE Consortium	AA204160
8248 F	8 021TA1FW	ESTs {IMAGE:643902}	643902 IMAGE Consortium	AA204090
8249 F	10 021TA1FW	IMAGE EST {IMAGE:644864}	644864 IMAGE Consortium	AA209551
8250 F	12 021TA1FW	ESTs {IMAGE:641687}	641687 IMAGE Consortium	AA209990
8251 H	2 021TA1FW	IMAGE EST {IMAGE:657918}	657918 IMAGE Consortium	AA199272
8252 H	4 021TA1FW	ESTs {IMAGE:657766}	657766 IMAGE Consortium	AA199188
8253 H	6 021TA1FW	ESTs {IMAGE:643905}	643905 IMAGE Consortium	AA203809
8254 H	8 021TA1FW	ESTs {IMAGE:641918}	641918 IMAGE Consortium	AA209904
8255 H	10 021TA1FW	ESTs {IMAGE:652666}	652666 IMAGE Consortium	AA216849
8256 H	12 021TA1FW	IMAGE EST {IMAGE:652904}	652904 IMAGE Consortium	AA217081
8257 B	2 0210A1FX	Pore forming protein {IMAGE:641679}	641679 IMAGE Consortium	AA209962
8258 B	4 0210A1FX	IMAGE EST {IMAGE:641660}	641660 IMAGE Consortium	AA209977
8259 B	6 0210A1FX	ESTs {IMAGE:642120}	642120 IMAGE Consortium	AA212128
8260 B	8 0210A1FX	Hemochromatosis {IMAGE:652689}	652689 IMAGE Consortium	AA217236
8261 B	10 0210A1FX	ESTs {IMAGE:652744}	652744 IMAGE Consortium	AA217352
8262 B	12 0210A1FX	Mus musculus formin binding protein 21 mRNA, complete cds {IMAGE:653252}	653252 IMAGE Consortium	AA217435
8263 D	2 0210A1FX	IMAGE EST {IMAGE:642168}	642168 IMAGE Consortium	AA212168
8264 D	4 0210A1FX	Mouse gene for mu chain association protein (8hs20) {IMAGE:652535}	652535 IMAGE Consortium	AA217465
8265 D	6 0210A1FX	IMAGE EST {IMAGE:652698}	652698 IMAGE Consortium	AA217366
8266 D	8 0210A1FX	ESTs {IMAGE:653244}	653244 IMAGE Consortium	AA217322
8267 D	10 0210A1FX	ESTs, Weakly similar to QUINONE OXIDOREDUCTASE [M.musculus] {IMAGE:653324}	653324 IMAGE Consortium	AA241313
8268 D	12 0210A1FX	IMAGE EST {IMAGE:642741}	642741 IMAGE Consortium	AA185567
8269 F	2 0210A1FX	ESTs {IMAGE:652559}	652559 IMAGE Consortium	AA217491
8270 F	4 0210A1FX	ESTs {IMAGE:653125}	653125 IMAGE Consortium	AA216878
8271 F	6 0210A1FX	Son of sevenless homolog 2, (Drosophila) {IMAGE:642732}	642732 IMAGE Consortium	AA184845
8272 F	8 0210A1FX	IMAGE EST {IMAGE:643326}	643326 IMAGE Consortium	AA189603
8273 F	10 0210A1FX	ESTs {IMAGE:643431}	643431 IMAGE Consortium	AA186173
8274 F	12 0210A1FX	ESTs {IMAGE:643673}	643673 IMAGE Consortium	AA204593
8275 H	2 0210A1FX	ESTs, Highly similar to HYPOTHETICAL TRP-ASP REPEATS CONTAINING PROTEIN IN PGI1-KTR4 INTERGENIC REGION [Saccharomyces cerevisiae] {IMAGE:657961}	657961 IMAGE Consortium	AA198335
8276 H	4 0210A1FX	IMAGE EST {IMAGE:644356}	644356 IMAGE Consortium	AA265225
8277 H	6 0210A1FX	ESTs, Weakly similar to C05D9.6 gene product [C.elegans] {IMAGE:641752}	641752 IMAGE Consortium	AA210034
8278 H	8 0210A1FX	ESTs {IMAGE:641685}	641685 IMAGE Consortium	AA209989

8279 H	10 0210A1FX	ESTs, Highly similar to PUTATIVE UBIQUITIN CARBOXYL-TERMINAL HYDROLASE C13A11.04C [Schizosaccharomyces pombe] {IMAGE:642148}	642148 IMAGE Consortium	AA212162
8280 H	12 0210A1FX	IMAGE EST {IMAGE:653016}	653016 IMAGE Consortium	AA217294
8281 B	2 0217A1FY	IMAGE EST {IMAGE:850960}	850960 IMAGE Consortium	AA462281
8282 B	4 0217A1FY	ESTs {IMAGE:851649}	851649 IMAGE Consortium	AA500751
8283 B	6 0217A1FY	Guanine nucleotide binding protein, alpha 11 {IMAGE:850418}	850418 IMAGE Consortium	AA451269
8284 B	8 0217A1FY	ESTs {IMAGE:850821}	850821 IMAGE Consortium	AA451125
8285 B	10 0217A1FY	IMAGE EST {IMAGE:851638}	851638 IMAGE Consortium	AA500724
8286 B	12 0217A1FY	ESTs, Highly similar to EF-7 [M.musculus] {IMAGE:850717}	850717 IMAGE Consortium	AA451058
8287 D	2 0217A1FY	IMAGE EST {IMAGE:850395}	850395 IMAGE Consortium	AA451259
8288 D	4 0217A1FY	IMAGE EST {IMAGE:850976}	850976 IMAGE Consortium	AA462387
8289 D	6 0217A1FY	Mus musculus ras-related protein (rab18) mRNA, complete cds {IMAGE:851551}	851551 IMAGE Consortium	AA461854
8290 D	8 0217A1FY	Mus musculus Xin mRNA, complete cds {IMAGE:860360}	860360 IMAGE Consortium	AA501052
8291 D	10 0217A1FY	Mus musculus putative histone deacetylase (HD1) mRNA, partial cds {IMAGE:850354}	850354 IMAGE Consortium	AA451208
8292 D	12 0217A1FY	Pyruvate decarboxylase {IMAGE:850104}	850104 IMAGE Consortium	AA450665
8293 F	2 0217A1FY	ESTs {IMAGE:860181}	860181 IMAGE Consortium	AA500861
8294 F	4 0217A1FY	ESTs {IMAGE:850389}	850389 IMAGE Consortium	AA451244
8295 F	6 0217A1FY	ESTs, Highly similar to CLATHRIN COAT ASSEMBLY PROTEIN AP47 HOMOLOG 1 [R.norvegicus] {IMAGE:851573}	851573 IMAGE Consortium	AA461912
8296 F	8 0217A1FY	Kras oncogene-associated gene {IMAGE:860087}	860087 IMAGE Consortium	AA500799
8297 F	10 0217A1FY	Mus musculus mRNA for DEDD protein {IMAGE:874041}	874041 IMAGE Consortium	AA471746
8298 F	12 0217A1FY	ESTs, Weakly similar to F35G2.2 [C.elegans] {IMAGE:864753}	864753 IMAGE Consortium	AA444621
8299 H	2 0217A1FY	IMAGE EST {IMAGE:874154}	874154 IMAGE Consortium	AA472271
8300 H	4 0217A1FY	ESTs, Highly similar to hNop56 [H.sapiens] {IMAGE:874146}	874146 IMAGE Consortium	AA471760
8301 H	6 0217A1FY	IMAGE EST {IMAGE:874293}	874293 IMAGE Consortium	AA472536
8302 H	8 0217A1FY	ESTs {IMAGE:871732}	871732 IMAGE Consortium	AA462499
8303 H	10 0217A1FY	ESTs, Highly similar to ENDOTHELIAL ACTIN-BINDING PROTEIN [Homo sapiens] {IMAGE:873058}	873058 IMAGE Consortium	AA466198
8304 H	12 0217A1FY	IMAGE EST {IMAGE:873590}	873590 IMAGE Consortium	AA472764
8305 B	2 021EA1FZ	IMAGE EST {IMAGE:643844}	643844 IMAGE Consortium	AA204206
8306 B	4 021EA1FZ	ESTs {IMAGE:644358}	644358 IMAGE Consortium	AA265226
8307 B	6 021EA1FZ	ESTs, Highly similar to similar to human ZFY protein. [H.sapiens] {IMAGE:641892}	641892 IMAGE Consortium	AA209887
8308 B	8 021EA1FZ	ESTs {IMAGE:641800}	641800 IMAGE Consortium	AA210167
8309 B	10 021EA1FZ	IMAGE EST {IMAGE:641733}	641733 IMAGE Consortium	AA210013
8310 B	12 021EA1FZ	IMAGE EST {IMAGE:642294}	642294 IMAGE Consortium	AA212408
8311 D	2 021EA1FZ	ESTs {IMAGE:658663}	658663 IMAGE Consortium	AA230850

8312 D	4 021EA1FZ	Mus musculus metalloprotease/disintegrin/cysteine rich protein precursor (MDC9) cDNA, complete cds {IMAGE:644907}	644907 IMAGE Consortium	AA210306
8313 D	6 021EA1FZ	IMAGE EST {IMAGE:644993}	644993 IMAGE Consortium	AA210406
8314 D	8 021EA1FZ	ESTs {IMAGE:642275}	642275 IMAGE Consortium	AA212551
8315 D	10 021EA1FZ	ESTs {IMAGE:642066}	642066 IMAGE Consortium	AA210182
8316 D	12 021EA1FZ	ESTs {IMAGE:642482}	642482 IMAGE Consortium	AA184855
8317 F	2 021EA1FZ	ESTs {IMAGE:847226}	847226 IMAGE Consortium	AA420104
8318 F	4 021EA1FZ	Mus musculus EBI-1 ligand chemokine mRNA, complete cds {IMAGE:832043}	832043 IMAGE Consortium	AA444730
8319 F	6 021EA1FZ	ESTs {IMAGE:832393}	832393 IMAGE Consortium	AA427149
8320 F	8 021EA1FZ	IMAGE EST {IMAGE:846847}	846847 IMAGE Consortium	AA422494
8321 F	10 021EA1FZ	IMAGE EST {IMAGE:832774}	832774 IMAGE Consortium	AA427119
8322 F	12 021EA1FZ	IMAGE EST {IMAGE:846525}	846525 IMAGE Consortium	AA433547
8323 H	2 021EA1FZ	IMAGE EST {IMAGE:846131}	846131 IMAGE Consortium	AA415942
8324 H	4 021EA1FZ	IMAGE EST {IMAGE:848403}	848403 IMAGE Consortium	AA445678
8325 H	6 021EA1FZ	ESTs, Highly similar to NADH-UBIQUINONE OXIDOREDUCTASE B22 SUBUNIT [Bos taurus] {IMAGE:846155}	846155 IMAGE Consortium	AA415725
8326 H	8 021EA1FZ	ESTs, Highly similar to G10 PROTEIN [Xenopus laevis] {IMAGE:832007}	832007 IMAGE Consortium	AA444533
8327 H	10 021EA1FZ	IMAGE EST {IMAGE:832589}	832589 IMAGE Consortium	AA423708
8328 H	12 021EA1FZ	ESTs {IMAGE:847202}	847202 IMAGE Consortium	AA420054
8329 B	2 021EA1G0	ESTs {IMAGE:874003}	874003 IMAGE Consortium	AA472186
8330 B	4 021EA1G0	IMAGE EST {IMAGE:870808}	870808 IMAGE Consortium	AA461990
8331 B	6 021EA1G0	Mus musculus calcium-binding protein S100A1 mRNA, complete cds {IMAGE:872869}	872869 IMAGE Consortium	AA466432
8332 B	8 021EA1G0	Mus musculus mRNA for TOM1 protein {IMAGE:873940}	873940 IMAGE Consortium	AA475177
8333 B	10 021EA1G0	ESTs {IMAGE:874588}	874588 IMAGE Consortium	AA472424
8334 B	12 021EA1G0	IMAGE EST {IMAGE:874705}	874705 IMAGE Consortium	AA472383
8335 D	2 021EA1G0	ESTs, Highly similar to RNA polymerase transcriptional regulation mediator [H.sapiens] {IMAGE:871272}	871272 IMAGE Consortium	AA462971
8336 D	4 021EA1G0	IMAGE EST {IMAGE:871674}	871674 IMAGE Consortium	AA462347
8337 D	6 021EA1G0	Pancreatitis-associated protein {IMAGE:873502}	873502 IMAGE Consortium	AA473899
8338 D	8 021EA1G0	IMAGE EST {IMAGE:874030}	874030 IMAGE Consortium	AA472200
8339 D	10 021EA1G0	CD36 antigen {IMAGE:864409}	864409 IMAGE Consortium	AA458178
8340 D	12 021EA1G0	ESTs, Moderately similar to Similarity to S.pombe hypothetical protein C2F7.02C [C.elegans] {IMAGE:874232}	874232 IMAGE Consortium	AA472410
8341 F	2 021EA1G0	ESTs {IMAGE:874847}	874847 IMAGE Consortium	AA474904
8342 F	4 021EA1G0	Natural resistance-associated macrophage protein 2 {IMAGE:872793}	872793 IMAGE Consortium	AA463173
8343 F	6 021EA1G0	Cystatin B {IMAGE:873417}	873417 IMAGE Consortium	AA473768
8344 F	8 021EA1G0	IMAGE EST {IMAGE:890422}	890422 IMAGE Consortium	AA509548

8345 F	10 021EA1G0	ESTs {IMAGE:891052}	891052 IMAGE Consortium	AA510110
8346 F	12 021EA1G0	IMAGE EST {IMAGE:890999}	890999 IMAGE Consortium	AA511242
8347 H	2 021EA1G0	Mus musculus CRP-ductin-alpha mRNA, complete cds {IMAGE:888707}	888707 IMAGE Consortium	AA498773
8348 H	4 021EA1G0	Myeloid cell leukemia sequence 1 {IMAGE:889253}	889253 IMAGE Consortium	AA498218
8349 H	6 021EA1G0	ESTs {IMAGE:890716}	890716 IMAGE Consortium	AA510008
8350 H	8 021EA1G0	ESTs {IMAGE:890726}	890726 IMAGE Consortium	AA510013
8351 H	10 021EA1G0	Mus musculus hematopoietic lineage switch 2 (HLS2) mRNA, complete cds {IMAGE:876463}	876463 IMAGE Consortium	AA475930
8352 H	12 021EA1G0	ESTs, Weakly similar to MATRIN 3 [R.norvegicus] {IMAGE:876381}	876381 IMAGE Consortium	AA475805
8353 B	2 021LA1G1	ESTs {IMAGE:876876}	876876 IMAGE Consortium	AA497484
8354 B	4 021LA1G1	IMAGE EST {IMAGE:902096}	902096 IMAGE Consortium	AA516559
8355 B	6 021LA1G1	IMAGE EST {IMAGE:889437}	889437 IMAGE Consortium	AA498455
8356 B	8 021LA1G1	IMAGE EST {IMAGE:890607}	890607 IMAGE Consortium	AA509765
8357 B	10 021LA1G1	IMAGE EST {IMAGE:876166}	876166 IMAGE Consortium	AA476157
8358 B	12 021LA1G1	IMAGE EST {IMAGE:876655}	876655 IMAGE Consortium	AA499263
8359 D	2 021LA1G1	IMAGE EST {IMAGE:876646}	876646 IMAGE Consortium	AA476079
8360 D	4 021LA1G1	ESTs, Weakly similar to C32E8.3 [C.elegans] {IMAGE:888544}	888544 IMAGE Consortium	AA498788
8361 D	6 021LA1G1	Heat shock protein, 74 kDa {IMAGE:889543}	889543 IMAGE Consortium	AA498713
8362 D	8 021LA1G1	ESTs {IMAGE:890707}	890707 IMAGE Consortium	AA510004
8363 D	10 021LA1G1	Mus musculus transcription factor GIF mRNA, complete cds {IMAGE:876145}	876145 IMAGE Consortium	AA476158
8364 D	12 021LA1G1	ESTs, Weakly similar to coded for by C. elegans cDNA CEESD64F [C.elegans] {IMAGE:876361}	876361 IMAGE Consortium	AA475795
8365 F	2 021LA1G1	Guanylate cyclase activator 2 (guanylin 2, intestinal, heatstable) {IMAGE:889440}	889440 IMAGE Consortium	AA498457
8366 F	4 021LA1G1	BRAIN PROTEIN H5 {IMAGE:889281}	889281 IMAGE Consortium	AA498275
8367 F	6 021LA1G1	Tissue inhibitor of metalloproteinase 2 {IMAGE:902923}	902923 IMAGE Consortium	AA518165
8368 F	8 021LA1G1	Mus musculus GTP binding protein (GTP2) mRNA, complete cds {IMAGE:890444}	890444 IMAGE Consortium	AA509565
8369 F	10 021LA1G1	Selectin, platelet (p-selectin) ligand {IMAGE:876033}	876033 IMAGE Consortium	AA475311
8370 F	12 021LA1G1	RAS-RELATED C3 BOTULINUM TOXIN SUBSTRATE 2 {IMAGE:891108}	891108 IMAGE Consortium	AA510140
8371 H	2 021LA1G1	ESTs {IMAGE:891260}	891260 IMAGE Consortium	AA510873
8372 H	4 021LA1G1	IMAGE EST {IMAGE:888624}	888624 IMAGE Consortium	AA498801
8373 H	6 021LA1G1	ESTs, Highly similar to S-ADENOSYLMETHIONINE SYNTHETASE GAMMA FORM [Homo sapiens] {IMAGE:889659}	889659 IMAGE Consortium	AA498312
8374 H	8 021LA1G1	ESTs {IMAGE:876106}	876106 IMAGE Consortium	AA475528
8375 H	10 021LA1G1	ESTs {IMAGE:891110}	891110 IMAGE Consortium	AA510141
8376 H	12 021LA1G1	ESTs, Moderately similar to homologue of Drosophila Fat protein [H.sapiens] {IMAGE:888553}	888553 IMAGE Consortium	AA498605

8377 B	2 021SA1G2	IMAGE EST {IMAGE:949480}	949480 IMAGE Consortium	AA537676
8378 B	4 021SA1G2	IMAGE EST {IMAGE:935377}	935377 IMAGE Consortium	AA543277
8379 B	6 021SA1G2	Period homolog (Drosophila) {IMAGE:949546}	949546 IMAGE Consortium	AA537133
8380 B	8 021SA1G2	Mus domesticus nuclear binding factor NF2d9 mRNA, complete cds {IMAGE:949534}	949534 IMAGE Consortium	AA537117
8381 B	10 021SA1G2	ESTs, Weakly similar to PROTEIN PHOSPHATASES PP1 REGULATORY SUBUNIT SDS22 [Schizosaccharomyces pombe] {IMAGE:948648}	948648 IMAGE Consortium	AA543106
8382 B	12 021SA1G2	ESTs, Weakly similar to SERINE/THREONINE-PROTEIN KINASE PCTAIRE-3 [M.musculus] {IMAGE:920069}	920069 IMAGE Consortium	AA499941
8383 D	2 021SA1G2	ESTs {IMAGE:949003}	949003 IMAGE Consortium	AA543796
8384 D	4 021SA1G2	IMAGE EST {IMAGE:949266}	949266 IMAGE Consortium	AA544916
8385 D	6 021SA1G2	Mus musculus gob-5 mRNA, complete cds {IMAGE:904618}	904618 IMAGE Consortium	AA518112
8386 D	8 021SA1G2	ESTs {IMAGE:948588}	948588 IMAGE Consortium	AA544722
8387 D	10 021SA1G2	ESTs {IMAGE:920414}	920414 IMAGE Consortium	AA538456
8388 D	12 021SA1G2	ESTs, Highly similar to CLATHRIN COAT ASSEMBLY PROTEIN AP47 HOMOLOG 1 [R.norvegicus] {IMAGE:935378}	935378 IMAGE Consortium	AA543269
8389 F	2 021SA1G2	Mouse alpha-B crystallin mRNA {IMAGE:672201}	672201 IMAGE Consortium	AA231358
8390 F	4 021SA1G2	TRANSPLANTATION ANTIGEN P35B {IMAGE:660586}	660586 IMAGE Consortium	AA222875
8391 F	6 021SA1G2	ESTs {IMAGE:676153}	676153 IMAGE Consortium	AA209002
8392 F	8 021SA1G2	IMAGE EST {IMAGE:669869}	669869 IMAGE Consortium	AA250443
8393 F	10 021SA1G2	Histocompatibility 2, class II, locus Ma {IMAGE:669933}	669933 IMAGE Consortium	AA230649
8394 F	12 021SA1G2	IMAGE EST {IMAGE:677555}	677555 IMAGE Consortium	AA213002
8395 H	2 021SA1G2	Neurofilament, light polypeptide {IMAGE:660991}	660991 IMAGE Consortium	AA253725
8396 H	4 021SA1G2	Pore forming protein {IMAGE:676539}	676539 IMAGE Consortium	AA208865
8397 H	6 021SA1G2	Degenerative spermatocyte homolog (Drosophila) {IMAGE:676578}	676578 IMAGE Consortium	AA209017
8398 H	8 021SA1G2	IMAGE EST {IMAGE:677091}	677091 IMAGE Consortium	AA212707
8399 H	10 021SA1G2	ESTs, Moderately similar to glutaminase [R.norvegicus] {IMAGE:677076}	677076 IMAGE Consortium	AA212816
8400 H	12 021SA1G2	Mus musculus Nip2l mRNA, complete cds {IMAGE:677412}	677412 IMAGE Consortium	AA212780
8401 B	2 021ZA1G3	Axin {IMAGE:922015}	922015 IMAGE Consortium	AA512276
8402 B	4 021ZA1G3	IMAGE EST {IMAGE:949246}	949246 IMAGE Consortium	AA544895
8403 B	6 021ZA1G3	Mus musculus groucho-related gene 4 protein (Grg4) mRNA, partial cds {IMAGE:949722}	949722 IMAGE Consortium	AA537430
8404 B	8 021ZA1G3	IMAGE EST {IMAGE:535763}	535763 IMAGE Consortium	AA108185
8405 B	10 021ZA1G3	IMAGE EST {IMAGE:948689}	948689 IMAGE Consortium	AA543145
8406 B	12 021ZA1G3	IMAGE EST {IMAGE:920335}	920335 IMAGE Consortium	AA538376
8407 D	2 021ZA1G3	Heme oxygenase (decycling) 2 {IMAGE:567485}	567485 IMAGE Consortium	AA183852
8408 D	4 021ZA1G3	IMAGE EST {IMAGE:948685}	948685 IMAGE Consortium	AA543143
8409 D	6 021ZA1G3	ESTs {IMAGE:948909}	948909 IMAGE Consortium	AA544963
8410 D	8 021ZA1G3	IMAGE EST {IMAGE:949512}	949512 IMAGE Consortium	AA537107



8411 D	10 021ZA1G3	ESTs {IMAGE:935570}	935570 IMAGE Consortium	AA543573
8412 D	12 021ZA1G3	IMAGE EST {IMAGE:949644}	949644 IMAGE Consortium	AA537291
8413 F	2 021ZA1G3	Mus musculus Balb/c zinc finger protein PZF (Pzf) mRNA, complete cds {IMAGE:947593}	947593 IMAGE Consortium	AA543497
8414 F	4 021ZA1G3	IMAGE EST {IMAGE:948662}	948662 IMAGE Consortium	AA543113
8415 F	6 021ZA1G3	Histocompatibility 2, D region locus 1 {IMAGE:920425}	920425 IMAGE Consortium	AA538511
8416 F	8 021ZA1G3	ESTs {IMAGE:948871}	948871 IMAGE Consortium	AA544881
8417 F	10 021ZA1G3	IMAGE EST {IMAGE:949078}	949078 IMAGE Consortium	AA543925
8418 F	12 021ZA1G3	IMAGE EST {IMAGE:949656}	949656 IMAGE Consortium	AA537342
8419 H	2 021ZA1G3	ESTs {IMAGE:949617}	949617 IMAGE Consortium	AA537277
8420 H	4 021ZA1G3	IMAGE EST {IMAGE:947786}	947786 IMAGE Consortium	AA542081
8421 H	6 021ZA1G3	IMAGE EST {IMAGE:905403}	905403 IMAGE Consortium	AA529046
8422 H	8 021ZA1G3	ESTs {IMAGE:948381}	948381 IMAGE Consortium	AA544354
8423 H	10 021ZA1G3	Interferon gamma receptor {IMAGE:920516}	920516 IMAGE Consortium	AA541842
8424 H	12 021ZA1G3	M.musculus mRNA for Mpgc60 protein {IMAGE:922096}	922096 IMAGE Consortium	AA512376
8425 B	2 0216A1G4	ESTs, Weakly similar to ORF YPL149w [S.cerevisiae] {IMAGE:660954}	660954 IMAGE Consortium	AA230682
8426 B	4 0216A1G4	ESTs {IMAGE:670080}	670080 IMAGE Consortium	AA221300
8427 B	6 0216A1G4	ESTs, Moderately similar to HYPOTHETICAL 167.8 KD PROTEIN CCE1-CAP1 INTERGENIC REGION [Saccharomyces cerevisiae] {IMAGE:669999}	669999 IMAGE Consortium	AA253938
8428 B	8 0216A1G4	ESTs, Highly similar to PROBABLE 3-OXOADIPATE COA-TRANSFERASE SUBUNIT B [Bacillus subtilis] {IMAGE:670658}	670658 IMAGE Consortium	AA230896
8429 B	10 0216A1G4	ESTs {IMAGE:677592}	677592 IMAGE Consortium	AA212893
8430 B	12 0216A1G4	ESTs, Highly similar to GMP SYNTHASE [H.sapiens] {IMAGE:658714}	658714 IMAGE Consortium	AA221877
8431 D	2 0216A1G4	ESTs, Moderately similar to similar to human DNA-binding protein 5. [H.sapiens] {IMAGE:670557}	670557 IMAGE Consortium	AA221914
8432 D	4 0216A1G4	Midkine {IMAGE:671218}	671218 IMAGE Consortium	AA222497
8433 D	6 0216A1G4	ESTs {IMAGE:671845}	671845 IMAGE Consortium	AA242310
8434 D	8 0216A1G4	ESTs {IMAGE:671663}	671663 IMAGE Consortium	AA241785
8435 D	10 0216A1G4	ESTs {IMAGE:671940}	671940 IMAGE Consortium	AA242702
8436 D	12 0216A1G4	ESTs {IMAGE:672351}	672351 IMAGE Consortium	AA231566
8437 F	2 0216A1G4	ESTs, Highly similar to ELONGATION FACTOR 1-GAMMA [Homo sapiens] {IMAGE:671075}	671075 IMAGE Consortium	AA240994
8438 F	4 0216A1G4	IMAGE EST {IMAGE:658922}	658922 IMAGE Consortium	AA218316
8439 F	6 0216A1G4	M.musculus p16K gene for 16 kDa protein {IMAGE:671667}	671667 IMAGE Consortium	AA240897
8440 F	8 0216A1G4	IMAGE EST {IMAGE:671774}	671774 IMAGE Consortium	AA242235
8441 F	10 0216A1G4	ESTs {IMAGE:660220}	660220 IMAGE Consortium	AA220440
8442 F	12 0216A1G4	Mus musculus (clone pMNF-66) 66 kDa neurofilament protein NF-66 mRNA, complete cds {IMAGE:660756}	660756 IMAGE Consortium	AA218283
8443 H	2 0216A1G4	IMAGE EST {IMAGE:659939}	659939 IMAGE Consortium	AA219986

8444 H	4 0216A1G4	IMAGE EST {IMAGE:660030}	660030 IMAGE Consortium	AA220636
8445 H	6 0216A1G4	IMAGE EST {IMAGE:672496}	672496 IMAGE Consortium	AA231302
8446 H	8 0216A1G4	ESTs {IMAGE:672906}	672906 IMAGE Consortium	AA231433
8447 H	10 0216A1G4	ESTs, Highly similar to HYPOTHETICAL 167.8 KD PROTEIN CCE1-CAP1 INTERGENIC REGION [ <i>Saccharomyces cerevisiae</i> ] {IMAGE:660932}	660932 IMAGE Consortium	AA216947
8448 H	12 0216A1G4	ESTs {IMAGE:669953}	669953 IMAGE Consortium	AA230655
8449 B	2 021DA1G5	ESTs {IMAGE:660997}	660997 IMAGE Consortium	AA253853
8450 B	4 021DA1G5	<i>M.musculus</i> mRNA for casein kinase I-alpha {IMAGE:661005}	661005 IMAGE Consortium	AA230698
8451 B	6 021DA1G5	Hydroxysteroid 17-beta dehydrogenase 1 {IMAGE:677046}	677046 IMAGE Consortium	AA213048
8452 B	8 021DA1G5	IMAGE EST {IMAGE:670742}	670742 IMAGE Consortium	AA230829
8453 B	10 021DA1G5	IMAGE EST {IMAGE:670775}	670775 IMAGE Consortium	AA230950
8454 B	12 021DA1G5	Guanidinoacetate methyltransferase {IMAGE:671133}	671133 IMAGE Consortium	AA222404
8455 D	2 021DA1G5	<i>Mus musculus</i> cytochrome P450 monooxygenase (CYP2J5) mRNA, complete cds {IMAGE:676630}	676630 IMAGE Consortium	AA212435
8456 D	4 021DA1G5	IMAGE EST {IMAGE:676879}	676879 IMAGE Consortium	AA212961
8457 D	6 021DA1G5	IMAGE EST {IMAGE:677575}	677575 IMAGE Consortium	AA212892
8458 D	8 021DA1G5	ESTs {IMAGE:677564}	677564 IMAGE Consortium	AA213082
8459 D	10 021DA1G5	ESTs {IMAGE:659193}	659193 IMAGE Consortium	AA218384
8460 D	12 021DA1G5	<i>Mus musculus</i> ret-11 mRNA, complete cds {IMAGE:659182}	659182 IMAGE Consortium	AA218103
8461 F	2 021DA1G5	ESTs {IMAGE:677519}	677519 IMAGE Consortium	AA213240
8462 F	4 021DA1G5	<i>Mus musculus</i> mRNA for AMY-1, complete cds {IMAGE:658871}	658871 IMAGE Consortium	AA217989
8463 F	6 021DA1G5	Insulin-like growth factor binding protein 5 {IMAGE:671661}	671661 IMAGE Consortium	AA241784
8464 F	8 021DA1G5	Small proline-rich protein 1A {IMAGE:672405}	672405 IMAGE Consortium	AA230988
8465 F	10 021DA1G5	TALIN {IMAGE:676376}	676376 IMAGE Consortium	AA208883
8466 F	12 021DA1G5	Retinoid X receptor alpha {IMAGE:676526}	676526 IMAGE Consortium	AA209592
8467 H	2 021DA1G5	ESTs, Weakly similar to C30B5.2 [ <i>C.elegans</i> ] {IMAGE:660608}	660608 IMAGE Consortium	AA222930
8468 H	4 021DA1G5	IMAGE EST {IMAGE:676305}	676305 IMAGE Consortium	AA208876
8469 H	6 021DA1G5	CALGRANULIN A {IMAGE:669813}	669813 IMAGE Consortium	AA230451
8470 H	8 021DA1G5	IMAGE EST {IMAGE:670344}	670344 IMAGE Consortium	AA221233
8471 H	10 021DA1G5	IMAGE EST {IMAGE:678745}	678745 IMAGE Consortium	AA239481
8472 H	12 021DA1G5	IMAGE EST {IMAGE:678827}	678827 IMAGE Consortium	AA250590
8473 B	2 021KA1G6	ESTs {IMAGE:694351}	694351 IMAGE Consortium	AA220059
8474 B	4 021KA1G6	IMAGE EST {IMAGE:681137}	681137 IMAGE Consortium	AA242115
8475 B	6 021KA1G6	VITAMIN K-DEPENDENT PROTEIN S PRECURSOR {IMAGE:681615}	681615 IMAGE Consortium	AA237828
8476 B	8 021KA1G6	IMAGE EST {IMAGE:679326}	679326 IMAGE Consortium	AA244526
8477 B	10 021KA1G6	IMAGE EST {IMAGE:692699}	692699 IMAGE Consortium	AA239666
8478 B	12 021KA1G6	ESTs, Highly similar to TYROSINE AMINOTRANSFERASE [ <i>Homo sapiens</i> ] {IMAGE:680259}	680259 IMAGE Consortium	AA238289

8479 D	2 021KA1G6	ESTs {IMAGE:679244}	679244 IMAGE Consortium	AA244536
8480 D	4 021KA1G6	IMAGE EST {IMAGE:680307}	680307 IMAGE Consortium	AA238305
8481 D	6 021KA1G6	IMAGE EST {IMAGE:680149}	680149 IMAGE Consortium	AA237895
8482 D	8 021KA1G6	Urate oxidase {IMAGE:680254}	680254 IMAGE Consortium	AA237345
8483 D	10 021KA1G6	IMAGE EST {IMAGE:680454}	680454 IMAGE Consortium	AA237176
8484 D	12 021KA1G6	ESTs {IMAGE:680419}	680419 IMAGE Consortium	AA238230
8485 F	2 021KA1G6	ESTs {IMAGE:692963}	692963 IMAGE Consortium	AA238472
8486 F	4 021KA1G6	Amino acid transporter, cationic 2 (low affinity) {IMAGE:693244}	693244 IMAGE Consortium	AA250170
8487 F	6 021KA1G6	ESTs, Moderately similar to hTOM34p [H.sapiens] {IMAGE:693114}	693114 IMAGE Consortium	AA250161
8488 F	8 021KA1G6	IMAGE EST {IMAGE:693392}	693392 IMAGE Consortium	AA245312
8489 F	10 021KA1G6	IMAGE EST {IMAGE:693767}	693767 IMAGE Consortium	AA265282
8490 F	12 021KA1G6	Mus musculus beta-galactoside binding lectin mRNA, complete cds {IMAGE:680815}	680815 IMAGE Consortium	AA250039
8491 H	2 021KA1G6	Proprotein convertase subtilisin/kexin type 3 {IMAGE:680455}	680455 IMAGE Consortium	AA237179
8492 H	4 021KA1G6	ESTs, Weakly similar to C06G3.11 [C.elegans] {IMAGE:680349}	680349 IMAGE Consortium	AA238154
8493 H	6 021KA1G6	IMAGE EST {IMAGE:681105}	681105 IMAGE Consortium	AA242539
8494 H	8 021KA1G6	Extracellular matrix protein 1 {IMAGE:678765}	678765 IMAGE Consortium	AA237378
8495 H	10 021KA1G6	Mus musculus CAMP metalloproteinase mRNA, complete cds {IMAGE:692257}	692257 IMAGE Consortium	AA238244
8496 H	12 021KA1G6	PROTEIN C PRECURSOR {IMAGE:692609}	692609 IMAGE Consortium	AA250213
8497 B	2 021RA1G7	Sterol O-acyltransferase 1 {IMAGE:680485}	680485 IMAGE Consortium	AA237241
8498 B	4 021RA1G7	ESTs {IMAGE:681009}	681009 IMAGE Consortium	AA250079
8499 B	6 021RA1G7	Mus musculus BAP-135 homolog (Diws1t) mRNA, complete cds {IMAGE:694365}	694365 IMAGE Consortium	AA220113
8500 B	8 021RA1G7	ESTs {IMAGE:681446}	681446 IMAGE Consortium	AA244741
8501 B	10 021RA1G7	ESTs {IMAGE:681500}	681500 IMAGE Consortium	AA237628
8502 B	12 021RA1G7	IMAGE EST {IMAGE:679356}	679356 IMAGE Consortium	AA244770
8503 D	2 021RA1G7	ESTs {IMAGE:692546}	692546 IMAGE Consortium	AA250615
8504 D	4 021RA1G7	ESTs {IMAGE:692799}	692799 IMAGE Consortium	AA239913
8505 D	6 021RA1G7	IMAGE EST {IMAGE:680146}	680146 IMAGE Consortium	AA237870
8506 D	8 021RA1G7	Mus musculus von Hippel-Lindau binding protein homolog (vbp-1) mRNA, complete cds {IMAGE:693287}	693287 IMAGE Consortium	AA397091
8507 D	10 021RA1G7	IMAGE EST {IMAGE:693712}	693712 IMAGE Consortium	AA265270
8508 D	12 021RA1G7	ESTs {IMAGE:680687}	680687 IMAGE Consortium	AA237469
8509 F	2 021RA1G7	GLUTAMINE SYNTHETASE {IMAGE:693146}	693146 IMAGE Consortium	AA253677
8510 F	4 021RA1G7	IMAGE EST {IMAGE:693248}	693248 IMAGE Consortium	AA396996
8511 F	6 021RA1G7	ESTs, Highly similar to MITOCHONDRIAL INNER MEMBRANE PROTEASE SUBUNIT 1 [Saccharomyces cerevisiae] {IMAGE:693456}	693456 IMAGE Consortium	AA245155
8512 F	8 021RA1G7	ESTs, Moderately similar to KIAA0409 [H.sapiens]	693893 IMAGE Consortium	AA238815

		{IMAGE:693893}		
8513 F	10 021RA1G7	Hemolytic complement {IMAGE:678523}	678523 IMAGE Consortium	AA237264
8514 F	12 021RA1G7	Presenilin 1 {IMAGE:681670}	681670 IMAGE Consortium	AA237995
8515 H	2 021RA1G7	Mus musculus protein co-factor mRNA, complete cds {IMAGE:694133}	694133 IMAGE Consortium	AA239009
8516 H	4 021RA1G7	Asialoglycoprotein receptor 1 {IMAGE:693880}	693880 IMAGE Consortium	AA238685
8517 H	6 021RA1G7	ESTs {IMAGE:681314}	681314 IMAGE Consortium	AA241936
8518 H	8 021RA1G7	IMAGE EST {IMAGE:678511}	678511 IMAGE Consortium	AA237127
8519 H	10 021RA1G7	IMAGE EST {IMAGE:679011}	679011 IMAGE Consortium	AA245806
8520 H	12 021RA1G7	ESTs {IMAGE:678831}	678831 IMAGE Consortium	AA250458
8521 B	2 021YA1G8	IMAGE EST {IMAGE:679088}	679088 IMAGE Consortium	AA245657
8522 B	4 021YA1G8	ESTs {IMAGE:692545}	692545 IMAGE Consortium	AA250659
8523 B	6 021YA1G8	ESTs, Highly similar to CLATHRIN COAT ASSEMBLY PROTEIN AP17 [Rattus norvegicus] {IMAGE:692299}	692299 IMAGE Consortium	AA260594
8524 B	8 021YA1G8	ESTs, Highly similar to serine protease [H.sapiens] {IMAGE:692768}	692768 IMAGE Consortium	AA239856
8525 B	10 021YA1G8	IMAGE EST {IMAGE:692940}	692940 IMAGE Consortium	AA238462
8526 B	12 021YA1G8	Voltage-dependent anion channel 3 {IMAGE:680191}	680191 IMAGE Consortium	AA237920
8527 D	2 021YA1G8	ESTs, Highly similar to UBIQUITIN-CONJUGATING ENZYME E2-17 KD [Drosophila melanogaster] {IMAGE:718282}	718282 IMAGE Consortium	AA261605
8528 D	4 021YA1G8	Zinc finger protein 93 {IMAGE:718197}	718197 IMAGE Consortium	AA266357
8529 D	6 021YA1G8	Indian hedgehog homolog, (Drosophila) {IMAGE:699460}	699460 IMAGE Consortium	AA245525
8530 D	8 021YA1G8	ESTs, Moderately similar to H-2 class I histocompatibility antigen T15 [M.musculus] {IMAGE:699253}	699253 IMAGE Consortium	AA245560
8531 D	10 021YA1G8	ESTs, Highly similar to MITOCHONDRIAL ELONGATION FACTOR TS PRECURSOR [Homo sapiens] {IMAGE:699237}	699237 IMAGE Consortium	AA245481
8532 D	12 021YA1G8	IMAGE EST {IMAGE:717121}	717121 IMAGE Consortium	AA266040
8533 F	2 021YA1G8	IMAGE EST {IMAGE:695246}	695246 IMAGE Consortium	AA220484
8534 F	4 021YA1G8	M.musculus mRNA for Pr22 protein {IMAGE:716630}	716630 IMAGE Consortium	AA265396
8535 F	6 021YA1G8	IMAGE EST {IMAGE:716834}	716834 IMAGE Consortium	AA266389
8536 F	8 021YA1G8	ESTs {IMAGE:696543}	696543 IMAGE Consortium	AA239746
8537 F	10 021YA1G8	IMAGE EST {IMAGE:717466}	717466 IMAGE Consortium	AA403429
8538 F	12 021YA1G8	IMAGE EST {IMAGE:717683}	717683 IMAGE Consortium	AA250385
8539 H	2 021YA1G8	Mus musculus mRNA for 1-acyl-sn-glycerol-3-phosphate acyltransferase, complete cds {IMAGE:699479}	699479 IMAGE Consortium	AA245633
8540 H	4 021YA1G8	IMAGE EST {IMAGE:716941}	716941 IMAGE Consortium	AA266097
8541 H	6 021YA1G8	Mus musculus transcobalmin II (Tcn2) mRNA, complete cds {IMAGE:697010}	697010 IMAGE Consortium	AA220699
8542 H	8 021YA1G8	Hemoglobin alpha, adult chain 1 {IMAGE:696858}	696858 IMAGE Consortium	AA220584
8543 H	10 021YA1G8	IMAGE EST {IMAGE:697374}	697374 IMAGE Consortium	AA242197
8544 H	12 021YA1G8	IMAGE EST {IMAGE:717883}	717883 IMAGE Consortium	AA250555
8545 B	2 021YA1G9	Bcl2 homologous antagonist/killer {IMAGE:696909}	696909 IMAGE Consortium	AA220617

8546 B	4 0215A1G9	ESTs, Highly similar to RAS-RELATED PROTEIN RAB-25 [Oryctolagus cuniculus] {IMAGE:697383}	697383 IMAGE Consortium	AA244806
8547 B	6 0215A1G9	ESTs {IMAGE:718110}	718110 IMAGE Consortium	AA266400
8548 B	8 0215A1G9	IMAGE EST {IMAGE:717977}	717977 IMAGE Consortium	AA261171
8549 B	10 0215A1G9	ESTs {IMAGE:718375}	718375 IMAGE Consortium	AA267726
8550 B	12 0215A1G9	ESTs {IMAGE:695037}	695037 IMAGE Consortium	AA221124
8551 D	2 0215A1G9	Mus musculus mRNA capping enzyme (MCE) mRNA, complete cds {IMAGE:552577}	552577 IMAGE Consortium	AA096583
8552 D	4 0215A1G9	Finkel-Biskis-Reilly murine sarcoma virus (FBR-MuSV) ubiquitously expressed (fox derived) {IMAGE:698100}	698100 IMAGE Consortium	AA239437
8553 D	6 0215A1G9	ESTs {IMAGE:699289}	699289 IMAGE Consortium	AA245396
8554 D	8 0215A1G9	Mus musculus alpha(1,3)fucosyltransferase Fuc-TVII gene, complete cds {IMAGE:699343}	699343 IMAGE Consortium	AA245378
8555 D	10 0215A1G9	IMAGE EST {IMAGE:699280}	699280 IMAGE Consortium	AA245545
8556 D	12 0215A1G9	IMAGE EST {IMAGE:716523}	716523 IMAGE Consortium	AA265488
8557 F	2 0215A1G9	IMAGE EST {IMAGE:718543}	718543 IMAGE Consortium	AA266610
8558 F	4 0215A1G9	ESTs {IMAGE:718690}	718690 IMAGE Consortium	AA265695
8559 F	6 0215A1G9	ESTs, Weakly similar to ORF YDL015c [S.cerevisiae] {IMAGE:695279}	695279 IMAGE Consortium	AA220458
8560 F	8 0215A1G9	Mouse protein disulfide isomerase-related protein (ERp72) mRNA, complete cds {IMAGE:695696}	695696 IMAGE Consortium	AA222205
8561 F	10 0215A1G9	IMAGE EST {IMAGE:696681}	696681 IMAGE Consortium	AA240266
8562 F	12 0215A1G9	ESTs, Highly similar to HYPOTHETICAL 52.8 KD PROTEIN T05E11.5 IN CHROMOSOME IV [Caenorhabditis elegans] {IMAGE:717316}	717316 IMAGE Consortium	AA403866
8563 H	2 0215A1G9	Surfactant associated protein C {IMAGE:695697}	695697 IMAGE Consortium	AA222208
8564 H	4 0215A1G9	M.musculus mRNA for ribosomal protein S5 {IMAGE:696769}	696769 IMAGE Consortium	AA240279
8565 H	6 0215A1G9	IMAGE EST {IMAGE:696635}	696635 IMAGE Consortium	AA241108
8566 H	8 0215A1G9	IMAGE EST {IMAGE:697208}	697208 IMAGE Consortium	AA221820
8567 H	10 0215A1G9	ESTs {IMAGE:697537}	697537 IMAGE Consortium	AA250652
8568 H	12 0215A1G9	Aconitase 1 {IMAGE:697949}	697949 IMAGE Consortium	AA238899
8569 B	2 021PA1GA	IMAGE EST {IMAGE:719455}	719455 IMAGE Consortium	AA255243
8570 B	4 021PA1GA	ESTs {IMAGE:722330}	722330 IMAGE Consortium	AA254237
8571 B	6 021PA1GA	ESTs {IMAGE:722193}	722193 IMAGE Consortium	AA254032
8572 B	8 021PA1GA	IMAGE EST {IMAGE:720014}	720014 IMAGE Consortium	AA260980
8573 B	10 021PA1GA	ESTs {IMAGE:722944}	722944 IMAGE Consortium	AA254380
8574 B	12 021PA1GA	ESTs, Highly similar to casein kinase II substrate [R.norvegicus] {IMAGE:720647}	720647 IMAGE Consortium	AA261368
8575 D	2 021PA1GA	Trans-acting transcription factor 1 {IMAGE:719204}	719204 IMAGE Consortium	AA267163
8576 D	4 021PA1GA	ESTs {IMAGE:719840}	719840 IMAGE Consortium	AA260427
8577 D	6 021PA1GA	IMAGE EST {IMAGE:720096}	720096 IMAGE Consortium	AA254822
8578 D	8 021PA1GA	ESTs {IMAGE:722849}	722849 IMAGE Consortium	AA254370
8579 D	10 021PA1GA	Mus musculus cleavage and polyadenylation specificity factor	723260 IMAGE Consortium	AA267638

		(MCPSF) mRNA, complete cds {IMAGE:723260}		
8580 D	12 021PA1GA	IMAGE EST {IMAGE:733344}	733344 IMAGE Consortium	AA261326
8581 F	2 021PA1GA	ESTs {IMAGE:721937}	721937 IMAGE Consortium	AA267979
8582 F	4 021PA1GA	IMAGE EST {IMAGE:719579}	719579 IMAGE Consortium	AA254859
8583 F	6 021PA1GA	ESTs, Highly similar to LEUCINE-RICH ALPHA-2-GLYCOPROTEIN [Homo sapiens] {IMAGE:720207}	720207 IMAGE Consortium	AA255171
8584 F	8 021PA1GA	IMAGE EST {IMAGE:720554}	720554 IMAGE Consortium	AA261696
8585 F	10 021PA1GA	ESTs, Highly similar to COMPLEMENT C1R COMPONENT PRECURSOR [Homo sapiens] {IMAGE:720566}	720566 IMAGE Consortium	AA261393
8586 F	12 021PA1GA	IMAGE EST {IMAGE:723261}	723261 IMAGE Consortium	AA267772
8587 H	2 021PA1GA	ESTs {IMAGE:722403}	722403 IMAGE Consortium	AA254186
8588 H	4 021PA1GA	IMAGE EST {IMAGE:722648}	722648 IMAGE Consortium	AA260702
8589 H	6 021PA1GA	IMAGE EST {IMAGE:720567}	720567 IMAGE Consortium	AA261397
8590 H	8 021PA1GA	ESTs {IMAGE:720772}	720772 IMAGE Consortium	AA266835
8591 H	10 021PA1GA	ESTs, Highly similar to PHOSPHORYLASE B KINASE GAMMA CATALYTIC CHAIN, TESTIS ISOFORM [Rattus norvegicus] {IMAGE:720780}	720780 IMAGE Consortium	AA266813
8592 H	12 021PA1GA	IMAGE EST {IMAGE:720665}	720665 IMAGE Consortium	AA267268
8593 B	2 021WA1GB	IMAGE EST {IMAGE:697370}	697370 IMAGE Consortium	AA242195
8594 B	4 021WA1GB	IMAGE EST {IMAGE:697433}	697433 IMAGE Consortium	AA253808
8595 B	6 021WA1GB	ESTs {IMAGE:697662}	697662 IMAGE Consortium	AA238618
8596 B	8 021WA1GB	IMAGE EST {IMAGE:718372}	718372 IMAGE Consortium	AA267721
8597 B	10 021WA1GB	ESTs, Weakly similar to CIRP [M.musculus] {IMAGE:718640}	718640 IMAGE Consortium	AA265671
8598 B	12 021WA1GB	ESTs {IMAGE:718716}	718716 IMAGE Consortium	AA265716
8599 D	2 021WA1GB	ESTs {IMAGE:716698}	716698 IMAGE Consortium	AA265234
8600 D	4 021WA1GB	Carnitine palmitoyltransferase 1, liver {IMAGE:717056}	717056 IMAGE Consortium	AA267525
8601 D	6 021WA1GB	ESTs {IMAGE:696524}	696524 IMAGE Consortium	AA239508
8602 D	8 021WA1GB	IMAGE EST {IMAGE:717904}	717904 IMAGE Consortium	AA250558
8603 D	10 021WA1GB	IMAGE EST {IMAGE:718196}	718196 IMAGE Consortium	AA266353
8604 D	12 021WA1GB	IMAGE EST {IMAGE:718425}	718425 IMAGE Consortium	AA266782
8605 F	2 021WA1GB	Mouse carboxyesterase mRNA {IMAGE:719734}	719734 IMAGE Consortium	AA254921
8606 F	4 021WA1GB	ESTs, Moderately similar to JAK3 [M.musculus] {IMAGE:722568}	722568 IMAGE Consortium	AA260505
8607 F	6 021WA1GB	ESTs {IMAGE:722775}	722775 IMAGE Consortium	AA254552
8608 F	8 021WA1GB	ESTs, Moderately similar to ISOLEUCYL-TRNA SYNTHETASE [Staphylococcus aureus] {IMAGE:723128}	723128 IMAGE Consortium	AA267557
8609 F	10 021WA1GB	IMAGE EST {IMAGE:723066}	723066 IMAGE Consortium	AA254409
8610 F	12 021WA1GB	ESTs {IMAGE:720951}	720951 IMAGE Consortium	AA266737
8611 H	2 021WA1GB	IMAGE EST {IMAGE:721933}	721933 IMAGE Consortium	AA267977
8612 H	4 021WA1GB	IMAGE EST {IMAGE:719573}	719573 IMAGE Consortium	AA254697
8613 H	6 021WA1GB	ESTs {IMAGE:719615}	719615 IMAGE Consortium	AA254957
8614 H	8 021WA1GB	Mus musculus poly(ADP-ribose) glycohydrolase (Parg) mRNA, complete cds {IMAGE:722592}	722592 IMAGE Consortium	AA260570

8615 H	10 021WA1GB	IMAGE EST {IMAGE:723297}	723297 IMAGE Consortium	AA267826
8616 H	12 021WA1GB	ESTs {IMAGE:723258}	723258 IMAGE Consortium	AA267637
8617 B	2 0213A1GC	GATA-binding protein 2 {IMAGE:719592}	719592 IMAGE Consortium	AA255034
8618 B	4 0213A1GC	ESTs {IMAGE:722362}	722362 IMAGE Consortium	AA254109
8619 B	6 0213A1GC	IMAGE EST {IMAGE:722625}	722625 IMAGE Consortium	AA260445
8620 B	8 0213A1GC	ESTs {IMAGE:720844}	720844 IMAGE Consortium	AA267166
8621 B	10 0213A1GC	ESTs {IMAGE:720828}	720828 IMAGE Consortium	AA266957
8622 B	12 0213A1GC	ESTs {IMAGE:720857}	720857 IMAGE Consortium	AA267057
8623 D	2 0213A1GC	ESTs {IMAGE:720031}	720031 IMAGE Consortium	AA255275
8624 D	4 0213A1GC	ESTs, Weakly similar to cDNA EST EMBL:Z14593 comes from this gene [C.elegans] {IMAGE:722581}	722581 IMAGE Consortium	AA260576
8625 D	6 0213A1GC	ESTs, Weakly similar to PROTEIN DISULFIDE ISOMERASE PRECURSOR [M.musculus] {IMAGE:722782}	722782 IMAGE Consortium	AA254546
8626 D	8 0213A1GC	IMAGE EST {IMAGE:720988}	720988 IMAGE Consortium	AA267590
8627 D	10 0213A1GC	IMAGE EST {IMAGE:733517}	733517 IMAGE Consortium	AA259873
8628 D	12 0213A1GC	ESTs {IMAGE:720958}	720958 IMAGE Consortium	AA266859
8629 F	2 0213A1GC	IMAGE EST {IMAGE:720032}	720032 IMAGE Consortium	AA260985
8630 F	4 0213A1GC	Mago-nashi homolog, proliferation-associated (Drosophila) {IMAGE:722821}	722821 IMAGE Consortium	AA254528
8631 F	6 0213A1GC	IMAGE EST {IMAGE:722954}	722954 IMAGE Consortium	AA254268
8632 F	8 0213A1GC	ESTs {IMAGE:720678}	720678 IMAGE Consortium	AA267274
8633 F	10 0213A1GC	ESTs, Highly similar to RAS-RELATED PROTEIN RAB-5A [Canis familiaris] {IMAGE:734064}	734064 IMAGE Consortium	AA269366
8634 F	12 0213A1GC	ESTs {IMAGE:733834}	733834 IMAGE Consortium	AA268155
8635 H	2 0213A1GC	ESTs {IMAGE:720284}	720284 IMAGE Consortium	AA261483
8636 H	4 0213A1GC	ESTs {IMAGE:720774}	720774 IMAGE Consortium	AA266836
8637 H	6 0213A1GC	IMAGE EST {IMAGE:733661}	733661 IMAGE Consortium	AA259502
8638 H	8 0213A1GC	ESTs, Highly similar to PROTEIN PHOSPHATASE PP2A, 65 KD REGULATORY SUBUNIT, ALPHA ISOFORM [Homo sapiens] {IMAGE:733882}	733882 IMAGE Consortium	AA268171
8639 H	10 0213A1GC	ESTs, Highly similar to PROPIONYL-COA CARBOXYLASE BETA CHAIN PRECURSOR [R.norvegicus] {IMAGE:719007}	719007 IMAGE Consortium	AA267533
8640 H	12 0213A1GC	Calcyclin {IMAGE:721906}	721906 IMAGE Consortium	AA267952
8641 B	2 021AA1GD	Ubiquitin conjugating enzyme E3A {IMAGE:721372}	721372 IMAGE Consortium	AA267222
8642 B	4 021AA1GD	ESTs {IMAGE:721239}	721239 IMAGE Consortium	AA267015
8643 B	6 021AA1GD	ESTs {IMAGE:721421}	721421 IMAGE Consortium	AA267569
8644 B	8 021AA1GD	Mus musculus cdc2/CDC28-like protein kinase 3 (Clk3) mRNA, complete cds {IMAGE:734121}	734121 IMAGE Consortium	AA268598
8645 B	10 021AA1GD	ESTs, Highly similar to KIAA0398 [H.sapiens] {IMAGE:721931}	721931 IMAGE Consortium	AA267976
8646 B	12 021AA1GD	ESTs, Highly similar to HISTIDINE-RICH GLYCOPROTEIN PRECURSOR [Homo sapiens] {IMAGE:719667}	719667 IMAGE Consortium	AA255115
8647 D	2 021AA1GD	IMAGE EST {IMAGE:721637}	721637 IMAGE Consortium	AA260679

8648 D	4 021AA1GD	Inter-alpha trypsin inhibitor, heavy chain 1 {IMAGE:734289}	734289 IMAGE Consortium	AA268664
8649 D	6 021AA1GD	IMAGE EST {IMAGE:734688}	734688 IMAGE Consortium	AA261160
8650 D	8 021AA1GD	ESTs {IMAGE:734660}	734660 IMAGE Consortium	AA259534
8651 D	10 021AA1GD	ESTs {IMAGE:735058}	735058 IMAGE Consortium	AA259894
8652 D	12 021AA1GD	C-reactive protein, petaxin related {IMAGE:737941}	737941 IMAGE Consortium	AA271451
8653 F	2 021AA1GD	ESTs, Moderately similar to K10B2.4 [C.elegans] {IMAGE:747674}	747674 IMAGE Consortium	AA268773
8654 F	4 021AA1GD	ESTs, Weakly similar to C43E11.9 [C.elegans] {IMAGE:747789}	747789 IMAGE Consortium	AA274849
8655 F	6 021AA1GD	ESTs {IMAGE:737390}	737390 IMAGE Consortium	AA277583
8656 F	8 021AA1GD	Mus musculus AMP activated protein kinase mRNA, complete cds {IMAGE:734780}	734780 IMAGE Consortium	AA259329
8657 F	10 021AA1GD	IMAGE EST {IMAGE:734939}	734939 IMAGE Consortium	AA261222
8658 F	12 021AA1GD	ESTs {IMAGE:734995}	734995 IMAGE Consortium	AA260155
8659 H	2 021AA1GD	ESTs {IMAGE:734338}	734338 IMAGE Consortium	AA268694
8660 H	4 021AA1GD	IMAGE EST {IMAGE:734665}	734665 IMAGE Consortium	AA261209
8661 H	6 021AA1GD	IMAGE EST {IMAGE:734829}	734829 IMAGE Consortium	AA259386
8662 H	8 021AA1GD	IMAGE EST {IMAGE:735227}	735227 IMAGE Consortium	AA259552
8663 H	10 021AA1GD	ESTs {IMAGE:734972}	734972 IMAGE Consortium	AA260104
8664 H	12 021AA1GD	ESTs {IMAGE:735607}	735607 IMAGE Consortium	AA270170
8665 B	2 021HA1GE	ESTs {IMAGE:735773}	735773 IMAGE Consortium	AA270087
8666 B	4 021HA1GE	ESTs, Weakly similar to Similarity to S.pombe hypothetical protein C1D4.09C [C.elegans] {IMAGE:735647}	735647 IMAGE Consortium	AA270218
8667 B	6 021HA1GE	ESTs {IMAGE:736229}	736229 IMAGE Consortium	AA270412
8668 B	8 021HA1GE	IMAGE EST {IMAGE:736263}	736263 IMAGE Consortium	AA270432
8669 B	10 021HA1GE	Mouse mRNA for alpha-2,3-sialyltransferase, complete cds {IMAGE:736562}	736562 IMAGE Consortium	AA270471
8670 B	12 021HA1GE	ESTs, Highly similar to QUEUEINE TRNA-RIBOSYLTRANSFERASE [H.sapiens] {IMAGE:736721}	736721 IMAGE Consortium	AA270950
8671 D	2 021HA1GE	IMAGE EST {IMAGE:746594}	746594 IMAGE Consortium	AA268006
8672 D	4 021HA1GE	ESTs, Weakly similar to cDNA EST EMBL:D71941 comes from this gene [C.elegans] {IMAGE:746649}	746649 IMAGE Consortium	AA268144
8673 D	6 021HA1GE	M.musculus mRNA for calpain-like protease {IMAGE:747101}	747101 IMAGE Consortium	AA260476
8674 D	8 021HA1GE	ESTs {IMAGE:747038}	747038 IMAGE Consortium	AA260280
8675 D	10 021HA1GE	ESTs, Highly similar to ALPHA-L-IDURONIDASE PRECURSOR [M.musculus] {IMAGE:747385}	747385 IMAGE Consortium	AA268228
8676 D	12 021HA1GE	Mus musculus cis-retinol androgen dehydrogenase 1 mRNA, complete cds {IMAGE:747207}	747207 IMAGE Consortium	AA260409
8677 F	2 021HA1GE	IMAGE EST {IMAGE:735557}	735557 IMAGE Consortium	AA270035
8678 F	4 021HA1GE	ESTs, Moderately similar to HYPOTHETICAL PROTEIN KIAA0039 [H.sapiens] {IMAGE:735374}	735374 IMAGE Consortium	AA269653
8679 F	6 021HA1GE	IMAGE EST {IMAGE:746506}	746506 IMAGE Consortium	AA260199
8680 F	8 021HA1GE	IMAGE EST {IMAGE:747127}	747127 IMAGE Consortium	AA260483



8681 F	10 021HA1GE	ESTs {IMAGE:747111}	747111 IMAGE Consortium	AA274981
8682 F	12 021HA1GE	IMAGE EST {IMAGE:747362}	747362 IMAGE Consortium	AA268218
8683 H	2 021HA1GE	ESTs, Weakly similar to PROTEIN PHOSPHATASE PP2A, 72 KD REGULATORY SUBUNIT [H.sapiens] {IMAGE:735357}	735357 IMAGE Consortium	AA269494
8684 H	4 021HA1GE	DIK1-like homolog (Drosophila) {IMAGE:746888}	746888 IMAGE Consortium	AA274459
8685 H	6 021HA1GE	IMAGE EST {IMAGE:736017}	736017 IMAGE Consortium	AA270263
8686 H	8 021HA1GE	ESTs {IMAGE:747327}	747327 IMAGE Consortium	AA274922
8687 H	10 021HA1GE	IMAGE EST {IMAGE:736579}	736579 IMAGE Consortium	AA270632
8688 H	12 021HA1GE	IMAGE EST {IMAGE:736858}	736858 IMAGE Consortium	AA271373
8689 B	2 021OA1GF	ESTs {IMAGE:737596}	737596 IMAGE Consortium	AA277482
8690 B	4 021OA1GF	ESTs {IMAGE:737800}	737800 IMAGE Consortium	AA277324
8691 B	6 021OA1GF	IMAGE EST {IMAGE:734949}	734949 IMAGE Consortium	AA261227
8692 B	8 021OA1GF	ESTs {IMAGE:746383}	746383 IMAGE Consortium	AA259960
8693 B	10 021OA1GF	Growth factor receptor bound protein 10 {IMAGE:746564}	746564 IMAGE Consortium	AA260248
8694 B	12 021OA1GF	T-cell acute lymphocytic leukemia 1 {IMAGE:736275}	736275 IMAGE Consortium	AA270232
8695 D	2 021OA1GF	IMAGE EST {IMAGE:737620}	737620 IMAGE Consortium	AA277507
8696 D	4 021OA1GF	ESTs, Moderately similar to tetratricopeptide repeat protein [H.sapiens] {IMAGE:735061}	735061 IMAGE Consortium	AA259927
8697 D	6 021OA1GF	IMAGE EST {IMAGE:738185}	738185 IMAGE Consortium	AA271059
8698 D	8 021OA1GF	ESTs, Highly similar to KIAA0017 [H.sapiens] {IMAGE:735617}	735617 IMAGE Consortium	AA269584
8699 D	10 021OA1GF	ESTs {IMAGE:735673}	735673 IMAGE Consortium	AA269627
8700 D	12 021OA1GF	IMAGE EST {IMAGE:735883}	735883 IMAGE Consortium	AA269699
8701 F	2 021OA1GF	ESTs {IMAGE:734870}	734870 IMAGE Consortium	AA259394
8702 F	4 021OA1GF	ESTs, Highly similar to PROCOLLAGEN-LYSINE,2-OXOGLUTARATE 5-DIOXYGENASE PRECURSOR [Homo sapiens] {IMAGE:735312}	735312 IMAGE Consortium	AA270001
8703 F	6 021OA1GF	IMAGE EST {IMAGE:735674}	735674 IMAGE Consortium	AA269686
8704 F	8 021OA1GF	ESTs, Weakly similar to contains similarity to ATP/GTP-binding site motif [C.elegans] {IMAGE:746916}	746916 IMAGE Consortium	AA274822
8705 F	10 021OA1GF	ESTs, Highly similar to PROBABLE PHOSPHOSERINE AMINOTRANSFERASE [Oryctolagus cuniculus] {IMAGE:746853}	746853 IMAGE Consortium	AA274533
8706 F	12 021OA1GF	ESTs, Weakly similar to coded for by C. elegans cDNA yk16b1.3 [C.elegans] {IMAGE:736416}	736416 IMAGE Consortium	AA270716
8707 H	2 021OA1GF	ESTs, Weakly similar to nuclear factor YC [M.musculus] {IMAGE:735528}	735528 IMAGE Consortium	AA270201
8708 H	4 021OA1GF	IMAGE EST {IMAGE:738239}	738239 IMAGE Consortium	AA271106
8709 H	6 021OA1GF	IMAGE EST {IMAGE:736205}	736205 IMAGE Consortium	AA270404
8710 H	8 021OA1GF	ESTs {IMAGE:747180}	747180 IMAGE Consortium	AA260397
8711 H	10 021OA1GF	IMAGE EST {IMAGE:747167}	747167 IMAGE Consortium	AA260293
8712 H	12 021OA1GF	Phosphatidylinositol glycan, class R {IMAGE:737364}	737364 IMAGE Consortium	AA277571

8713 B	2	021VA1GG	DNA primase, p49 subunit {IMAGE:746463}	746463	IMAGE Consortium	AA259900
8714 B	4	021VA1GG	Integrin alpha 6 {IMAGE:736738}	736738	IMAGE Consortium	AA270921
8715 B	6	021VA1GG	ESTs, Weakly similar to cDNA EST EMBL:T01157 comes from this gene [C.elegans] {IMAGE:747672}	747672	IMAGE Consortium	AA268772
8716 B	8	021VA1GG	ESTs {IMAGE:737174}	737174	IMAGE Consortium	AA272566
8717 B	10	021VA1GG	ESTs {IMAGE:747941}	747941	IMAGE Consortium	AA260950
8718 B	12	021VA1GG	IMAGE EST {IMAGE:748322}	748322	IMAGE Consortium	AA274649
8719 D	2	021VA1GG	IMAGE EST {IMAGE:749776}	749776	IMAGE Consortium	AA288611
8720 D	4	021VA1GG	ESTs {IMAGE:762483}	762483	IMAGE Consortium	AA274160
8721 D	6	021VA1GG	ESTs {IMAGE:750411}	750411	IMAGE Consortium	AA389961
8722 D	8	021VA1GG	ESTs {IMAGE:748466}	748466	IMAGE Consortium	AA268202
8723 D	10	021VA1GG	Cytochrome P450, 2b9, phenobarbital inducible, type a {IMAGE:748569}	748569	IMAGE Consortium	AA269533
8724 D	12	021VA1GG	Mus musculus mRNA for alpha-albumine protein {IMAGE:748648}	748648	IMAGE Consortium	AA270289
8725 F	2	021VA1GG	IMAGE EST {IMAGE:749941}	749941	IMAGE Consortium	AA388787
8726 F	4	021VA1GG	ESTs {IMAGE:748015}	748015	IMAGE Consortium	AA275014
8727 F	6	021VA1GG	IMAGE EST {IMAGE:748027}	748027	IMAGE Consortium	AA260317
8728 F	8	021VA1GG	ESTs, Highly similar to phosphatidylserine-binding phosphoprotein [H.sapiens] {IMAGE:750503}	750503	IMAGE Consortium	AA472899
8729 F	10	021VA1GG	IMAGE EST {IMAGE:750878}	750878	IMAGE Consortium	AA290206
8730 F	12	021VA1GG	IMAGE EST {IMAGE:751348}	751348	IMAGE Consortium	AA290417
8731 H	2	021VA1GG	ESTs {IMAGE:750205}	750205	IMAGE Consortium	AA396360
8732 H	4	021VA1GG	ESTs {IMAGE:750222}	750222	IMAGE Consortium	AA390084
8733 H	6	021VA1GG	ESTs {IMAGE:750495}	750495	IMAGE Consortium	AA390032
8734 H	8	021VA1GG	ESTs {IMAGE:750649}	750649	IMAGE Consortium	AA290108
8735 H	10	021VA1GG	IMAGE EST {IMAGE:748435}	748435	IMAGE Consortium	AA272859
8736 H	12	021VA1GG	ESTs, Highly similar to NADH-UBIQUINONE OXIDOREDUCTASE 13 KD-B SUBUNIT [Bos taurus] {IMAGE:748891}	748891	IMAGE Consortium	AA288040
8737 B	2	0212A1GH	IMAGE EST {IMAGE:749897}	749897	IMAGE Consortium	AA389271
8738 B	4	0212A1GH	ESTs {IMAGE:747968}	747968	IMAGE Consortium	AA260962
8739 B	6	0212A1GH	ESTs, Moderately similar to DEATH-ASSOCIATED PROTEIN 1 [H.sapiens] {IMAGE:748546}	748546	IMAGE Consortium	AA269524
8740 B	8	0212A1GH	Mus musculus NSD1 protein mRNA, complete cds {IMAGE:748555}	748555	IMAGE Consortium	AA269575
8741 B	10	0212A1GH	IMAGE EST {IMAGE:751560}	751560	IMAGE Consortium	AA395937
8742 B	12	0212A1GH	IMAGE EST {IMAGE:748939}	748939	IMAGE Consortium	AA288144
8743 D	2	0212A1GH	Mus musculus SKD1 protein mRNA, complete cds {IMAGE:762176}	762176	IMAGE Consortium	AA277275
8744 D	4	0212A1GH	ESTs, Moderately similar to cDNA EST yk400h8.3 comes from this gene [C.elegans] {IMAGE:750558}	750558	IMAGE Consortium	AA472933
8745 D	6	0212A1GH	Mus musculus Btk locus, alpha-D-galactosidase A (Ags), ribosomal protein (L44L), and Bruton's tyrosine kinase (Btk)	750282	IMAGE Consortium	AA396783

8746 D	8 0212A1GH	genes, complete cds {IMAGE:750282} Homolog of human MLLT2 unidentified gene {IMAGE:750930}	750930 IMAGE Consortium	AA289139
8747 D	10 0212A1GH	ESTs {IMAGE:751729}	751729 IMAGE Consortium	AA396132
8748 D	12 0212A1GH	IMAGE EST {IMAGE:751385}	751385 IMAGE Consortium	AA396312
8749 F	2 0212A1GH	Cytochrome P450, steroid inducible 3a11 {IMAGE:748318}	748318 IMAGE Consortium	AA268120
8750 F	4 0212A1GH	IMAGE EST {IMAGE:749053}	749053 IMAGE Consortium	AA288998
8751 F	6 0212A1GH	ESTs {IMAGE:751409}	751409 IMAGE Consortium	AA396362
8752 F	8 0212A1GH	ESTs {IMAGE:749313}	749313 IMAGE Consortium	AA288555
8753 F	10 0212A1GH	IMAGE EST {IMAGE:752144}	752144 IMAGE Consortium	AA289947
8754 F	12 0212A1GH	Glycosylation dependent cell adhesion molecule 1 {IMAGE:749605}	749605 IMAGE Consortium	AA288467
8755 H	2 0212A1GH	Mouse murinoglobulin (MUG2) mRNA, complete cds {IMAGE:748895}	748895 IMAGE Consortium	AA288042
8756 H	4 0212A1GH	IMAGE EST {IMAGE:751901}	751901 IMAGE Consortium	AA396249
8757 H	6 0212A1GH	IMAGE EST {IMAGE:749396}	749396 IMAGE Consortium	AA289616
8758 H	8 0212A1GH	IMAGE EST {IMAGE:752396}	752396 IMAGE Consortium	AA290033
8759 H	10 0212A1GH	ESTs, Highly similar to ACETYL-COA ACETYLTRANSFERASE PRECURSOR, MITOCHONDRIAL [Homo sapiens] {IMAGE:761668}	761668 IMAGE Consortium	AA272067
8760 H	12 0212A1GH	ESTs {IMAGE:749926}	749926 IMAGE Consortium	AA389278
8761 B	2 0219A1GI	ESTs {IMAGE:765304}	765304 IMAGE Consortium	AA275236
8762 B	4 0219A1GI	ESTs {IMAGE:765722}	765722 IMAGE Consortium	AA277076
8763 B	6 0219A1GI	ESTs {IMAGE:765660}	765660 IMAGE Consortium	AA275228
8764 B	8 0219A1GI	ESTs, Highly similar to ASPARTOACYLASE [Homo sapiens] {IMAGE:766467}	766467 IMAGE Consortium	AA275673
8765 B	10 0219A1GI	ESTs {IMAGE:763174}	763174 IMAGE Consortium	AA271288
8766 B	12 0219A1GI	Annexin VI, p68 {IMAGE:775597}	775597 IMAGE Consortium	AA275924
8767 D	2 0219A1GI	IMAGE EST {IMAGE:765770}	765770 IMAGE Consortium	AA275347
8768 D	4 0219A1GI	IMAGE EST {IMAGE:765766}	765766 IMAGE Consortium	AA275203
8769 D	6 0219A1GI	ESTs, Highly similar to HYPOTHETICAL PROTEIN S164 [H.sapiens] {IMAGE:766564}	766564 IMAGE Consortium	AA274725
8770 D	8 0219A1GI	IMAGE EST {IMAGE:763103}	763103 IMAGE Consortium	AA271331
8771 D	10 0219A1GI	M.musculus mRNA for ribosomal protein S5 {IMAGE:775645}	775645 IMAGE Consortium	AA275884
8772 D	12 0219A1GI	CD81 ANTIGEN {IMAGE:775893}	775893 IMAGE Consortium	AA276336
8773 F	2 0219A1GI	IMAGE EST {IMAGE:777319}	777319 IMAGE Consortium	AA276912
8774 F	4 0219A1GI	IMAGE EST {IMAGE:777472}	777472 IMAGE Consortium	AA276533
8775 F	6 0219A1GI	IMAGE EST {IMAGE:765890}	765890 IMAGE Consortium	AA277912
8776 F	8 0219A1GI	IMAGE EST {IMAGE:765935}	765935 IMAGE Consortium	AA277948
8777 F	10 0219A1GI	Mus musculus uridine kinase mRNA, partial cds {IMAGE:766505}	766505 IMAGE Consortium	AA275763
8778 F	12 0219A1GI	IMAGE EST {IMAGE:763597}	763597 IMAGE Consortium	AA285951
8779 H	2 0219A1GI	ESTs {IMAGE:777439}	777439 IMAGE Consortium	AA276605

8780 H	4 0219A1GI	Mus musculus uterine mRNA, complete cds {IMAGE:762425}	762425 IMAGE Consortium	AA274119
8781 H	6 0219A1GI	ESTs {IMAGE:762532}	762532 IMAGE Consortium	AA271468
8782 H	8 0219A1GI	Mus musculus EXO70 protein (Exo70) mRNA, complete cds {IMAGE:762591}	762591 IMAGE Consortium	AA271561
8783 H	10 0219A1GI	ESTs {IMAGE:763232}	763232 IMAGE Consortium	AA271984
8784 H	12 0219A1GI	ESTs, Highly similar to U2 SMALL NUCLEAR RIBONUCLEOPROTEIN A' [Homo sapiens] {IMAGE:763635}	763635 IMAGE Consortium	AA285965
8785 B	2 021GA1GJ	ESTs, Moderately similar to rA9 [R.norvegicus] {IMAGE:749259}	749259 IMAGE Consortium	AA290444
8786 B	4 021GA1GJ	Chemokine (C-C) receptor 2 {IMAGE:749361}	749361 IMAGE Consortium	AA289476
8787 B	6 021GA1GJ	(2'-5')OLIGOADENYLATE SYNTHETASE 1 {IMAGE:752301}	752301 IMAGE Consortium	AA290052
8788 B	8 021GA1GJ	IMAGE EST {IMAGE:761573}	761573 IMAGE Consortium	AA272334
8789 B	10 021GA1GJ	ESTs {IMAGE:750129}	750129 IMAGE Consortium	AA396418
8790 B	12 021GA1GJ	ESTs {IMAGE:750360}	750360 IMAGE Consortium	AA396734
8791 D	2 021GA1GJ	Ecotropic viral integration site 2 {IMAGE:749410}	749410 IMAGE Consortium	AA289559
8792 D	4 021GA1GJ	ESTs {IMAGE:752482}	752482 IMAGE Consortium	AA290355
8793 D	6 021GA1GJ	ESTs {IMAGE:761622}	761622 IMAGE Consortium	AA272372
8794 D	8 021GA1GJ	ESTs {IMAGE:749890}	749890 IMAGE Consortium	AA389155
8795 D	10 021GA1GJ	IMAGE EST {IMAGE:762422}	762422 IMAGE Consortium	AA274228
8796 D	12 021GA1GJ	ESTs, Moderately similar to hypertension-induced protein S-A [R.norvegicus] {IMAGE:748351}	748351 IMAGE Consortium	AA274663
8797 F	2 021GA1GJ	ESTs {IMAGE:749775}	749775 IMAGE Consortium	AA396291
8798 F	4 021GA1GJ	IMAGE EST {IMAGE:761702}	761702 IMAGE Consortium	AA272167
8799 F	6 021GA1GJ	IMAGE EST {IMAGE:762339}	762339 IMAGE Consortium	AA273964
8800 F	8 021GA1GJ	ESTs, Highly similar to PROBABLE PROTEIN DISULFIDE ISOMERASE P5 PRECURSOR [Mesocricetus auratus] {IMAGE:762159}	762159 IMAGE Consortium	AA277271
8801 F	10 021GA1GJ	ESTs {IMAGE:750475}	750475 IMAGE Consortium	AA472890
8802 F	12 021GA1GJ	ESTs {IMAGE:750876}	750876 IMAGE Consortium	AA290205
8803 H	2 021GA1GJ	ESTs, Weakly similar to C05D9.6 gene product [C.elegans] {IMAGE:752343}	752343 IMAGE Consortium	AA290001
8804 H	4 021GA1GJ	IMAGE EST {IMAGE:752183}	752183 IMAGE Consortium	AA289070
8805 H	6 021GA1GJ	IMAGE EST {IMAGE:761718}	761718 IMAGE Consortium	AA272023
8806 H	8 021GA1GJ	ESTs {IMAGE:749844}	749844 IMAGE Consortium	AA389143
8807 H	10 021GA1GJ	IMAGE EST {IMAGE:762256}	762256 IMAGE Consortium	AA277407
8808 H	12 021GA1GJ	IMAGE EST {IMAGE:762191}	762191 IMAGE Consortium	AA277314
8809 B	2 021NA1GK	ESTs, Weakly similar to ORF YGL231c [S.cerevisiae] {IMAGE:777257}	777257 IMAGE Consortium	AA276947
8810 B	4 021NA1GK	IMAGE EST {IMAGE:762868}	762868 IMAGE Consortium	AA271792
8811 B	6 021NA1GK	IMAGE EST {IMAGE:766446}	766446 IMAGE Consortium	AA278130
8812 B	8 021NA1GK	Myosin heavy chain, cardiac muscle, adult {IMAGE:763396}	763396 IMAGE Consortium	AA272186
8813 B	10 021NA1GK	IMAGE EST {IMAGE:763526}	763526 IMAGE Consortium	AA285765
8814 B	12 021NA1GK	ESTs, Moderately similar to N-ACETYLLACTOSAMINE	763319 IMAGE Consortium	AA285646

		SYNTHASE [Homo sapiens] {IMAGE:763319}		
8815 D	2 021NA1GK	T-complex testis expressed 1 {IMAGE:762306}	762306 IMAGE Consortium	AA277421
8816 D	4 021NA1GK	Mouse mRNA for AF1q, complete cds {IMAGE:762773}	762773 IMAGE Consortium	AA271708
8817 D	6 021NA1GK	Receptor interacting protein {IMAGE:763021}	763021 IMAGE Consortium	AA271576
8818 D	8 021NA1GK	IMAGE EST {IMAGE:763540}	763540 IMAGE Consortium	AA285771
8819 D	10 021NA1GK	IMAGE EST {IMAGE:763885}	763885 IMAGE Consortium	AA285896
8820 D	12 021NA1GK	Mus musculus mRNA for putative tripeptidyl peptidase I {IMAGE:776048}	776048 IMAGE Consortium	AA276023
8821 F	2 021NA1GK	Cytochrome P450, 2e1, ethanol inducible {IMAGE:762788}	762788 IMAGE Consortium	AA271588
8822 F	4 021NA1GK	ESTs {IMAGE:763263}	763263 IMAGE Consortium	AA271950
8823 F	6 021NA1GK	ESTs {IMAGE:763575}	763575 IMAGE Consortium	AA285930
8824 F	8 021NA1GK	ESTs, Weakly similar to Similarity with yeast gene L3502.1 [C.elegans] {IMAGE:776138}	776138 IMAGE Consortium	AA276168
8825 F	10 021NA1GK	IMAGE EST {IMAGE:764643}	764643 IMAGE Consortium	AA273907
8826 F	12 021NA1GK	ESTs {IMAGE:765100}	765100 IMAGE Consortium	AA273893
8827 H	2 021NA1GK	IMAGE EST {IMAGE:775841}	775841 IMAGE Consortium	AA276064
8828 H	4 021NA1GK	Mus musculus kruppel-related zinc finger protein (Emzf1) mRNA, complete cds {IMAGE:764361}	764361 IMAGE Consortium	AA277447
8829 H	6 021NA1GK	ESTs {IMAGE:764681}	764681 IMAGE Consortium	AA274037
8830 H	8 021NA1GK	IMAGE EST {IMAGE:777053}	777053 IMAGE Consortium	AA276251
8831 H	10 021NA1GK	IMAGE EST {IMAGE:776876}	776876 IMAGE Consortium	AA273553
8832 H	12 021NA1GK	ESTs {IMAGE:777316}	777316 IMAGE Consortium	AA276904
8833 B	2 021UA1GL	IMAGE EST {IMAGE:775722}	775722 IMAGE Consortium	AA275684
8834 B	4 021UA1GL	IMAGE EST {IMAGE:776075}	776075 IMAGE Consortium	AA276035
8835 B	6 021UA1GL	ESTs, Highly similar to Zis [R.norvegicus] {IMAGE:764790}	764790 IMAGE Consortium	AA274193
8836 B	8 021UA1GL	ESTs, Weakly similar to No definition line found [C.elegans] {IMAGE:776467}	776467 IMAGE Consortium	AA275754
8837 B	10 021UA1GL	IMAGE EST {IMAGE:776792}	776792 IMAGE Consortium	AA419634
8838 B	12 021UA1GL	IMAGE EST {IMAGE:776807}	776807 IMAGE Consortium	AA419651
8839 D	2 021UA1GL	IMAGE EST {IMAGE:763553}	763553 IMAGE Consortium	AA285513
8840 D	4 021UA1GL	Protein phosphatase 1, catalytic subunit, beta isoform {IMAGE:764170}	764170 IMAGE Consortium	AA285841
8841 D	6 021UA1GL	ESTs {IMAGE:764398}	764398 IMAGE Consortium	AA277176
8842 D	8 021UA1GL	ESTs {IMAGE:764625}	764625 IMAGE Consortium	AA273981
8843 D	10 021UA1GL	IMAGE EST {IMAGE:764803}	764803 IMAGE Consortium	AA273684
8844 D	12 021UA1GL	ESTs, Weakly similar to HYPOTHETICAL PROTEIN F44G4.1 IN CHROMOSOME II [C.elegans] {IMAGE:765177}	765177 IMAGE Consortium	AA273955
8845 F	2 021UA1GL	Renin 1 structural {IMAGE:776250}	776250 IMAGE Consortium	AA276457
8846 F	4 021UA1GL	Mus musculus Ras-binding protein SUR-8 mRNA, complete cds {IMAGE:776500}	776500 IMAGE Consortium	AA286039
8847 F	6 021UA1GL	IMAGE EST {IMAGE:776564}	776564 IMAGE Consortium	AA275889
8848 F	8 021UA1GL	IMAGE EST {IMAGE:765140}	765140 IMAGE Consortium	AA273262
8849 F	10 021UA1GL	IMAGE EST {IMAGE:777236}	777236 IMAGE Consortium	AA276891

8850 F	12 021UA1GL	IMAGE EST {IMAGE:808315}	808315 IMAGE Consortium	AA433039
8851 H	2 021UA1GL	ESTs {IMAGE:820254}	820254 IMAGE Consortium	AA423133
8852 H	4 021UA1GL	IMAGE EST {IMAGE:820582}	820582 IMAGE Consortium	AA423488
8853 H	6 021UA1GL	IMAGE EST {IMAGE:831991}	831991 IMAGE Consortium	AA444525
8854 H	8 021UA1GL	ESTs {IMAGE:777308}	777308 IMAGE Consortium	AA276960
8855 H	10 021UA1GL	Cytochrome P450, 2e1, ethanol inducible {IMAGE:777580}	777580 IMAGE Consortium	AA276835
8856 H	12 021UA1GL	Complement component factor h {IMAGE:777640}	777640 IMAGE Consortium	AA276818
8857 B	2 0211A1GM	M.musculus adipocyte-specific mRNA, partial cds {IMAGE:831665}	831665 IMAGE Consortium	AA466094
8858 B	4 0211A1GM	Protein kinase C, delta {IMAGE:777583}	777583 IMAGE Consortium	AA276844
8859 B	6 0211A1GM	IMAGE EST {IMAGE:777516}	777516 IMAGE Consortium	AA276640
8860 B	8 0211A1GM	ESTs {IMAGE:817924}	817924 IMAGE Consortium	AA466508
8861 B	10 0211A1GM	ESTs, Highly similar to GMP REDUCTASE [Homo sapiens] {IMAGE:818880}	818880 IMAGE Consortium	AA434881
8862 B	12 0211A1GM	ESTs {IMAGE:819125}	819125 IMAGE Consortium	AA426768
8863 D	2 0211A1GM	Mus musculus mRNA for beta-1,4-galactosyltransferase II, complete cds {IMAGE:808719}	808719 IMAGE Consortium	AA467331
8864 D	4 0211A1GM	IMAGE EST {IMAGE:817954}	817954 IMAGE Consortium	AA466530
8865 D	6 0211A1GM	IMAGE EST {IMAGE:790705}	790705 IMAGE Consortium	AA388563
8866 D	8 0211A1GM	IMAGE EST {IMAGE:818653}	818653 IMAGE Consortium	AA466649
8867 D	10 0211A1GM	ESTs {IMAGE:820186}	820186 IMAGE Consortium	AA423210
8868 D	12 0211A1GM	ESTs, Weakly similar to integrase [M.musculus] {IMAGE:820336}	820336 IMAGE Consortium	AA423205
8869 F	2 0211A1GM	ESTs {IMAGE:809031}	809031 IMAGE Consortium	AA466039
8870 F	4 0211A1GM	ESTs {IMAGE:790857}	790857 IMAGE Consortium	AA388512
8871 F	6 0211A1GM	Mus musculus arginase II mRNA, complete cds {IMAGE:820157}	820157 IMAGE Consortium	AA423144
8872 F	8 0211A1GM	ESTs, Weakly similar to ORF YOR173w [S.cerevisiae] {IMAGE:803399}	803399 IMAGE Consortium	AA473067
8873 F	10 0211A1GM	ESTs {IMAGE:806753}	806753 IMAGE Consortium	AA404124
8874 F	12 0211A1GM	ESTs, Moderately similar to unknown protein CIT987SK_2A8_1 [H.sapiens] {IMAGE:831732}	831732 IMAGE Consortium	AA444926
8875 H	2 0211A1GM	ESTs, Highly similar to PHOSPHORYLASE B KINASE ALPHA REGULATORY CHAIN, LIVER ISOFORM [Oryctolagus cuniculus] {IMAGE:820277}	820277 IMAGE Consortium	AA423239
8876 H	4 0211A1GM	IMAGE EST {IMAGE:806777}	806777 IMAGE Consortium	AA403810
8877 H	6 0211A1GM	ESTs {IMAGE:831799}	831799 IMAGE Consortium	AA444640
8878 H	8 0211A1GM	ESTs {IMAGE:777795}	777795 IMAGE Consortium	AA286382
8879 H	10 0211A1GM	IMAGE EST {IMAGE:777584}	777584 IMAGE Consortium	AA276837
8880 H	12 0211A1GM	ESTs {IMAGE:777543}	777543 IMAGE Consortium	AA276754
8881 B	2 0218A1GN	IMAGE EST {IMAGE:820535}	820535 IMAGE Consortium	AA423392
8882 B	4 0218A1GN	ESTs, Weakly similar to 39E1.i [D.melanogaster] {IMAGE:807957}	807957 IMAGE Consortium	AA432925

8883 B	6 0218A1GN	ESTs {IMAGE:777633}	777633 IMAGE Consortium	AA276732
8884 B	8 0218A1GN	IMAGE EST {IMAGE:777857}	777857 IMAGE Consortium	AA286518
8885 B	10 0218A1GN	ESTs {IMAGE:777814}	777814 IMAGE Consortium	AA286504
8886 B	12 0218A1GN	IMAGE EST {IMAGE:818420}	818420 IMAGE Consortium	AA434854
8887 D	2 0218A1GN	IMAGE EST {IMAGE:777605}	777605 IMAGE Consortium	AA276859
8888 D	4 0218A1GN	IMAGE EST {IMAGE:777666}	777666 IMAGE Consortium	AA276825
8889 D	6 0218A1GN	ESTs {IMAGE:809062}	809062 IMAGE Consortium	AA466077
8890 D	8 0218A1GN	IMAGE EST {IMAGE:818790}	818790 IMAGE Consortium	AA467382
8891 D	10 0218A1GN	ATPase, Na <sup>+</sup> /K <sup>+</sup> transporting, beta 2 polypeptide {IMAGE:803337}	803337 IMAGE Consortium	AA473010
8892 D	12 0218A1GN	IMAGE EST {IMAGE:820316}	820316 IMAGE Consortium	AA423248
8893 F	2 0218A1GN	ESTs {IMAGE:818253}	818253 IMAGE Consortium	AA435375
8894 F	4 0218A1GN	Peroxisomal sarcosine oxidase {IMAGE:790721}	790721 IMAGE Consortium	AA388435
8895 F	6 0218A1GN	IMAGE EST {IMAGE:820417}	820417 IMAGE Consortium	AA423317
8896 F	8 0218A1GN	ESTs, Weakly similar to K08F11.3 [C.elegans] {IMAGE:820430}	820430 IMAGE Consortium	AA423312
8897 F	10 0218A1GN	ESTs, Highly similar to HYPOTHETICAL 138.8 KD PROTEIN C11D3.14C IN CHROMOSOME I [Schizosaccharomyces pombe] {IMAGE:831818}	831818 IMAGE Consortium	AA444650
8898 F	12 0218A1GN	ESTs {IMAGE:777238}	777238 IMAGE Consortium	AA276892
8899 H	2 0218A1GN	Promyelocytic leukemia {IMAGE:820560}	820560 IMAGE Consortium	AA423337
8900 H	4 0218A1GN	IMAGE EST {IMAGE:831676}	831676 IMAGE Consortium	AA467020
8901 H	6 0218A1GN	ESTs {IMAGE:831951}	831951 IMAGE Consortium	AA444485
8902 H	8 0218A1GN	ESTs {IMAGE:808377}	808377 IMAGE Consortium	AA466705
8903 H	10 0218A1GN	IMAGE EST {IMAGE:808922}	808922 IMAGE Consortium	AA465808
8904 H	12 0218A1GN	Interleukin 4 receptor, alpha {IMAGE:777811}	777811 IMAGE Consortium	AA286506
8905 B	2 021FA1GO			
8906 B	4 021FA1GO			
8907 B	6 021FA1GO			
8908 B	8 021FA1GO			
8909 B	10 021FA1GO			
8910 B	12 021FA1GO			
8911 D	2 021FA1GO			
8912 D	4 021FA1GO			
8913 D	6 021FA1GO			
8914 D	8 021FA1GO			
8915 D	10 021FA1GO			
8916 D	12 021FA1GO			
8917 F	2 021FA1GO			
8918 F	4 021FA1GO			
8919 F	6 021FA1GO			
8920 F	8 021FA1GO			
8921 F	10 021FA1GO			

8922 F	12 021FA1GO			
8923 H	2 021FA1GO			
8924 H	4 021FA1GO			
8925 H	6 021FA1GO			
8926 H	8 021FA1GO			
8927 H	10 021FA1GO			
8928 H	12 021FA1GO			
8929 B	2 021MA1GP	ESTs, Moderately similar to RAS-RELATED PROTEIN RABB [Dictyostelium discoideum] {IMAGE:522605}	522605 IMAGE Consortium	AA068057
8930 B	4 021MA1GP	ESTs {IMAGE:523241}	523241 IMAGE Consortium	AA067119
8931 B	6 021MA1GP	IMAGE EST {IMAGE:523686}	523686 IMAGE Consortium	AA067149
8932 B	8 021MA1GP	Mus musculus RRM RNA binding protein GRY-RBP mRNA, complete cds {IMAGE:524304}	524304 IMAGE Consortium	AA066208
8933 B	10 021MA1GP	IMAGE EST {IMAGE:524253}	524253 IMAGE Consortium	AA067887
8934 B	12 021MA1GP	ESTs {IMAGE:535645}	535645 IMAGE Consortium	AA108103
8935 D	2 021MA1GP	ESTs, Highly similar to CYSTATHIONINE GAMMA-LYASE [Homo sapiens] {IMAGE:525119}	525119 IMAGE Consortium	AA096870
8936 D	4 021MA1GP	Mus musculus protein kinase C inhibitor (mPKCI) mRNA, complete cds {IMAGE:533117}	533117 IMAGE Consortium	AA068901
8937 D	6 021MA1GP	ESTs {IMAGE:533415}	533415 IMAGE Consortium	AA105717
8938 D	8 021MA1GP	IMAGE EST {IMAGE:534003}	534003 IMAGE Consortium	AA071831
8939 D	10 021MA1GP	Mus musculus hemin-sensitive initiation factor 2 alpha kinase mRNA, complete cds {IMAGE:534096}	534096 IMAGE Consortium	AA072620
8940 D	12 021MA1GP	ESTs, Highly similar to PRE-MRNA SPLICING FACTOR PRP6 [Saccharomyces cerevisiae] {IMAGE:535535}	535535 IMAGE Consortium	AA087611
8941 F	2 021MA1GP	Cyclin G {IMAGE:523713}	523713 IMAGE Consortium	AA067318
8942 F	4 021MA1GP	ESTs, Moderately similar to nuclear pore complex glycoprotein p62 [M.musculus] {IMAGE:524367}	524367 IMAGE Consortium	AA065748
8943 F	6 021MA1GP	Protein kinase, DNA activated, catalytic polypeptide {IMAGE:524350}	524350 IMAGE Consortium	AA067927
8944 F	8 021MA1GP	ESTs {IMAGE:535577}	535577 IMAGE Consortium	AA110853
8945 F	10 021MA1GP	IMAGE EST {IMAGE:536248}	536248 IMAGE Consortium	AA073714
8946 F	12 021MA1GP	ESTs, Highly similar to HYPOTHETICAL 36.7 KD PROTEIN AH6.2 IN CHROMOSOME II [Caenorhabditis elegans] {IMAGE:532139}	532139 IMAGE Consortium	AA068366
8947 H	2 021MA1GP	ESTs, Weakly similar to L-ASPARAGINASE [Lupinus arboreus] {IMAGE:550910}	550910 IMAGE Consortium	AA097901
8948 H	4 021MA1GP	ESTs {IMAGE:551164}	551164 IMAGE Consortium	AA080385
8949 H	6 021MA1GP	ESTs, Highly similar to CHROMOSOME REGION MAINTENANCE PROTEIN 1 [Schizosaccharomyces pombe] {IMAGE:533299}	533299 IMAGE Consortium	AA105546
8950 H	8 021MA1GP	IMAGE EST {IMAGE:534269}	534269 IMAGE Consortium	AA072765
8951 H	10 021MA1GP	Lamin A {IMAGE:524119}	524119 IMAGE Consortium	AA066180
8952 H	12 021MA1GP	ESTs {IMAGE:524631}	524631 IMAGE Consortium	AA068125



8953 B	2 021TA1GQ	IMAGE EST {IMAGE:525168}	525168 IMAGE Consortium	AA097025
8954 B	4 021TA1GQ	ESTs, Weakly similar to ANKYRIN, BRAIN VARIANT 2 [Homo sapiens] {IMAGE:536633}	536633 IMAGE Consortium	AA080159
8955 B	6 021TA1GQ	Mus musculus steroid receptor coactivator-1 (mSRC-1) mRNA, complete cds {IMAGE:523588}	523588 IMAGE Consortium	AA097341
8956 B	8 021TA1GQ	Mus musculus putative endo/exonuclease MmMre11b (MmMRE11) mRNA, alternative splicing product, complete cds {IMAGE:524361}	524361 IMAGE Consortium	AA065745
8957 B	10 021TA1GQ	ESTs {IMAGE:535432}	535432 IMAGE Consortium	AA087579
8958 B	12 021TA1GQ	ESTs {IMAGE:535602}	535602 IMAGE Consortium	AA110858
8959 D	2 021TA1GQ	IMAGE EST {IMAGE:536419}	536419 IMAGE Consortium	AA073941
8960 D	4 021TA1GQ	IMAGE EST {IMAGE:522596}	522596 IMAGE Consortium	AA068008
8961 D	6 021TA1GQ	ESTs {IMAGE:523046}	523046 IMAGE Consortium	AA066439
8962 D	8 021TA1GQ	ESTs, Highly similar to HYPOTHETICAL 64.3 KD GTP-BINDING PROTEIN C02F5.3 IN CHROMOSOME III [Caenorhabditis elegans] {IMAGE:523945}	523945 IMAGE Consortium	AA071987
8963 D	10 021TA1GQ	IMAGE EST {IMAGE:523726}	523726 IMAGE Consortium	AA067346
8964 D	12 021TA1GQ	Mus musculus ajuba (Ajuba) mRNA, complete cds {IMAGE:535665}	535665 IMAGE Consortium	AA106932
8965 F	2 021TA1GQ	ESTs, Weakly similar to protein co-factor [M.musculus] {IMAGE:535340}	535340 IMAGE Consortium	AA087796
8966 F	4 021TA1GQ	ESTs, Moderately similar to biphenyl hydrolase-related protein [H.sapiens] {IMAGE:524682}	524682 IMAGE Consortium	AA068497
8967 F	6 021TA1GQ	IMAGE EST {IMAGE:536574}	536574 IMAGE Consortium	AA073843
8968 F	8 021TA1GQ	Mus musculus C6-I proteasome chain mRNA, complete cds {IMAGE:550898}	550898 IMAGE Consortium	AA098608
8969 F	10 021TA1GQ	ESTs, Highly similar to 60S RIBOSOMAL PROTEIN L15 [Rattus norvegicus] {IMAGE:532770}	532770 IMAGE Consortium	AA068842
8970 F	12 021TA1GQ	ESTs, Highly similar to HYPOTHETICAL 27.4 KD PROTEIN IN MER2-CPR7 INTERGENIC REGION [Saccharomyces cerevisiae] {IMAGE:427351}	427351 IMAGE Consortium	AA003173
8971 H	2 021TA1GQ	Renin 1 structural {IMAGE:661753}	661753 IMAGE Consortium	AA207872
8972 H	4 021TA1GQ	ESTs {IMAGE:652273}	652273 IMAGE Consortium	AA214955
8973 H	6 021TA1GQ	IMAGE EST {IMAGE:917776}	917776 IMAGE Consortium	AA497742
8974 H	8 021TA1GQ	MITOCHONDRIAL MATRIX PROTEIN P1 PRECURSOR {IMAGE:633625}	633625 IMAGE Consortium	AA184322
8975 H	10 021TA1GQ	IMAGE EST {IMAGE:651485}	651485 IMAGE Consortium	AA215069
8976 H	12 021TA1GQ	Leukemia inhibitory factor receptor {IMAGE:662150}	662150 IMAGE Consortium	AA207338
8977 B	2 0210A1GR	Mus musculus C6-I proteasome chain mRNA, complete cds {IMAGE:651274}	651274 IMAGE Consortium	AA214752
8978 B	4 0210A1GR	SODIUM/POTASSIUM-TRANSPORTING ATPASE GAMMA CHAIN {IMAGE:662933}	662933 IMAGE Consortium	AA208297
8979 B	6 0210A1GR	Mus musculus G-protein-like LRG-47 mRNA, complete cds {IMAGE:651927}	651927 IMAGE Consortium	AA214784

8980 B	8 0210A1GR	ESTs {IMAGE:656383}	656383 IMAGE Consortium	AA242517
8981 B	10 0210A1GR	Mus musculus mRNA for rabkinesin-6 {IMAGE:633209}	633209 IMAGE Consortium	AA177197
8982 B	12 0210A1GR	ESTs, Highly similar to ETn insert [M.musculus] {IMAGE:633484}	633484 IMAGE Consortium	AA184421
8983 D	2 0210A1GR	IMAGE EST {IMAGE:656113}	656113 IMAGE Consortium	AA239561
8984 D	4 0210A1GR	IMAGE EST {IMAGE:656145}	656145 IMAGE Consortium	AA241114
8985 D	6 0210A1GR	ESTs {IMAGE:651184}	651184 IMAGE Consortium	AA213286
8986 D	8 0210A1GR	Trans-acting transcription factor 1 {IMAGE:651823}	651823 IMAGE Consortium	AA212645
8987 D	10 0210A1GR	IMAGE EST {IMAGE:652000}	652000 IMAGE Consortium	AA214870
8988 D	12 0210A1GR	IMAGE EST {IMAGE:917949}	917949 IMAGE Consortium	AA499756
8989 F	2 0210A1GR	ESTs, Weakly similar to COATOMER GAMMA SUBUNIT [Saccharomyces cerevisiae] {IMAGE:656147}	656147 IMAGE Consortium	AA241132
8990 F	4 0210A1GR	IMAGE EST {IMAGE:633261}	633261 IMAGE Consortium	AA177260
8991 F	6 0210A1GR	IMAGE EST {IMAGE:651338}	651338 IMAGE Consortium	AA214947
8992 F	8 0210A1GR	ESTs, Highly similar to nuclear protein Skip [H.sapiens] {IMAGE:651683}	651683 IMAGE Consortium	AA212339
8993 F	10 0210A1GR	IMAGE EST {IMAGE:651622}	651622 IMAGE Consortium	AA212170
8994 F	12 0210A1GR	Upstream transcription factor 2 {IMAGE:732918}	732918 IMAGE Consortium	AA286166
8995 H	2 0210A1GR			
8996 H	4 0210A1GR			
8997 H	6 0210A1GR			
8998 H	8 0210A1GR			
8999 H	10 0210A1GR			
9000 H	12 0210A1GR			
9001 B	2 021Q9974	Control: Ratio 1:3 (Cy3:Cy5)	Ratio_A	Controls
9002 B	4 021Q9974	Control: Ratio 3:1 (Cy3:Cy5)	Ratio_B	Controls
9003 B	6 021Q9974	Control: Ratio 1:10 (Cy3:Cy5)	Ratio_C	Controls
9004 B	8 021Q9974	Control: Ratio 10:1 (Cy3:Cy5)	Ratio_D	Controls
9005 B	10 021Q9974	Control: Ratio 1:25 (Cy3:Cy5)	Ratio_E	Controls
9006 B	12 021Q9974	Control: Ratio 25:1 (Cy3:Cy5)	Ratio_F	Controls
9007 D	2 021Q9974	Control: Sensitivity 2pg	Sensitivity_A	Controls
9008 D	4 021Q9974	Control: Sensitivity 20pg	Sensitivity_B	Controls
9009 D	6 021Q9974	Control: Sensitivity 200pg	Sensitivity_C	Controls
9010 D	8 021Q9974	Control: Sensitivity 2000pg	Sensitivity_D	Controls
9011 D	10 021Q9974	Control: Buffer Only	Buffer_Only_Control	Controls
9012 D	12 021Q9974	Control: Buffer Only	Buffer_Only_Control	Controls
9013 F	2 021Q9974	Control: Internal	Internal_Control_A	Controls
9014 F	4 021Q9974	Control: Buffer Only	Buffer_Only_Control	Controls
9015 F	6 021Q9974	Control: Internal	Internal_Control_B	Controls
9016 F	8 021Q9974	Control: Buffer Only	Buffer_Only_Control	Controls
9017 F	10 021Q9974	Control: Internal	Internal_Control_C	Controls
9018 F	12 021Q9974	Control: Internal	Internal_Control_D	Controls
9019 H	2 021Q9974	Control: Complex Target (Mus musculus)	RNA_Quality_MM	Controls

9020 H	4 021Q9974	Control: Complex Target (Mus musculus)	RNA_Quality_MM	Controls
9021 H	6 021Q9974	Control: Complex Target (Mus musculus)	RNA_Quality_MM	Controls
9022 H	8 021Q9974	Control: Housekeeping (Alpha Tubulin)	Housekeeping_B	Controls
9023 H	10 021Q9974	Control: Housekeeping (Ribosomal S9)	Housekeeping_A	Controls
9024 H	12 021Q9974	Control: Housekeeping (23kD HBP)	Housekeeping_C	Controls

**Gem Mouse 2:** 10176 elements of 10200 total possible elements (some were excluded by the company due to technical considerations)

Location	PlateRow	PlateCol	PlatelD	GeneName	CloneID	CloneSource	AccessionNum
	1	A	1	0217AJ7T	Control: Ratio 1:3 (Cy3:Cy5)	Ratio_A	Controls
	2	A	3	0217AJ7T	Control: Ratio 3:1 (Cy3:Cy5)	Ratio_B	Controls
	3	A	5	0217AJ7T	Control: Ratio 1:10 (Cy3:Cy5)	Ratio_C	Controls
	4	A	7	0217AJ7T	Control: Ratio 10:1 (Cy3:Cy5)	Ratio_D	Controls
	5	A	9	0217AJ7T	Control: Ratio 1:25 (Cy3:Cy5)	Ratio_E	Controls
	6	A	11	0217AJ7T	Control: Ratio 25:1 (Cy3:Cy5)	Ratio_F	Controls
	7	C	1	0217AJ7T	Control: Sensitivity 2pg	Sensitivity_A	Controls
	8	C	3	0217AJ7T	Control: Sensitivity 20pg	Sensitivity_B	Controls
	9	C	5	0217AJ7T	Control: Sensitivity 200pg	Sensitivity_C	Controls
	10	C	7	0217AJ7T	Control: Sensitivity 2000pg	Sensitivity_D	Controls
	11	C	9	0217AJ7T	Control: Buffer Only	Buffer_Only_Control	Controls
	12	C	11	0217AJ7T	Control: Buffer Only	Buffer_Only_Control	Controls
	13	E	1	0217AJ7T	Control: Internal	Internal_Control_A	Controls
	14	E	3	0217AJ7T	Control: Buffer Only	Buffer_Only_Control	Controls
	15	E	5	0217AJ7T	Control: Internal	Internal_Control_B	Controls
	16	E	7	0217AJ7T	Control: Buffer Only	Buffer_Only_Control	Controls
	17	E	9	0217AJ7T	Control: Internal	Internal_Control_C	Controls
	18	E	11	0217AJ7T	Control: Internal	Internal_Control_D	Controls
	19	G	1	0217AJ7T	Control: Complex Target (Mus musculus)	RNA_Quality_MM	Controls
	20	G	3	0217AJ7T	Control: Complex Target (Mus musculus)	RNA_Quality_MM	Controls
	21	G	5	0217AJ7T	Control: Complex Target (Mus musculus)	RNA_Quality_MM	Controls
	22	G	7	0217AJ7T	Control: Housekeeping (Alpha Tubulin)	Housekeeping_B	Controls
	23	G	9	0217AJ7T	Control: Housekeeping (Ribosomal S9)	Housekeeping_A	Controls
	24	G	11	0217AJ7T	Control: Housekeeping (23kD HBP)	Housekeeping_C	Controls
	25	A	1	021JAMMK	peroxisome proliferator activated receptor alpha	313176	IMAGE Consortium W10495
	26	A	3	021JAMMK	ESTs, Weakly similar to putative poly(ADP-ribosyl) transferase [H.sapiens]	314211	IMAGE Consortium W09784
	27	A	5	021JAMMK	ESTs, Weakly similar to DEOXYRIBOSE-PHOSPHATE ALDOLASE [Escherichia coli]	315392	IMAGE Consortium AI327448
	28	A	7	021JAMMK	Notch gene homolog 4, (Drosophila)	317393	IMAGE Consortium W34046
	29	A	9	021JAMMK	caveolin 3	427435	IMAGE Consortium AA002425
	30	A	11	021JAMMK	ESTs, Weakly similar to acetyl LDL receptor [H.sapiens]	421505	IMAGE Consortium W99208
	31	C	1	021JAMMK	serine/arginine-rich protein specific kinase 2	314443	IMAGE Consortium AI605657
	32	C	3	021JAMMK	mast cell protease 6	314268	IMAGE Consortium W10007
	33	C	5	021JAMMK	cytochrome P450, 2b19	316138	IMAGE Consortium W12527
	34	C	7	021JAMMK	tyrosine kinase TYK2	316704	IMAGE Consortium W11201
	35	C	9	021JAMMK	distal-less homeobox 5	427483	IMAGE Consortium AI894025
	36	C	11	021JAMMK	ESTs	423080	IMAGE Consortium W98004
	37	E	1	021JAMMK	ESTs	303426	IMAGE Consortium AI627139

38 E	3 021JAMMK	ESTs, Highly similar to KIAA0456 protein [H.sapiens]	313586 IMAGE Consortium	AI605614
39 E	5 021JAMMK	ESTs, Highly similar to FKBP-RAPAMYCIN ASSOCIATED PROTEIN [Homo sapiens]	315428 IMAGE Consortium	AI327449
40 E	7 021JAMMK	ESTs	316867 IMAGE Consortium	AI607010
41 E	9 021JAMMK	adaptor protein complex AP-1, beta 1 subunit	437154 IMAGE Consortium	AA003413
42 E	11 021JAMMK	ESTs	421847 IMAGE Consortium	W91450
43 G	1 021JAMMK	ESTs	303562 IMAGE Consortium	AI627156
44 G	3 021JAMMK	ESTs	315026 IMAGE Consortium	W12565
45 G	5 021JAMMK	beta-site APP cleaving enzyme	314741 IMAGE Consortium	AI605684
46 G	7 021JAMMK	intercellular adhesion molecule 5, telencephalin	422226 IMAGE Consortium	AI893998
47 G	9 021JAMMK	TEA domain family member 3	437492 IMAGE Consortium	AI894030
48 G	11 021JAMMK	5-azacytidine induced gene 1	424580 IMAGE Consortium	W98281
49 A	1 021QAMML	DNA segment, Chr 11, KL Mohlke 47	539908 IMAGE Consortium	AI550286
50 A	3 021QAMML	ESTs, Highly similar to MELANOMA-ASSOCIATED ANTIGEN XP [Homo sapiens]	540466 IMAGE Consortium	AI892529
51 A	5 021QAMML	ESTs	539514 IMAGE Consortium	AA153180
52 A	7 021QAMML	ESTs	539104 IMAGE Consortium	AA117356
53 A	9 021QAMML	ESTs	539308 IMAGE Consortium	AA116953
54 A	11 021QAMML	ESTs	538839 IMAGE Consortium	AA123098
55 C	1 021QAMML	ESTs, Weakly similar to CG5181 gene product [D.melanogaster]	539995 IMAGE Consortium	AA122644
56 C	3 021QAMML	EST X83327	540488 IMAGE Consortium	AA162627
57 C	5 021QAMML	lactate dehydrogenase 1, A chain	539536 IMAGE Consortium	AI325819
58 C	7 021QAMML	ESTs	539119 IMAGE Consortium	AI550217
59 C	9 021QAMML	ESTs	539320 IMAGE Consortium	AA117509
60 C	11 021QAMML	DNA segment, Chr 11, ERATO Doi 603, expressed	539576 IMAGE Consortium	AA123079
61 E	1 021QAMML	chondroitin sulfate proteoglycan 6	540067 IMAGE Consortium	AI325825
62 E	3 021QAMML	ESTs	540518 IMAGE Consortium	AI550569
63 E	5 021QAMML	COP9 (constitutive photomorphogenic), subunit 5 (Arabidopsis)	539629 IMAGE Consortium	AA123224
64 E	7 021QAMML	ESTs	539141 IMAGE Consortium	AA116494
65 E	9 021QAMML	ELAV (embryonic lethal, abnormal vision, Drosophila)-like 1 (Hu antigen R)	539337 IMAGE Consortium	AA117058
66 E	11 021QAMML	ESTs, Moderately similar to AF133124_1 transcription factor IIIC63 [H.sapiens]	538871 IMAGE Consortium	AA120184
67 G	1 021QAMML	DNA primase, p49 subunit	540239 IMAGE Consortium	AI324469
68 G	3 021QAMML	ESTs, Moderately similar to AF182003_1 tyrosyl-DNA phosphodiesterase protein [H.sapiens]	540552 IMAGE Consortium	AA162701
69 G	5 021QAMML	ESTs	539680 IMAGE Consortium	AA117506
70 G	7 021QAMML	ESTs	539174 IMAGE Consortium	AA116742
71 G	9 021QAMML	ESTs	539393 IMAGE Consortium	AI550234
72 G	11 021QAMML	breast cancer 2	539644 IMAGE Consortium	AA123045
73 A	1 021XAMMM	ESTs	541049 IMAGE Consortium	AA120439

74 A	3	021XAMMM	ESTs	540628	IMAGE Consortium	AA162716
75 A	5	021XAMMM				
76 A	7	021XAMMM				
77 A	9	021XAMMM	sialyltransferase 1 (beta-galactoside alpha-2,6-sialyltransferase)	560014	IMAGE Consortium	AA265220
78 A	11	021XAMMM	guanine nucleotide binding protein (G protein), gamma 3 subunit	560312	IMAGE Consortium	AA210241
79 C	1	021XAMMM	ESTs	541140	IMAGE Consortium	AA162693
80 C	3	021XAMMM	ESTs, Moderately similar to ZINC FINGER PROTEIN 32 [Homo sapiens]	540666	IMAGE Consortium	AI892535
81 C	5	021XAMMM	ESTs, Weakly similar to CG5047 gene product [D.melanogaster]	559785	IMAGE Consortium	AA209780
82 C	7	021XAMMM	ESTs	560066	IMAGE Consortium	AA265328
83 C	9	021XAMMM	tropomodulin 3	560109	IMAGE Consortium	AA265361
84 C	11	021XAMMM	Mus musculus ORF1 for hypothetical protein, clone 424981	560318	IMAGE Consortium	AA210242
85 E	1	021XAMMM	ESTs, Moderately similar to AF200348_1 melanoma-associated antigen MG50 [H.sapiens]	541156	IMAGE Consortium	AA162049
86 E	3	021XAMMM	ESTs, Moderately similar to unknown [H.sapiens]	540682	IMAGE Consortium	AI427170
87 E	5	021XAMMM	ESTs, Highly similar to T13057 KIAA0729 protein - human [H.sapiens]	559792	IMAGE Consortium	AA209628
88 E	7	021XAMMM	ESTs	560097	IMAGE Consortium	AA265490
89 E	9	021XAMMM	ESTs, Weakly similar to UB5B_HUMAN UBIQUITIN-CONJUGATING ENZYME E2-17 KD 2 [R.norvegicus]	560157	IMAGE Consortium	AA210502
90 E	11	021XAMMM	ESTs, Weakly similar to A55817 cyclin-dependent kinase p130-PITSLRE - mouse [M.musculus]	560325	IMAGE Consortium	AA210299
91 G	1	021XAMMM	ESTs	541179	IMAGE Consortium	AA162996
92 G	3	021XAMMM	ESTs, Highly similar to HYPOTHETICAL ZINC FINGER PROTEIN ZK686.4 IN CHROMOSOME III [Caenorhabditis elegans]	540714	IMAGE Consortium	AI892537
93 G	5	021XAMMM	A kinase (PRKA) anchor protein 10	540982	IMAGE Consortium	AA120401
94 G	7	021XAMMM	nucleobindin 2	560115	IMAGE Consortium	AA265339
95 G	9	021XAMMM	ESTs, Moderately similar to KIAA1291 protein [H.sapiens]	560175	IMAGE Consortium	AA210448
96 G	11	021XAMMM	serine racemase	560366	IMAGE Consortium	AA210414
97 A	1	0214AMMN	DNA segment, Chr 14, University of California at Los Angeles 3	560431	IMAGE Consortium	AA210481
98 A	3	0214AMMN	peptidylprolyl isomerase C	560901	IMAGE Consortium	AA212126
99 A	5	0214AMMN	procollagen lysine, 2-oxoglutarate 5-dioxygenase 2	822625	IMAGE Consortium	AI549660
100 A	7	0214AMMN	ESTs, Highly similar to A26882 pIL2 hypothetical protein - rat [R.norvegicus]	822906	IMAGE Consortium	AA438128
101 A	9	0214AMMN	ESTs	822414	IMAGE Consortium	AA474127
102 A	11	0214AMMN	ESTs	822507	IMAGE Consortium	AA474088
103 C	1	0214AMMN	DNA methyltransferase 3B	560490	IMAGE Consortium	AA210568
104 C	3	0214AMMN	ESTs	806033	IMAGE Consortium	AA399804

105 C	5 0214AMMN	ESTs	822672 IMAGE Consortium	AI549666
106 C	7 0214AMMN	ESTs, Highly similar to GUANINE NUCLEOTIDE-BINDING PROTEIN G(K), ALPHA SUBUNIT [Rattus norvegicus]	822931 IMAGE Consortium	AA438152
107 C	9 0214AMMN	ESTs	822420 IMAGE Consortium	AI549636
108 C	11 0214AMMN	ESTs	822514 IMAGE Consortium	AI549650
109 E	1 0214AMMN	carbonic anhydrase 2	560505 IMAGE Consortium	AA210505
110 E	3 0214AMMN	ESTs, Highly similar to BN51 PROTEIN [Homo sapiens]	806044 IMAGE Consortium	AA399844
111 E	5 0214AMMN	ESTs	822684 IMAGE Consortium	AA474300
112 E	7 0214AMMN	ESTs	822935 IMAGE Consortium	AI549693
113 E	9 0214AMMN	ESTs, Highly similar to hypothetical protein [H.sapiens]	822425 IMAGE Consortium	AA474138
114 E	11 0214AMMN	Public domain EST	821851 IMAGE Consortium	
115 G	1 0214AMMN	DNA segment, Chr 13, ERATO Doi 328, expressed	560548 IMAGE Consortium	AA212023
116 G	3 0214AMMN	DNA segment, Chr 16, ERATO Doi 61, expressed	806080 IMAGE Consortium	AA399813
117 G	5 0214AMMN	ESTs	822695 IMAGE Consortium	AI549671
118 G	7 0214AMMN	ESTs	822940 IMAGE Consortium	AA438154
119 G	9 0214AMMN	ESTs, Weakly similar to p58 [M.musculus]	822435 IMAGE Consortium	AA474147
120 G	11 0214AMMN	ribonuclease P2	821862 IMAGE Consortium	AA413027
121 A	1 021BAMMO	trophoblast glycoprotein	821342 IMAGE Consortium	AI552351
122 A	3 021BAMMO	ESTs	806373 IMAGE Consortium	AA432755
123 A	5 021BAMMO	rho GTPase activating protein 5	821808 IMAGE Consortium	AI549615
124 A	7 021BAMMO	cyclin A2	865112 IMAGE Consortium	AA469716
125 A	9 021BAMMO	ESTs, Weakly similar to AF220191_1 uncharacterized hypothalamus protein HSMNP1 [H.sapiens]	865173 IMAGE Consortium	AA450468
126 A	11 021BAMMO	ESTs, Highly similar to MO25_MOUSE MO25 PROTEIN [M.musculus]	865230 IMAGE Consortium	AA450759
127 C	1 021BAMMO	RNA polymerase 1-4 (194 kDa subunit)	821355 IMAGE Consortium	AA438036
128 C	3 021BAMMO	glucosidase I gene	821689 IMAGE Consortium	AA412802
129 C	5 021BAMMO	ESTs, Moderately similar to T00374 hypothetical protein KIAA0648 - human [H.sapiens]	821825 IMAGE Consortium	AA474185
130 C	7 021BAMMO	proteasome (prosome, macropain) 26S subunit, ATPase 3	865139 IMAGE Consortium	AA450445
131 C	9 021BAMMO	ESTs, Weakly similar to CG6210 gene product [D.melanogaster]	865185 IMAGE Consortium	AA450518
132 C	11 021BAMMO	phosphoglycerate kinase 1	865240 IMAGE Consortium	AA450763
133 E	1 021BAMMO	ESTs, Weakly similar to CG6482 gene product [D.melanogaster]	821371 IMAGE Consortium	AA438042
134 E	3 021BAMMO	ESTs, Weakly similar to SMB2_MOUSE DNA-BINDING PROTEIN SMUBP-2 [M.musculus]	806388 IMAGE Consortium	AA432841
135 E	5 021BAMMO	ESTs, Weakly similar to TERA_RAT TRANSITIONAL ENDOPLASMIC RETICULUM ATPASE [R.norvegicus]	865415 IMAGE Consortium	AI643201
136 E	7 021BAMMO	ESTs	865141 IMAGE Consortium	AA450446
137 E	9 021BAMMO	transforming growth factor alpha regulated gene 4	865188 IMAGE Consortium	AA450513

138 E	11 021BAMMO	serine palmitoyltransferase, long chain base subunit 1	865262 IMAGE Consortium	AA450780
139 G	1 021BAMMO	ESTs, Highly similar to 45kDa splicing factor [H.sapiens]	821373 IMAGE Consortium	AA438043
140 G	3 021BAMMO	ESTs, Weakly similar to CG10686 gene product [D.melanogaster]	821722 IMAGE Consortium	AI552366
141 G	5 021BAMMO	nucleosome assembly protein 1-like 4	865045 IMAGE Consortium	AA469641
142 G	7 021BAMMO	development and differentiation enhancing	865146 IMAGE Consortium	AA450440
143 G	9 021BAMMO	small inducible cytokine A19	865193 IMAGE Consortium	AA450521
144 G	11 021BAMMO	FK506 binding protein 8 (38 kDa)	865268 IMAGE Consortium	AA450783
145 A	1 021IAMMP	eukaryotic translation initiation factor 3	865317 IMAGE Consortium	AA450861
146 A	3 021IAMMP	src associated in mitosis, 68 kDa	865381 IMAGE Consortium	AA444584
147 A	5 021IAMMP	Swi/SNF related matrix associated, actin dependent regulator of chromatin, subfamily a-like 1	865764 IMAGE Consortium	AA458425
148 A	7 021IAMMP	ESTs	906271 IMAGE Consortium	AA529511
149 A	9 021IAMMP	ESTs, Weakly similar to Lpe10p [S.cerevisiae]	908471 IMAGE Consortium	AA589042
150 A	11 021IAMMP	ESTs, Highly similar to SMALL NUCLEAR RIBONUCLEOPROTEIN SM D3 [Homo sapiens]	906527 IMAGE Consortium	AA522225
151 C	1 021IAMMP	proteasome (prosome, macropain) subunit, beta type 6	865327 IMAGE Consortium	AA450909
152 C	3 021IAMMP	novel nuclear protein 1	865390 IMAGE Consortium	AA444579
153 C	5 021IAMMP	rabaptin 5	906002 IMAGE Consortium	AA529597
154 C	7 021IAMMP	ESTs	906323 IMAGE Consortium	AA529567
155 C	9 021IAMMP	ESTs	908565 IMAGE Consortium	AA591537
156 C	11 021IAMMP	proteasome (prosome, macropain) subunit, beta type 1	906551 IMAGE Consortium	AI551919
157 E	1 021IAMMP	ESTs, Highly similar to SUCCINATE DEHYDROGENASE [Homo sapiens]	865330 IMAGE Consortium	AA450902
158 E	3 021IAMMP	ESTs	866140 IMAGE Consortium	AA473706
159 E	5 021IAMMP	ESTs	906028 IMAGE Consortium	AI553351
160 E	7 021IAMMP	ESTs, Weakly similar to Cys2/His2 zinc finger protein [R.norvegicus]	906438 IMAGE Consortium	AA529691
161 E	9 021IAMMP	ESTs, Highly similar to dipeptidyl peptidase III [R.norvegicus]	908733 IMAGE Consortium	AA591381
162 E	11 021IAMMP	FK506 binding protein 4 (59 kDa)	907309 IMAGE Consortium	AA671029
163 G	1 021IAMMP	brachyury	865333 IMAGE Consortium	AA450910
164 G	3 021IAMMP	transgelin 2	865426 IMAGE Consortium	AI643202
165 G	5 021IAMMP	ESTs	906045 IMAGE Consortium	AA529654
166 G	7 021IAMMP	ESTs, Moderately similar to ATP SYNTHASE COUPLING FACTOR B, MITOCHONDRIAL [Bos taurus]	905765 IMAGE Consortium	AA529296
167 G	9 021IAMMP	Mus musculus zinc transporter like 2 (Znt12) mRNA, complete cds	907016 IMAGE Consortium	AI551957
168 G	11 021IAMMP	Mus musculus brain cDNA, clone MNCb-1706	906662 IMAGE Consortium	AA522308
169 A	1 021PAMMQ	ESTs	907488 IMAGE Consortium	AA588991
170 A	3 021PAMMQ	leucine zipper-EF-hand containing transmembrane protein 1	906947 IMAGE Consortium	AA591648
171 A	5 021PAMMQ			



172 A	7 021PAMMQ			
173 A	9 021PAMMQ	ESTs, Weakly similar to cDNA EST EMBL:M88812 comes from this gene [C.elegans]	514205 IMAGE Consortium	AA063897
174 A	11 021PAMMQ	ESTs, Weakly similar to coded for by C. elegans cDNA cm21e6 [C.elegans]	515153 IMAGE Consortium	AI595292
175 C	1 021PAMMQ	ESTs, Weakly similar to PAGT_RAT POLYPEPTIDE N-ACETYLGLACTOSAMINYLTRANSFERASE [R.norvegicus]	906775 IMAGE Consortium	AA522404
176 C	3 021PAMMQ	DNA polymerase epsilon, subunit 2	906997 IMAGE Consortium	AA591707
177 C	5 021PAMMQ	cAMP responsive element modulator	513759 IMAGE Consortium	AI385961
178 C	7 021PAMMQ			
179 C	9 021PAMMQ	topoisomerase (DNA) II alpha	514291 IMAGE Consortium	AA063958
180 C	11 021PAMMQ	general transcription factor IIH, polypeptide 1 (62kD subunit)	515209 IMAGE Consortium	AI892212
181 E	1 021PAMMQ	ESTs, Moderately similar to unnamed protein product [H.sapiens]	907562 IMAGE Consortium	AA590510
182 E	3 021PAMMQ	ESTs, Highly similar to S-ADENOSYLMETHIONINE SYNTHETASE GAMMA FORM [Rattus norvegicus]	907725 IMAGE Consortium	AA555793
183 E	5 021PAMMQ			
184 E	7 021PAMMQ	ESTs	514802 IMAGE Consortium	AI595255
185 E	9 021PAMMQ	ESTs, Highly similar to unnamed protein product [H.sapiens]	514366 IMAGE Consortium	AI595141
186 E	11 021PAMMQ	ESTs, Highly similar to Y143_HUMAN HYPOTHETICAL PROTEIN KIAA0143 [H.sapiens]	514501 IMAGE Consortium	AA064544
187 G	1 021PAMMQ	ESTs	906868 IMAGE Consortium	AI551943
188 G	3 021PAMMQ	ESTs	908545 IMAGE Consortium	AA590557
189 G	5 021PAMMQ			
190 G	7 021PAMMQ	dihydropyrimidinase-like 3	514004 IMAGE Consortium	AA062324
191 G	9 021PAMMQ	stannin	514373 IMAGE Consortium	AA061097
192 G	11 021PAMMQ	ESTs	515271 IMAGE Consortium	AA064399
193 A	1 021WAMMR	histone deacetylase 5	515336 IMAGE Consortium	AA064466
194 A	3 021WAMMR	ESTs	515522 IMAGE Consortium	AI595010
195 A	5 021WAMMR	ESTs	515027 IMAGE Consortium	AI595281
196 A	7 021WAMMR			
197 A	9 021WAMMR	golgi SNAP receptor complex member 1	515734 IMAGE Consortium	AA065486
198 A	11 021WAMMR	ESTs, Highly similar to Bing1 [M.musculus]	515942 IMAGE Consortium	AA062251
199 C	1 021WAMMR	dual adaptor for phosphotyrosine and 3-phosphoinositides 1	515361 IMAGE Consortium	AA064488
200 C	3 021WAMMR	guanine nucleotide binding protein, alpha 13	515531 IMAGE Consortium	AI595012
201 C	5 021WAMMR	ESTs, Moderately similar to CG11985 gene product [D.melanogaster]	515058 IMAGE Consortium	AA061271
202 C	7 021WAMMR	ESTs, Weakly similar to E04F6.2 gene product [C.elegans]	516103 IMAGE Consortium	AA089000
203 C	9 021WAMMR	Mus musculus male meiosis specific protein Tcte2 (Tcte2)	515779 IMAGE Consortium	AI595521

204 C	11 021WAMMR	mRNA, complete cds ESTs, Moderately similar to HYPOTHETICAL 84.5 KD PROTEIN IN CPS1-FPP1 INTERGENIC REGION [Saccharomyces cerevisiae]	515971 IMAGE Consortium	AI595543
205 E	1 021WAMMR	ESTs, Weakly similar to predicted using Genefinder [C.elegans]	515379 IMAGE Consortium	AI595619
206 E	3 021WAMMR	elongation of very long chain fatty acids (FEN1/Elo2, SUR4/Elo3, yeast)-like 2	514817 IMAGE Consortium	AI595258
207 E	5 021WAMMR			
208 E	7 021WAMMR	ESTs, Moderately similar to ZINC FINGER PROTEIN HF.12 [Homo sapiens]	516197 IMAGE Consortium	AI892221
209 E	9 021WAMMR	heat shock protein, 70 kDa 2	515812 IMAGE Consortium	AI385995
210 E	11 021WAMMR	serine/threonine kinase 22B (spermiogenesis associated)	515986 IMAGE Consortium	AA118345
211 G	1 021WAMMR	eukaryotic translation initiation factor 2 alpha kinase 1	515387 IMAGE Consortium	AI595623
212 G	3 021WAMMR	ESTs	515554 IMAGE Consortium	AA065383
213 G	5 021WAMMR	ash2 (absent, small, or homeotic)-like (Drosophila)	515373 IMAGE Consortium	AI595616
214 G	7 021WAMMR	ESTs, Weakly similar to glutamine repeat protein-1 [M.musculus]	516274 IMAGE Consortium	AI596566
215 G	9 021WAMMR	Public domain EST	515825 IMAGE Consortium	
216 G	11 021WAMMR	protein-L-isoaspartate (D-aspartate) O-methyltransferase 1	516378 IMAGE Consortium	AI596574
217 A	1 0213AMMS	defensin beta 1	517293 IMAGE Consortium	AA065510
218 A	3 0213AMMS	ESTs	517425 IMAGE Consortium	AI597095
219 A	5 0213AMMS	ESTs	517493 IMAGE Consortium	AA059501
220 A	7 0213AMMS	peptidase 4	517525 IMAGE Consortium	AA066579
221 A	9 0213AMMS	ferritin heavy chain	517695 IMAGE Consortium	AI892243
222 A	11 0213AMMS	ESTs, Weakly similar to periplakin [M.musculus]	519076 IMAGE Consortium	AA106834
223 C	1 0213AMMS	nuclear autoantigenic sperm protein (histone-binding)	516571 IMAGE Consortium	AA105299
224 C	3 0213AMMS	phosphofructokinase, liver, B-type	517433 IMAGE Consortium	AA062378
225 C	5 0213AMMS	solute carrier family 12, member 1	517502 IMAGE Consortium	AA065574
226 C	7 0213AMMS			
227 C	9 0213AMMS	ESTs, Highly similar to SINGLE-STRANDED DNA-BINDING PROTEIN, MITOCHONDRIAL PRECURSOR [Rattus norvegicus]	517721 IMAGE Consortium	AI892245
228 C	11 0213AMMS	ESTs, Weakly similar to LIM PROTEIN RIL [Rattus norvegicus]	519108 IMAGE Consortium	AI325572
229 E	1 0213AMMS	cellular nucleic acid binding protein	517318 IMAGE Consortium	AA065512
230 E	3 0213AMMS	ESTs	517447 IMAGE Consortium	AA059596
231 E	5 0213AMMS	craniofacial development protein 1	517562 IMAGE Consortium	AA066410
232 E	7 0213AMMS	ESTs, Weakly similar to T14171 ataxin-2 - mouse [M.musculus]	517558 IMAGE Consortium	AA066616
233 E	9 0213AMMS	vanin 1	517746 IMAGE Consortium	AI597120
234 E	11 0213AMMS	ESTs	519159 IMAGE Consortium	AA106002

235 G	1 0213AMMS	cytochrome c oxidase, subunit Va	517328 IMAGE Consortium	AA062293
236 G	3 0213AMMS	ectonucleotide pyrophosphatase/phosphodiesterase 1	517450 IMAGE Consortium	AA065567
237 G	5 0213AMMS	succinate-Coenzyme A ligase, GDP-forming, beta subunit	516843 IMAGE Consortium	AA062367
238 G	7 0213AMMS	DNA segment, Chr 4, Wayne State University 125, expressed	517579 IMAGE Consortium	AA066502
239 G	9 0213AMMS	Mus musculus elafin-like protein I mRNA, complete cds	517757 IMAGE Consortium	AA066601
240 G	11 0213AMMS	ESTs, Highly similar to C38637 Ras inhibitor [H.sapiens]	519231 IMAGE Consortium	AA106210
241 A	1 021AAMMT	ESTs	519416 IMAGE Consortium	AI597354
242 A	3 021AAMMT	ESTs, Weakly similar to AF153613_1 peroxisomal D3,D2-enoyl-CoA isomerase [M.musculus]	518672 IMAGE Consortium	AA105274
243 A	5 021AAMMT	heat-responsive protein 12	518907 IMAGE Consortium	AA106879
244 A	7 021AAMMT	ESTs, Highly similar to PEROXISOMAL CARNITINE OCTANOYLTRANSFERASE [Rattus norvegicus]	519562 IMAGE Consortium	AA106580
245 A	9 021AAMMT	hypothetical protein	520117 IMAGE Consortium	AA117581
246 A	11 021AAMMT	ESTs, Highly similar to HYPOTHETICAL 25.7 KD PROTEIN IN MSH1-EPT1 INTERGENIC REGION [Saccharomyces cerevisiae]	520373 IMAGE Consortium	AA110928
247 C	1 021AAMMT	ATP-binding cassette, sub-family D (ALD), member 4	518750 IMAGE Consortium	AA105879
248 C	3 021AAMMT	interferon inducible protein 1	518695 IMAGE Consortium	AA105762
249 C	5 021AAMMT	ESTs, Moderately similar to KIDNEY ANDROGEN-REGULATED PROTEIN PRECURSOR [Mus musculus]	518930 IMAGE Consortium	AA106167
250 C	7 021AAMMT	interferon-related developmental regulator 1	519583 IMAGE Consortium	AI324408
251 C	9 021AAMMT	insulin receptor-related receptor	520173 IMAGE Consortium	AI552669
252 C	11 021AAMMT	solute carrier family 34 (sodium phosphate), member 1	520285 IMAGE Consortium	AA110887
253 E	1 021AAMMT	DNA segment, Chr 11, Wayne State University 68, expressed	518875 IMAGE Consortium	AA106733
254 E	3 021AAMMT	ESTs	518738 IMAGE Consortium	AA105771
255 E	5 021AAMMT	ESTs	518934 IMAGE Consortium	AA106169
256 E	7 021AAMMT	ESTs	519801 IMAGE Consortium	AA109528
257 E	9 021AAMMT	brain protein 44-like protein	520197 IMAGE Consortium	AA107936
258 E	11 021AAMMT	corticosteroid binding globulin	521016 IMAGE Consortium	AA066835
259 G	1 021AAMMT	cytochrome c oxidase, subunit VIIa 3	518896 IMAGE Consortium	AA106736
260 G	3 021AAMMT	Mus musculus evectin-2 (Evt2) mRNA, complete cds	518785 IMAGE Consortium	AA089173
261 G	5 021AAMMT	hemoglobin beta chain complex	518979 IMAGE Consortium	AA106071
262 G	7 021AAMMT	phospholipid scramblase 2	519938 IMAGE Consortium	AA110551
263 G	9 021AAMMT	CASP2 and RIPK1 domain containing adaptor with death domain	520277 IMAGE Consortium	AA109538
264 G	11 021AAMMT	FMS-like tyrosine kinase 1	521021 IMAGE Consortium	AI326556
265 A	1 021HAMMU	ESTs, Weakly similar to AF114379_1 cypher2 [M.musculus]	521169 IMAGE Consortium	AA065720
266 A	3 021HAMMU	alpha-2-macroglobulin	520709 IMAGE Consortium	AA066397
267 A	5 021HAMMU	sulfotransferase family, cytosolic, 2B, member 1	445155 IMAGE Consortium	AI322504
268 A	7 021HAMMU	ESTs, Moderately similar to Edp1 protein [M.musculus]	444899 IMAGE Consortium	AA016788

269 A	9 021HAMMU	integral membrane glycoprotein	523153 IMAGE Consortium	AA067284
270 A	11 021HAMMU	ESTs, Highly similar to ZINC FINGER PROTEIN 7 [Homo sapiens]	522715 IMAGE Consortium	AA087674
271 C	1 021HAMMU	macrophage activation 2	520463 IMAGE Consortium	AA105281
272 C	3 021HAMMU	Mus musculus phosphoglycerate mutase 1 (Pgam1) mRNA, complete cds	520733 IMAGE Consortium	AA065739
273 C	5 021HAMMU	guanine nucleotide binding protein, alpha q polypeptide	443239 IMAGE Consortium	AA015481
274 C	7 021HAMMU	crystallin, beta A1	439189 IMAGE Consortium	AA003874
275 C	9 021HAMMU	ESTs	523158 IMAGE Consortium	AA067279
276 C	11 021HAMMU	ESTs	522719 IMAGE Consortium	AI552805
277 E	1 021HAMMU	glutathione S-transferase, mu 1	521233 IMAGE Consortium	AA067003
278 E	3 021HAMMU	gap junction membrane channel protein beta 1	520762 IMAGE Consortium	AA065775
279 E	5 021HAMMU	goosecoid	441833 IMAGE Consortium	AA013883
280 E	7 021HAMMU	retinoic acid receptor, alpha	439287 IMAGE Consortium	AA003987
281 E	9 021HAMMU	eukaryotic translation initiation factor 2 alpha kinase 3	522441 IMAGE Consortium	AI552787
282 E	11 021HAMMU	ESTs, Highly similar to R27216_1 [H.sapiens]	522793 IMAGE Consortium	AA087438
283 G	1 021HAMMU	S-adenosylmethionine synthetase	521238 IMAGE Consortium	AA067002
284 G	3 021HAMMU	ESTs	520861 IMAGE Consortium	AI326356
285 G	5 021HAMMU	ESTs	444147 IMAGE Consortium	AA014807
286 G	7 021HAMMU	ESTs, Highly similar to REGULATOR OF G-PROTEIN SIGNALLING 2 [Homo sapiens]	439412 IMAGE Consortium	AA004108
287 G	9 021HAMMU	guanosine diphosphate (GDP) dissociation inhibitor 3	522453 IMAGE Consortium	AI325613
288 G	11 021HAMMU	presenilin 1	522798 IMAGE Consortium	AA087434
289 A	1 021OAMMV	ESTs, Highly similar to calcium-independent phospholipase A2 [H.sapiens]	522945 IMAGE Consortium	AA065782
290 A	3 021OAMMV	ESTs, Weakly similar to 60S RIBOSOMAL PROTEIN L30A [Saccharomyces cerevisiae]	521772 IMAGE Consortium	AI386067
291 A	5 021OAMMV	ESTs, Highly similar to ABC1 PROTEIN PRECURSOR [Saccharomyces cerevisiae]	521180 IMAGE Consortium	AI552731
292 A	7 021OAMMV	actopaxin	521622 IMAGE Consortium	AA066695
293 A	9 021OAMMV	keratin associated protein 12-4	523308 IMAGE Consortium	AA079946
294 A	11 021OAMMV	ESTs, Highly similar to AF151859_1 CGI-101 protein [H.sapiens]	523452 IMAGE Consortium	AA080205
295 C	1 021OAMMV	ESTs	522990 IMAGE Consortium	AI552850
296 C	3 021OAMMV			
297 C	5 021OAMMV	ESTs, Highly similar to CARBAMOYL-PHOSPHATE SYNTHASE [Rattus norvegicus]	522024 IMAGE Consortium	AI386236
298 C	7 021OAMMV	cytochrome P450, 2e1, ethanol inducible	521645 IMAGE Consortium	AI386063
299 C	9 021OAMMV	ESTs, Highly similar to HYPOTHETICAL 55.1 KD PROTEIN IN FAB1-PES4 INTERGENIC REGION [Saccharomyces cerevisiae]	523315 IMAGE Consortium	AA079906
300 C	11 021OAMMV	adipocyte protein aP2	523460 IMAGE Consortium	AA080270
301 E	1 021OAMMV	ral guanine nucleotide dissociation stimulator	522213 IMAGE Consortium	AA067971

302 E	3	021OAMMV	transcription factor EB	521847	IMAGE Consortium	AA106636
303 E	5	021OAMMV	cytochrome P450, 2a4	522059	IMAGE Consortium	AA106125
304 E	7	021OAMMV	eukaryotic translation initiation factor 2, subunit 3, structural gene X-linked	521675	IMAGE Consortium	AA066780
305 E	9	021OAMMV	ESTs, Weakly similar to U82695_2 expressed-Xq28STS protein [H.sapiens]	523337	IMAGE Consortium	AA079990
306 E	11	021OAMMV	ESTs	523567	IMAGE Consortium	AA097339
307 G	1	021OAMMV	glucocorticoid-induced leucine zipper	523039	IMAGE Consortium	AA066442
308 G	3	021OAMMV	eukaryotic translation elongation factor 2	521857	IMAGE Consortium	AI386071
309 G	5	021OAMMV	ESTs	521350	IMAGE Consortium	AA066941
310 G	7	021OAMMV	fatso	523141	IMAGE Consortium	AA067230
311 G	9	021OAMMV	Mus musculus NudeL mRNA, complete cds	523372	IMAGE Consortium	AA079916
312 G	11	021OAMMV	ESTs, Highly similar to LACTOPEROXIDASE [Homo sapiens]	522902	IMAGE Consortium	AI892299
313 A	1	021VAMMW	ribonucleotide reductase M1	523993	IMAGE Consortium	AA072017
314 A	3	021VAMMW	selenoprotein P, plasma, 1	524191	IMAGE Consortium	AA066225
315 A	5	021VAMMW	ESTs, Highly similar to AF161445_1 HSPC327 [H.sapiens]	523669	IMAGE Consortium	AA067304
316 A	7	021VAMMW	ESTs	523976	IMAGE Consortium	AA072083
317 A	9	021VAMMW	DNA segment, Chr 4, ERATO Doi 478, expressed	524655	IMAGE Consortium	AA068133
318 A	11	021VAMMW	ESTs	524862	IMAGE Consortium	AA097069
319 C	1	021VAMMW	fibroblast growth factor receptor 1	523025	IMAGE Consortium	AA067236
320 C	3	021VAMMW	ESTs	524211	IMAGE Consortium	AA066237
321 C	5	021VAMMW	ESTs, Moderately similar to T00362 hypothetical protein KIAA0675 - human [H.sapiens]	523734	IMAGE Consortium	AI613965
322 C	7	021VAMMW	protein S (alpha)	524437	IMAGE Consortium	AI326590
323 C	9	021VAMMW	ESTs, Highly similar to KIAA1223 protein [H.sapiens]	524683	IMAGE Consortium	AA068501
324 C	11	021VAMMW	DNA segment, Chr 16, human D22S680E, expressed	524932	IMAGE Consortium	AI326596
325 E	1	021VAMMW	semaF cytoplasmic domain associated protein 1	523099	IMAGE Consortium	AA067058
326 E	3	021VAMMW	ESTs, Moderately similar to similar to a C.elegans protein encoded in cosmid K12D12(Z49069) [H.sapiens]	524257	IMAGE Consortium	AA065683
327 E	5	021VAMMW	Hus1 homolog (S. pombe)	523771	IMAGE Consortium	AA067186
328 E	7	021VAMMW	retinoic acid induced 12	524467	IMAGE Consortium	AA068056
329 E	9	021VAMMW	acetyl-Coenzyme A dehydrogenase, medium chain	524703	IMAGE Consortium	AA068116
330 E	11	021VAMMW	histone gene complex 1	524976	IMAGE Consortium	AA089022
331 G	1	021VAMMW	EST	524007	IMAGE Consortium	AI428095
332 G	3	021VAMMW	helicase, lymphoid specific	524259	IMAGE Consortium	AA065684
333 G	5	021VAMMW	apoptosis inhibitory protein 5	523813	IMAGE Consortium	AA066960
334 G	7	021VAMMW	ESTs, Moderately similar to EHM2 [M.musculus]	524489	IMAGE Consortium	AA068097
335 G	9	021VAMMW	lymphocyte antigen 6 complex, locus D	524731	IMAGE Consortium	AI386262
336 G	11	021VAMMW	myristoylated alanine rich protein kinase C substrate	524982	IMAGE Consortium	AA089025
337 A	1	0212AMMX	calpain 4	524157	IMAGE Consortium	AA066140
338 A	3	0212AMMX	lysosomal membrane glycoprotein 1	524298	IMAGE Consortium	AA067897

339 A	5 0212AMMX	hydroxyacid oxidase (glycolate oxidase) 3	524645 IMAGE Consortium	AA068624
340 A	7 0212AMMX	ESTs, Weakly similar to type III collagen [M.musculus]	533220 IMAGE Consortium	AA068280
341 A	9 0212AMMX	Down syndrome critical region homolog 2 (human)	524878 IMAGE Consortium	AA088963
342 A	11 0212AMMX	ESTs, Highly similar to SEC PROTEIN [Homo sapiens]	533272 IMAGE Consortium	AA105453
343 C	1 0212AMMX	ESTs	524176 IMAGE Consortium	AA065620
344 C	3 0212AMMX	ESTs	524388 IMAGE Consortium	AA065755
345 C	5 0212AMMX	ESTs, Weakly similar to CG1534 gene product [D.melanogaster]	524668 IMAGE Consortium	AA068627
346 C	7 0212AMMX	ESTs, Highly similar to hypothetical protein [H.sapiens]	533243 IMAGE Consortium	AI551979
347 C	9 0212AMMX	ESTs, Weakly similar to S38783 integrin alpha chain - rat [R.norvegicus]	524896 IMAGE Consortium	AA088968
348 C	11 0212AMMX			
349 E	1 0212AMMX	ESTs, Weakly similar to transcription repressor [H.sapiens]	524182 IMAGE Consortium	AI614021
350 E	3 0212AMMX	calbindin-D9K	524394 IMAGE Consortium	AI386257
351 E	5 0212AMMX	ESTs, Highly similar to POLYPEPTIDE N-ACETYL GALACTOSAMINYLTRANSFERASE [Bos taurus]	524714 IMAGE Consortium	AA068195
352 E	7 0212AMMX	ESTs, Weakly similar to BUTY_MOUSE BUTYROPHILIN PRECURSOR [M.musculus]	532722 IMAGE Consortium	AI643140
353 E	9 0212AMMX	ESTs	532924 IMAGE Consortium	AA068297
354 E	11 0212AMMX	c-fos induced growth factor	533306 IMAGE Consortium	AA104537
355 G	1 0212AMMX	ESTs, Weakly similar to mp86 [M.musculus]	524228 IMAGE Consortium	AI326418
356 G	3 0212AMMX	septin 2	524422 IMAGE Consortium	AA067948
357 G	5 0212AMMX	ESTs, Weakly similar to CG12194 gene product [D.melanogaster]	524741 IMAGE Consortium	AA068206
358 G	7 0212AMMX	BTB and CNC homology 1	524824 IMAGE Consortium	AA088927
359 G	9 0212AMMX	ferredoxin 1	524945 IMAGE Consortium	AA096861
360 G	11 0212AMMX	DNA segment, Chr 15, ERATO Doi 81, expressed	534075 IMAGE Consortium	AA072596
361 A	1 0219AMMY	glucose-6-phosphatase, transport protein 1	534184 IMAGE Consortium	AI552049
362 A	3 0219AMMY	ESTs	533797 IMAGE Consortium	AI426638
363 A	5 0219AMMY	telomeric repeat binding factor 2	550831 IMAGE Consortium	AA098594
364 A	7 0219AMMY	muskelin 1, intracellular mediator containing kelch motifs	536663 IMAGE Consortium	AI550052
365 A	9 0219AMMY	ATPase, H+ transporting, lysosomal (vacuolar proton pump) 16kD	535528 IMAGE Consortium	AI325715
366 A	11 0219AMMY	SDP8 protein	535912 IMAGE Consortium	AA105517
367 C	1 0219AMMY	DNA polymerase alpha 2, 68 kDa	534195 IMAGE Consortium	AA072666
368 C	3 0219AMMY	ESTs	533816 IMAGE Consortium	AA071849
369 C	5 0219AMMY	ESTs	536287 IMAGE Consortium	AA073603
370 C	7 0219AMMY			
371 C	9 0219AMMY	ESTs	535568 IMAGE Consortium	AA108058
372 C	11 0219AMMY	dystrophia myotonica kinase, B15	476127 IMAGE Consortium	AA050380
373 E	1 0219AMMY	tyrosine 3-monooxygenase/tryptophan 5-monooxygenase	534215 IMAGE Consortium	AA072682

374 E	3 0219AMMY	activation protein, epsilon polypeptide uroporphyrinogen decarboxylase	533847 IMAGE Consortium	AA071596
375 E	5 0219AMMY	DNA segment, Chr 14, ERATO Doi 226, expressed	536415 IMAGE Consortium	AA073868
376 E	7 0219AMMY	poly (A) polymerase	535029 IMAGE Consortium	AA073307
377 E	9 0219AMMY	ESTs, Highly similar to AF067728_1 transactivating protein BRIDGE [R.norvegicus]	535629 IMAGE Consortium	AI552117
378 E	11 0219AMMY	myeloid ecotropic viral integration site-related gene 1	477909 IMAGE Consortium	AI323035
379 G	1 0219AMMY	cell line MC/9.IL4 derived transcript 1	534231 IMAGE Consortium	AA072751
380 G	3 0219AMMY	ESTs, Weakly similar to CD82_MOUSE CD82 ANTIGEN [M.musculus]	533937 IMAGE Consortium	AA072834
381 G	5 0219AMMY	alpha-methylacyl-Coenzyme A racemase	536523 IMAGE Consortium	AA073874
382 G	7 0219AMMY	nuclear transcription factor-Y gamma	535044 IMAGE Consortium	AA073194
383 G	9 0219AMMY	ribosomal protein L19	534910 IMAGE Consortium	AI892387
384 G	11 0219AMMY			
385 A	1 021GAMMZ	growth arrest specific	477480 IMAGE Consortium	AA048398
386 A	3 021GAMMZ	ESTs	477897 IMAGE Consortium	AA048477
387 A	5 021GAMMZ	potassium voltage gated channel, Shaw-related subfamily, member 1	493098 IMAGE Consortium	AA097374
388 A	7 021GAMMZ	ESTs	570573 IMAGE Consortium	AI587727
389 A	9 021GAMMZ	ESTs	493009 IMAGE Consortium	AI427754
390 A	11 021GAMMZ	procollagen, type IX, alpha 2	480602 IMAGE Consortium	AI595449
391 C	1 021GAMMZ	phosphodiesterase 4A, cAMP specific	478225 IMAGE Consortium	AI510059
392 C	3 021GAMMZ	deltex 1 homolog (Drosophila)	477914 IMAGE Consortium	AI510041
393 C	5 021GAMMZ	ESTs, Weakly similar to Luman [H.sapiens]	492422 IMAGE Consortium	AI595179
394 C	7 021GAMMZ	ESTs	572018 IMAGE Consortium	AI587822
395 C	9 021GAMMZ	feline sarcoma oncogene	481510 IMAGE Consortium	AA060312
396 C	11 021GAMMZ	non-agouti	482132 IMAGE Consortium	AI323098
397 E	1 021GAMMZ	eyes absent 3 homolog (Drosophila)	478302 IMAGE Consortium	AA049689
398 E	3 021GAMMZ	ESTs, Highly similar to transcription factor C1 [M.musculus]	479409 IMAGE Consortium	AI323056
399 E	5 021GAMMZ	ESTs, Highly similar to KIAA0169 protein [H.sapiens]	573444 IMAGE Consortium	AA118619
400 E	7 021GAMMZ	ESTs	570607 IMAGE Consortium	AI587730
401 E	9 021GAMMZ	ESTs, Weakly similar to CARBONIC ANHYDRASE III [Homo sapiens]	482940 IMAGE Consortium	AI385507
402 E	11 021GAMMZ	ESTs	480745 IMAGE Consortium	AI595457
403 G	1 021GAMMZ	ESTs, Highly similar to EFB3_MOUSE EPHRIN-B3 PRECURSOR [M.musculus]	476970 IMAGE Consortium	AI510011
404 G	3 021GAMMZ	polycystic kidney disease 1 homolog	479447 IMAGE Consortium	AA049302
405 G	5 021GAMMZ	amyloid beta (A4) precursor protein-binding, family A, member 3	572836 IMAGE Consortium	AI892846
406 G	7 021GAMMZ	ESTs, Highly similar to inwardly rectifying potassium channel Kir5.1 [M.musculus]	572121 IMAGE Consortium	AI587823
407 G	9 021GAMMZ	neuronal protein	479918 IMAGE Consortium	AA051560

408 G	11 021GAMMZ	peptidylglycine alpha-amidating monooxygenase	482270 IMAGE Consortium	AI323101
409 A	1 021GAMN0	receptor (calcitonin) activity modifying protein 1	483161 IMAGE Consortium	AA063790
410 A	3 021GAMN0	ESTs	574161 IMAGE Consortium	AI594994
411 A	5 021GAMN0	EST	573827 IMAGE Consortium	AI452125
412 A	7 021GAMN0	RAD52 homolog, ( <i>S. cerevisiae</i> )	575467 IMAGE Consortium	AA119520
413 A	9 021GAMN0	ATP-binding cassette, sub-family C (CFTR/MRP), member 5a	987788 IMAGE Consortium	AA570953
414 A	11 021GAMN0	cytochrome P450, 2d10	1001079 IMAGE Consortium	AA771393
415 C	1 021GAMN0	crystallin, gamma B	482548 IMAGE Consortium	AI385493
416 C	3 021GAMN0	ESTs	573580 IMAGE Consortium	AA119371
417 C	5 021GAMN0	small inducible cytokine A2	573898 IMAGE Consortium	AI323202
418 C	7 021GAMN0	ESTs	576271 IMAGE Consortium	AI551472
419 C	9 021GAMN0	phenylalkylamine Ca <sup>2+</sup> antagonist (emopamil) binding protein	987831 IMAGE Consortium	AA571047
420 C	11 021GAMN0	ESTs	1001131 IMAGE Consortium	AA771433
421 E	1 021GAMN0	TEA domain family member 4	483286 IMAGE Consortium	AA061482
422 E	3 021GAMN0	ESTs	575028 IMAGE Consortium	AI530538
423 E	5 021GAMN0	potassium voltage gated channel, shaker related subfamily, member 3	575326 IMAGE Consortium	AA119270
424 E	7 021GAMN0	ESTs	575572 IMAGE Consortium	AA124822
425 E	9 021GAMN0	ESTs, Weakly similar to unnamed protein product [ <i>M.musculus</i> ]	987861 IMAGE Consortium	AA571070
426 E	11 021GAMN0	ESTs	1002026 IMAGE Consortium	AA684275
427 G	1 021GAMN0	Mus musculus DNMT1 associated protein-1 (Dmap1) mRNA, complete cds	481887 IMAGE Consortium	AA059689
428 G	3 021GAMN0	ESTs, Highly similar to RIBOSE-PHOSPHATE PYROPHOSPHOKINASE I [ <i>Homo sapiens</i> ; <i>Rattus norvegicus</i> ]	575082 IMAGE Consortium	AA120675
429 G	5 021GAMN0	ESTs	574627 IMAGE Consortium	AA118384
430 G	7 021GAMN0	ESTs, Highly similar to serine protease [ <i>H.sapiens</i> ]	576361 IMAGE Consortium	AI536198
431 G	9 021GAMN0	tryptophan 2,3-dioxygenase	987444 IMAGE Consortium	AA572623
432 G	11 021GAMN0	ESTs, Weakly similar to dJ75K24.1 [ <i>H.sapiens</i> ]	1001364 IMAGE Consortium	AA771615
433 A	1 021NAMN1	ESTs	1001446 IMAGE Consortium	AI587944
434 A	3 021NAMN1	ESTs	1001713 IMAGE Consortium	AI504075
435 A	5 021NAMN1	ESTs	1002620 IMAGE Consortium	AA596764
436 A	7 021NAMN1	histidyl tRNA synthetase	1021237 IMAGE Consortium	AA636154
437 A	9 021NAMN1	transcription factor 4	1020812 IMAGE Consortium	AA645758
438 A	11 021NAMN1	clathrin adaptor protein mu3A	1002198 IMAGE Consortium	AA608117
439 C	1 021NAMN1	ESTs, Weakly similar to AF234765_1 serine-arginine-rich splicing regulatory protein SRRP86 [ <i>R.norvegicus</i> ]	1001473 IMAGE Consortium	AA666485
440 C	3 021NAMN1	ESTs	1001725 IMAGE Consortium	AA666767
441 C	5 021NAMN1	coactivator-associated arginine methyltransferase 1	1002622 IMAGE Consortium	AA596765
442 C	7 021NAMN1	DNA segment, Chr 11, ERATO Doi 730, expressed	1020564 IMAGE Consortium	AI594144



443 C	9 021NAMN1	ESTs	1020833 IMAGE Consortium	AA645782
444 C	11 021NAMN1	ESTs, Highly similar to AF155352_1 ankyrin repeat-containing protein Asb-1 [M.musculus]	1002206 IMAGE Consortium	AA608121
445 E	1 021NAMN1	ESTs, Highly similar to tetraspan NET-6 [H.sapiens]	1001499 IMAGE Consortium	AA666505
446 E	3 021NAMN1	CCAAT/enhancer binding protein (C/EBP), gamma	1001742 IMAGE Consortium	AI587987
447 E	5 021NAMN1	ESTs	1002689 IMAGE Consortium	AI606354
448 E	7 021NAMN1	beta-2 microglobulin	1020580 IMAGE Consortium	AI594147
449 E	9 021NAMN1	ESTs, Moderately similar to R30923_1 [H.sapiens]	1020875 IMAGE Consortium	AA645930
450 E	11 021NAMN1	galactose-1-phosphate uridyl transferase	1002214 IMAGE Consortium	AA596430
451 G	1 021NAMN1	ESTs	1001541 IMAGE Consortium	AA666573
452 G	3 021NAMN1	ESTs	1001809 IMAGE Consortium	AI587999
453 G	5 021NAMN1	ESTs	1002756 IMAGE Consortium	AA607476
454 G	7 021NAMN1	ESTs, Moderately similar to Glutamyl tRNA synthetase [H.sapiens]	1020618 IMAGE Consortium	AA647905
455 G	9 021NAMN1	protease, serine, 25	1020887 IMAGE Consortium	AA645936
456 G	11 021NAMN1	ESTs	1002249 IMAGE Consortium	AI606309
457 A	1 021UAMN2	ESTs, Highly similar to Tera [M.musculus]	1002413 IMAGE Consortium	AA596499
458 A	3 021UAMN2	ESTs	1002697 IMAGE Consortium	AA607448
459 A	5 021UAMN2	baculoviral IAP repeat-containing 5	1038683 IMAGE Consortium	AA671290
460 A	7 021UAMN2	branched chain ketoacid dehydrogenase E1, alpha polypeptide	1038945 IMAGE Consortium	AA624850
461 A	9 021UAMN2	ESTs, Highly similar to ALCOHOL DEHYDROGENASE [Anas platyrhynchos]	1023849 IMAGE Consortium	AA866943
462 A	11 021UAMN2	voltage-dependent anion channel 2	1038815 IMAGE Consortium	AA624465
463 C	1 021UAMN2	ESTs, Moderately similar to DHSD_HUMAN SUCCINATE DEHYDROGENASE [H.sapiens]	1002453 IMAGE Consortium	AA596582
464 C	3 021UAMN2	Braf transforming gene	1002723 IMAGE Consortium	AA607463
465 C	5 021UAMN2	myosin binding protein C, cardiac	1038704 IMAGE Consortium	AA671319
466 C	7 021UAMN2	ESTs	1023333 IMAGE Consortium	AA673542
467 C	9 021UAMN2	lectin, galactose binding, soluble 7	1023899 IMAGE Consortium	AA864136
468 C	11 021UAMN2	ESTs	1038856 IMAGE Consortium	AA624553
469 E	1 021UAMN2	ESTs	1002463 IMAGE Consortium	AA596525
470 E	3 021UAMN2	mitogen activated protein kinase kinase kinase 3	1002768 IMAGE Consortium	AA607484
471 E	5 021UAMN2	signal recognition particle receptor, B subunit	1038773 IMAGE Consortium	AA671503
472 E	7 021UAMN2	periplakin	1023414 IMAGE Consortium	AA647175
473 E	9 021UAMN2	ESTs	1039025 IMAGE Consortium	AA671156
474 E	11 021UAMN2	ESTs, Weakly similar to ras-related protein [M.musculus]	1038883 IMAGE Consortium	AA624579
475 G	1 021UAMN2	ESTs, Weakly similar to myelin transcription factor 1 [M.musculus]	1002477 IMAGE Consortium	AA596587
476 G	3 021UAMN2	actin-related protein 11 homolog (S. cerevisiae)	1021054 IMAGE Consortium	AA645795
477 G	5 021UAMN2	integrin linked kinase	1038823 IMAGE Consortium	AA624474
478 G	7 021UAMN2	ESTs	1023585 IMAGE Consortium	AA647096
479 G	9 021UAMN2	aconitase 2, mitochondrial	1039155 IMAGE Consortium	AA671470

480 G	11 021UAMN2	hemoglobin Y, beta-like embryonic chain	1038954 IMAGE Consortium	AA624843
481 A	1 0211AMN3	ESTs	1040023 IMAGE Consortium	AA612121
482 A	3 0211AMN3	ESTs	1040499 IMAGE Consortium	AI594517
483 A	5 0211AMN3	ESTs, Weakly similar to CG4749 gene product [D.melanogaster]	1077547 IMAGE Consortium	AI594830
484 A	7 0211AMN3	group specific component	1077718 IMAGE Consortium	AA822029
485 A	9 0211AMN3	erythrocyte protein band 7.2	1077484 IMAGE Consortium	AA821885
486 A	11 0211AMN3	ESTs, Weakly similar to KIAA0294 [H.sapiens]	1040200 IMAGE Consortium	AA792288
487 C	1 0211AMN3	nuclear factor I/X	1040038 IMAGE Consortium	AA612117
488 C	3 0211AMN3	murinoglobulin 1	1091724 IMAGE Consortium	AA670733
489 C	5 0211AMN3	ESTs, Moderately similar to T00335 hypothetical protein KIAA0564 - human [H.sapiens]	1077563 IMAGE Consortium	AI594834
490 C	7 0211AMN3	collagen binding protein 1	1077734 IMAGE Consortium	AA822044
491 C	9 0211AMN3	small inducible cytokine A6	1077529 IMAGE Consortium	AA821932
492 C	11 0211AMN3	procollagen, type V, alpha 1	1040230 IMAGE Consortium	AA792297
493 E	1 0211AMN3	radical fringe gene homolog, (Drosophila)	1040071 IMAGE Consortium	AA615228
494 E	3 0211AMN3	ESTs, Weakly similar to A55071 hydrogen peroxide-inducible protein hic-5 - mouse [M.musculus]	1091798 IMAGE Consortium	AI593658
495 E	5 0211AMN3	Mus musculus zinc finger protein 94 (Zfp94) mRNA, complete cds	1077575 IMAGE Consortium	AA822076
496 E	7 0211AMN3	WW domain binding protein 4	1077755 IMAGE Consortium	AA822111
497 E	9 0211AMN3	islet neogenesis associated protein-related protein	1077556 IMAGE Consortium	AI594831
498 E	11 0211AMN3	ESTs	1040369 IMAGE Consortium	AA612189
499 G	1 0211AMN3	ESTs, Highly similar to unnamed protein product [H.sapiens]	1040119 IMAGE Consortium	AA615254
500 G	3 0211AMN3	ESTs, Moderately similar to AF161418_1 HSPC300 [H.sapiens]	1077402 IMAGE Consortium	AA733472
501 G	5 0211AMN3	nuclear receptor subfamily 1, group I, member 3	1077589 IMAGE Consortium	AA821956
502 G	7 0211AMN3	isocitrate dehydrogenase 3 (NAD+), gamma	1077761 IMAGE Consortium	AA822114
503 G	9 0211AMN3	paraoxonase 1	1077579 IMAGE Consortium	AA821953
504 G	11 0211AMN3	ESTs	1040517 IMAGE Consortium	AA612414
505 A	1 0218AMN4	ESTs	1040667 IMAGE Consortium	AA606318
506 A	3 0218AMN4	cytochrome P450, 2b9, phenobarbitol inducible, type a	1092486 IMAGE Consortium	AA656694
507 A	5 0218AMN4	ESTs, Highly similar to GMP REDUCTASE [Homo sapiens]	1092146 IMAGE Consortium	AA670938
508 A	7 0218AMN4	fatty acid amide hydrolase	1092351 IMAGE Consortium	AA822583
509 A	9 0218AMN4	chitinase 3-like 3	1148798 IMAGE Consortium	AA673731
510 A	11 0218AMN4	ESTs, Highly similar to KIAA0983 protein [H.sapiens]	1148500 IMAGE Consortium	AA646456
511 C	1 0218AMN4	lung carcinoma myc related oncogene 1	1040702 IMAGE Consortium	AA606331
512 C	3 0218AMN4	Mus musculus partial mRNA for myosin heavy chain IIB (MYHC-IIB gene)	1092501 IMAGE Consortium	AA656712
513 C	5 0218AMN4	protein related to DAC and cerberus	1092199 IMAGE Consortium	AA755273
514 C	7 0218AMN4	ESTs	1139924 IMAGE Consortium	AA645533

515 C	9 0218AMN4	ESTs	1140255 IMAGE Consortium	AI591801
516 C	11 0218AMN4	Mouse mRNA for TI-227	1148534 IMAGE Consortium	AA672766
517 E	1 0218AMN4	autocrine motility factor receptor	1040709 IMAGE Consortium	AI594561
518 E	3 0218AMN4	cytochrome P450, 2a5	1092653 IMAGE Consortium	AA674177
519 E	5 0218AMN4	ESTs, Highly similar to interleukin 17 receptor□ [M.musculus]	1139646 IMAGE Consortium	AA646250
520 E	7 0218AMN4	ESTs, Moderately similar to AF151830_1 CGI-72 protein [H.sapiens]	1139960 IMAGE Consortium	AI591771
521 E	9 0218AMN4	ESTs	1140350 IMAGE Consortium	AA646153
522 E	11 0218AMN4	ESTs	1148545 IMAGE Consortium	AA672782
523 G	1 0218AMN4	ESTs	1040747 IMAGE Consortium	AI594566
524 G	3 0218AMN4	nuclear receptor subfamily 2, group H, member 2	1092732 IMAGE Consortium	AA674318
525 G	5 0218AMN4	eukaryotic translation elongation factor 1 alpha 2	1092217 IMAGE Consortium	AA755371
526 G	7 0218AMN4			
527 G	9 0218AMN4	TG interacting factor	1139605 IMAGE Consortium	AA645995
528 G	11 0218AMN4	Mus musculus mRNA for erythroid differentiation regulator, partial	1148549 IMAGE Consortium	AA672784
529 A	1 021FAMN5			
530 A	3 021FAMN5			
531 A	5 021FAMN5	Mus musculus E2F-like transcriptional repressor protein mRNA, complete cds	1149675 IMAGE Consortium	AA656128
532 A	7 021FAMN5	lymphocyte protein tyrosine kinase	1148872 IMAGE Consortium	AA717185
533 A	9 021FAMN5	ESTs, Weakly similar to ML64_MOUSE MLN 64 PROTEIN [M.musculus]	1149054 IMAGE Consortium	AA655683
534 A	11 021FAMN5	ESTs	1149142 IMAGE Consortium	AA655780
535 C	1 021FAMN5			
536 C	3 021FAMN5			
537 C	5 021FAMN5	DNA segment, Chr 10, Wayne State University 52, expressed	1149725 IMAGE Consortium	AA656168
538 C	7 021FAMN5	a disintegrin and metalloprotease domain 8	1148891 IMAGE Consortium	AA655872
539 C	9 021FAMN5	ESTs	1149062 IMAGE Consortium	AA655686
540 C	11 021FAMN5	ESTs	1149156 IMAGE Consortium	AA655797
541 E	1 021FAMN5			
542 E	3 021FAMN5			
543 E	5 021FAMN5	ESTs, Highly similar to AC004882_4 match to human [H.sapiens]	1149773 IMAGE Consortium	AA674117
544 E	7 021FAMN5	ESTs, Moderately similar to T17273 hypothetical protein DKFZp434C0935.1 - human [H.sapiens]	1148910 IMAGE Consortium	AI644883
545 E	9 021FAMN5	ESTs	1149080 IMAGE Consortium	AA655893
546 E	11 021FAMN5	ESTs	1149166 IMAGE Consortium	AI644932
547 G	1 021FAMN5			
548 G	3 021FAMN5			
549 G	5 021FAMN5	ESTs	1149797 IMAGE Consortium	AA672752

550 G	7 021FAMN5	ESTs, Highly similar to PROTEOLIPID PROTEIN PPA1 [Saccharomyces cerevisiae]	1148950 IMAGE Consortium	AI644887
551 G	9 021FAMN5	ESTs	1149084 IMAGE Consortium	AA655894
552 G	11 021FAMN5	Fus-associated protein with serine-arginine repeats	1149585 IMAGE Consortium	AA655921
553 A	1 021MAMN6	ESTs	1149786 IMAGE Consortium	AA674112
554 A	3 021MAMN6	ADP-ribosylation factor 2	1150391 IMAGE Consortium	AA930772
555 A	5 021MAMN6	syntaxin 8	1150967 IMAGE Consortium	AA791461
556 A	7 021MAMN6	ESTs	1193454 IMAGE Consortium	AA726848
557 A	9 021MAMN6	enhancer of zeste homolog 1 (Drosophila)	1193604 IMAGE Consortium	AA726943
558 A	11 021MAMN6	ESTs	1193719 IMAGE Consortium	AA727065
559 C	1 021MAMN6	integrin alpha L (Cd11a)	1149832 IMAGE Consortium	AA795625
560 C	3 021MAMN6	ESTs	1150466 IMAGE Consortium	AA930278
561 C	5 021MAMN6	amelogenin	1193028 IMAGE Consortium	AA711607
562 C	7 021MAMN6	ESTs, Highly similar to ARGR_HUMAN ARGININE-RICH PROTEIN [H.sapiens]	1193468 IMAGE Consortium	AA726855
563 C	9 021MAMN6	protein tyrosine phosphatase, receptor type, C	1193610 IMAGE Consortium	AA726945
564 C	11 021MAMN6	elongation of very long chain fatty acids (FEN1/Elo2, SUR4/Elo3, yeast)-like 1	1209104 IMAGE Consortium	AA726040
565 E	1 021MAMN6	monoglyceride lipase	1149905 IMAGE Consortium	AA822622
566 E	3 021MAMN6	ESTs, Moderately similar to AF272844_1 11 kDa secreted protein precursor [M.musculus]	1150548 IMAGE Consortium	AA930321
567 E	5 021MAMN6	ESTs, Moderately similar to ATP SYNTHASE COUPLING FACTOR B, MITOCHONDRIAL [Bos taurus]	1193036 IMAGE Consortium	AA711611
568 E	7 021MAMN6	retinoblastoma-like 2	1193484 IMAGE Consortium	AA726872
569 E	9 021MAMN6	ESTs	1193630 IMAGE Consortium	AA727040
570 E	11 021MAMN6	ESTs, Weakly similar to JE0086 SH3-domain binding protein Sab - human [H.sapiens]	1209198 IMAGE Consortium	AA726103
571 G	1 021MAMN6	ESTs	1149953 IMAGE Consortium	AA880056
572 G	3 021MAMN6	ESTs, Moderately similar to AF082535_1 type III multi-pass transmembrane protein [R.norvegicus]	1150609 IMAGE Consortium	AA930453
573 G	5 021MAMN6	ESTs, Weakly similar to serine/threonine kinase [M.musculus]	1193174 IMAGE Consortium	AA718431
574 G	7 021MAMN6	ESTs, Highly similar to 40S RIBOSOMAL PROTEIN S23 [Homo sapiens; Rattus norvegicus]	1193487 IMAGE Consortium	AA726883
575 G	9 021MAMN6	RAN binding protein 9	1193649 IMAGE Consortium	AA726970
576 G	11 021MAMN6	protease, serine, 18	1209218 IMAGE Consortium	AA726122
577 A	1 021TAMN7	activating transcription factor 7 interacting protein	1197184 IMAGE Consortium	AA733711
578 A	3 021TAMN7	ESTs	1198153 IMAGE Consortium	AA725922
579 A	5 021TAMN7	ESTs	1209135 IMAGE Consortium	AA726074
580 A	7 021TAMN7	DNA segment, Chr 8, Wayne State University 108, expressed	1209955 IMAGE Consortium	AA727383
581 A	9 021TAMN7	serine protease inhibitor 2-1	1210259 IMAGE Consortium	AA727334
582 A	11 021TAMN7	ESTs	1223178 IMAGE Consortium	AA734109

583 C	1 021TAMN7	SWI/SNF related, matrix associated, actin dependent regulator of chromatin, subfamily e, member 1	1197251 IMAGE Consortium	AA681910
584 C	3 021TAMN7	ESTs	1198186 IMAGE Consortium	AA725930
585 C	5 021TAMN7	ribosomal protein L10A	1209161 IMAGE Consortium	AA726096
586 C	7 021TAMN7	matrix metalloproteinase 14 (membrane-inserted)	1209994 IMAGE Consortium	AA727448
587 C	9 021TAMN7	kelch-like ECH-associated protein 1	1210306 IMAGE Consortium	AA727391
588 C	11 021TAMN7	delta-6 fatty acid desaturase	1223224 IMAGE Consortium	AA734182
589 E	1 021TAMN7	ESTs, Weakly similar to AF104033_1 MUEL protein [M.musculus]	1197280 IMAGE Consortium	AA681922
590 E	3 021TAMN7	solute carrier family 27 (fatty acid transporter), member 4	1197490 IMAGE Consortium	AA794409
591 E	5 021TAMN7	ESTs, Highly similar to KERATIN, TYPE II CYTOSKELETAL 4 [Homo sapiens]	1209230 IMAGE Consortium	AA726127
592 E	7 021TAMN7	ESTs	1210026 IMAGE Consortium	AA727504
593 E	9 021TAMN7	ESTs, Weakly similar to ECHM_RAT ENOYL-COA HYDRATASE, MITOCHONDRIAL PRECURSOR [R.norvegicus]	1210371 IMAGE Consortium	AA727486
594 E	11 021TAMN7	Tnf receptor associated factor 4	1224079 IMAGE Consortium	AA759867
595 G	1 021TAMN7	rho, GDP dissociation inhibitor (GDI) beta	1198016 IMAGE Consortium	AA727108
596 G	3 021TAMN7	ESTs	1198217 IMAGE Consortium	AA725961
597 G	5 021TAMN7	glioblastoma amplified sequence	1209286 IMAGE Consortium	AA726195
598 G	7 021TAMN7	Mus musculus 11 kDa secreted protein precursor, mRNA, complete cds	1210040 IMAGE Consortium	AA727521
599 G	9 021TAMN7	ESTs	1210499 IMAGE Consortium	AA727700
600 G	11 021TAMN7	ESTs, Highly similar to TALIN [Mus musculus]	1223439 IMAGE Consortium	AA733912
601 A	1 0210AMN8	ESTs	1224367 IMAGE Consortium	AA755771
602 A	3 0210AMN8	UBIQUITIN-LIKE PROTEIN GDX	1211618 IMAGE Consortium	AA655428
603 A	5 0210AMN8	cysteine-rich protein 2	1222894 IMAGE Consortium	AA667906
604 A	7 0210AMN8	differentially expressed in B16F10 2	597004 IMAGE Consortium	AA145454
605 A	9 0210AMN8			
606 A	11 0210AMN8	ESTs	596790 IMAGE Consortium	AA146546
607 C	1 0210AMN8	fibrillin 2	1224369 IMAGE Consortium	AA755772
608 C	3 0210AMN8	ESTs	1211650 IMAGE Consortium	AA656314
609 C	5 0210AMN8			
610 C	7 0210AMN8	ESTs, Weakly similar to AF196304_1 SUMO-1-specific protease [H.sapiens]	597030 IMAGE Consortium	AA145463
611 C	9 0210AMN8	ESTs	597293 IMAGE Consortium	AA146011
612 C	11 0210AMN8	lymphocyte antigen 68	596813 IMAGE Consortium	AA145088
613 E	1 0210AMN8	thioredoxin	1222999 IMAGE Consortium	AA795772
614 E	3 0210AMN8	Mus musculus LAG protein (Lag) mRNA, complete cds	1211698 IMAGE Consortium	AA656394
615 E	5 0210AMN8	ESTs	1211133 IMAGE Consortium	AA655158
616 E	7 0210AMN8	ESTs	597075 IMAGE Consortium	AA145479
617 E	9 0210AMN8			
618 E	11 0210AMN8	ESTs	596847 IMAGE Consortium	AA145097

619 G	1 0210AMN8	ESTs, Highly similar to FIBRINOGEN-LIKE PROTEIN [Homo sapiens]	1223761 IMAGE Consortium	AA815777
620 G	3 0210AMN8			
621 G	5 0210AMN8	ankyrin 1, erythroid	1210461 IMAGE Consortium	AA727627
622 G	7 0210AMN8	Public domain EST	596360 IMAGE Consortium	
623 G	9 0210AMN8	ESTs	596630 IMAGE Consortium	AA146521
624 G	11 0210AMN8	ESTs	596933 IMAGE Consortium	AA145291
625 A	1 0217AMN9	Eph receptor B4	1224545 IMAGE Consortium	AA755959
626 A	3 0217AMN9	bystin-like	1224760 IMAGE Consortium	AA756169
627 A	5 0217AMN9	ESTs, Moderately similar to TRYPSINOGEN [Bos taurus]	1227211 IMAGE Consortium	AA738672
628 A	7 0217AMN9	fragile X mental retardation gene 2, autosomal homolog	1227477 IMAGE Consortium	AA738780
629 A	9 0217AMN9	ESTs	1227763 IMAGE Consortium	AA760167
630 A	11 0217AMN9	Public domain EST	1228163 IMAGE Consortium	
631 C	1 0217AMN9	ESTs, Moderately similar to polyamine modulated factor-1 [H.sapiens]	1224568 IMAGE Consortium	AA756009
632 C	3 0217AMN9	vacuolar adenosine triphosphatase subunit C	1224815 IMAGE Consortium	AA789873
633 C	5 0217AMN9	submaxillary gland androgen regulated protein 1	1227261 IMAGE Consortium	AA756749
634 C	7 0217AMN9	hypothetical protein, clone 2-24	1227534 IMAGE Consortium	AA738910
635 C	9 0217AMN9	osmotic stress protein 94 kDa	1227792 IMAGE Consortium	AA760222
636 C	11 0217AMN9	keratin complex 1, acidic, gene 10	1228170 IMAGE Consortium	AA760250
637 E	1 0217AMN9	ESTs, Weakly similar to A41539 kinesin light chain A - rat [R.norvegicus]	1224608 IMAGE Consortium	AA756083
638 E	3 0217AMN9	peroxiredoxin 2	1227243 IMAGE Consortium	AA756741
639 E	5 0217AMN9	ESTs, Moderately similar to Y254_HUMAN HYPOTHETICAL PROTEIN KIAA0254 [H.sapiens]	1227311 IMAGE Consortium	AA756787
640 E	7 0217AMN9	ESTs	1227542 IMAGE Consortium	AI604316
641 E	9 0217AMN9	ESTs, Weakly similar to NUCLEAR AND CYTOPLASMIC POLYADENYLATED RNA-BINDING PROTEIN PUB1 [Saccharomyces cerevisiae]	1227795 IMAGE Consortium	AA760233
642 E	11 0217AMN9	ATPase, class VI, type 11A	1228225 IMAGE Consortium	AA760346
643 G	1 0217AMN9	ESTs	1224633 IMAGE Consortium	AA756118
644 G	3 0217AMN9	complement receptor related protein	1227295 IMAGE Consortium	AA756782
645 G	5 0217AMN9	beta-glucuronidase structural	1227326 IMAGE Consortium	AA760002
646 G	7 0217AMN9	ESTs	1227562 IMAGE Consortium	AA738934
647 G	9 0217AMN9			
648 G	11 0217AMN9	keratin complex 1, acidic, gene 15	1228244 IMAGE Consortium	AA760344
649 A	1 021RAMNA	ESTs, Weakly similar to glycine tyrosine-rich hair keratin protein [M.musculus]	1229526 IMAGE Consortium	AA763224
650 A	3 021RAMNA	ESTs	1230134 IMAGE Consortium	AA798936
651 A	5 021RAMNA	glioma-amplified sequence-41	1259959 IMAGE Consortium	AA855587
652 A	7 021RAMNA	ESTs, Weakly similar to /prediction	1261983 IMAGE Consortium	AA874142
653 A	9 021RAMNA	bone gamma-carboxyglutamate protein, related sequence 1	1261446 IMAGE Consortium	AA856059

654 A	11 021RAMNA	Mus musculus brain cDNA, clone MNCb-0467, similar to Homo sapiens cDNA FLJ10078, clone HEMBA1001869	1260282 IMAGE Consortium	AA855257
655 C	1 021RAMNA	ESTs, Highly similar to hADA2	1229644 IMAGE Consortium	AA763040
656 C	3 021RAMNA	caveolin, caveolae protein, 22 kDa	1230143 IMAGE Consortium	AA798948
657 C	5 021RAMNA	baculoviral IAP repeat-containing 2	1260101 IMAGE Consortium	AA840057
658 C	7 021RAMNA	ESTs	1261054 IMAGE Consortium	AA838855
659 C	9 021RAMNA	EST AA409339	1261464 IMAGE Consortium	AI465679
660 C	11 021RAMNA	Bloom syndrome homolog (human)	1260305 IMAGE Consortium	AA855289
661 E	1 021RAMNA	Mus musculus mRNA for stretch responsive muscle (X-chromosome) protein (Srmx gene)	1229575 IMAGE Consortium	AA763276
662 E	3 021RAMNA	transformation related protein 63	1230154 IMAGE Consortium	AA798953
663 E	5 021RAMNA	ESTs, Moderately similar to PMS1_HUMAN PMS1 PROTEIN HOMOLOG 1 [H.sapiens]	1242972 IMAGE Consortium	AA790369
664 E	7 021RAMNA	prostaglandin E receptor 1 (subtype EP1), 42kD	1261169 IMAGE Consortium	AA838963
665 E	9 021RAMNA	adaptor-related protein complex AP-1, mu subunit 1	1260811 IMAGE Consortium	AA855853
666 E	11 021RAMNA	RAD23a homolog (S. cerevisiae)	1260313 IMAGE Consortium	AA855304
667 G	1 021RAMNA	ESTs	1229655 IMAGE Consortium	AA763056
668 G	3 021RAMNA	ESTs, Weakly similar to DnaJ-like protein [M.musculus]	1230174 IMAGE Consortium	AA798972
669 G	5 021RAMNA	ESTs	1261521 IMAGE Consortium	AI465691
670 G	7 021RAMNA	nucleosome assembly protein 1-like 1	1262006 IMAGE Consortium	AA874061
671 G	9 021RAMNA	AT motif binding factor 1	1260868 IMAGE Consortium	AA840196
672 G	11 021RAMNA	ribophorin II	1260364 IMAGE Consortium	AI463951
673 A	1 021YAMNB	ESTs, Weakly similar to RGSE_MOUSE REGULATOR OF G-PROTEIN SIGNALING 14 [M.musculus]	1260708 IMAGE Consortium	AA855540
674 A	3 021YAMNB	histocompatibility 2, K region	1262564 IMAGE Consortium	AA839197
675 A	5 021YAMNB	ESTs	1277084 IMAGE Consortium	AA880938
676 A	7 021YAMNB	EST AI426782	1277592 IMAGE Consortium	AA880285
677 A	9 021YAMNB	ESTs	329617 IMAGE Consortium	AI451754
678 A	11 021YAMNB	dynein, cytoplasmic, intermediate chain 1	331202 IMAGE Consortium	AI390870
679 C	1 021YAMNB	EST	1260772 IMAGE Consortium	AI562092
680 C	3 021YAMNB	ESTs	1262752 IMAGE Consortium	AI465762
681 C	5 021YAMNB	ribosomal protein, large, P1	1277168 IMAGE Consortium	AA880995
682 C	7 021YAMNB	IMMUNOGLOBULIN J CHAIN PRECURSOR	1277614 IMAGE Consortium	AA880295
683 C	9 021YAMNB	Ngfi-A binding protein 2	329956 IMAGE Consortium	AI451768
684 C	11 021YAMNB	ESTs, Moderately similar to ADENYLATE CYCLASE, TYPE IV [Rattus norvegicus]	331379 IMAGE Consortium	AI327512
685 E	1 021YAMNB	max binding protein	1262246 IMAGE Consortium	AA920158
686 E	3 021YAMNB	ribosomal protein, mitochondrial, S25	1262774 IMAGE Consortium	AI036813
687 E	5 021YAMNB	ESTs	1277193 IMAGE Consortium	AA881025
688 E	7 021YAMNB	ESTs, Highly similar to HFE_MOUSE HEREDITARY HAEMOCHROMATOSIS PROTEIN HOMOLOG PRECURSOR [M.musculus]	1276978 IMAGE Consortium	AA880866
689 E	9 021YAMNB	prostate specific ets transcription factor	330181 IMAGE Consortium	AI451780

690 E	11 021YAMNB	ESTs	330782 IMAGE Consortium	AI390840
691 G	1 021YAMNB	ESTs	1262253 IMAGE Consortium	AA920171
692 G	3 021YAMNB	RAS-related C3 botulinum substrate 2	1276777 IMAGE Consortium	AA880744
693 G	5 021YAMNB	fatty acid binding protein 1, liver	1277312 IMAGE Consortium	AI466451
694 G	7 021YAMNB	aminolevulinic acid synthase 2, erythroid	1277758 IMAGE Consortium	AA880550
695 G	9 021YAMNB	ESTs	331080 IMAGE Consortium	AI390861
696 G	11 021YAMNB	ubiquitin carboxy-terminal hydrolase L1	331582 IMAGE Consortium	AI390890
697 A	1 0215AMNC	growth arrest specific 1	317615 IMAGE Consortium	W13100
698 A	3 0215AMNC	3-monooxygenase/tryptophan 5-monooxygenase activation protein, gamma polypeptide	318970 IMAGE Consortium	W34479
699 A	5 0215AMNC			
700 A	7 0215AMNC			
701 A	9 0215AMNC			
702 A	11 0215AMNC	elongation of very long chain fatty acids (FEN1/Elo2, SUR4/Elo3, yeast)-like 3	1280273 IMAGE Consortium	AA798103
703 C	1 0215AMNC	G protein-coupled receptor, family C, group 5, member B	318497 IMAGE Consortium	AI536352
704 C	3 0215AMNC			
705 C	5 0215AMNC			
706 C	7 0215AMNC			
707 C	9 0215AMNC			
708 C	11 0215AMNC	ESTs, Highly similar to INOSITOL 1,4,5-TRISPHOSPHATE-BINDING PROTEIN TYPE 1 RECEPTOR [Rattus norvegicus]	1280539 IMAGE Consortium	AA789907
709 E	1 0215AMNC	ESTs	317873 IMAGE Consortium	W13074
710 E	3 0215AMNC			
711 E	5 0215AMNC			
712 E	7 0215AMNC			
713 E	9 0215AMNC			
714 E	11 0215AMNC	ESTs, Highly similar to DEOXYHYPUSINE SYNTHASE [Homo sapiens]	1280554 IMAGE Consortium	AA789915
715 G	1 0215AMNC	ESTs, Weakly similar to DHBK_MOUSE PUTATIVE STEROID DEHYDROGENASE KIK-I□ [M.musculus]	317906 IMAGE Consortium	W12688
716 G	3 0215AMNC			
717 G	5 0215AMNC			
718 G	7 0215AMNC			
719 G	9 0215AMNC			
720 G	11 0215AMNC	ESTs, Weakly similar to CG14966 gene product [D.melanogaster]	1280702 IMAGE Consortium	AA790583
721 A	1 021CAMND	ESTs, Moderately similar to KIAA0385 [H.sapiens]	1278954 IMAGE Consortium	AA896621
722 A	3 021CAMND	ketoheokinase	1278411 IMAGE Consortium	AA895199
723 A	5 021CAMND	ESTs	1277900 IMAGE Consortium	AA882250
724 A	7 021CAMND	secretory leukocyte protease inhibitor	1367299 IMAGE Consortium	AI020539
725 A	9 021CAMND	ESTs	598820 IMAGE Consortium	AA163047



726 A	11	021CAMND	ESTs, Weakly similar to INTERFERON-INDUCIBLE PROTEIN 9-27 [Homo sapiens]	597616	IMAGE Consortium	AI386420
727 C	1	021CAMND	ESTs, Weakly similar to complement receptor CR2 [M.musculus]	1278119	IMAGE Consortium	AA882450
728 C	3	021CAMND	cyclin T1	1279179	IMAGE Consortium	AA981354
729 C	5	021CAMND	ribonuclease H1	1278705	IMAGE Consortium	AA895522
730 C	7	021CAMND	signal recognition particle 54 kDa	1350086	IMAGE Consortium	AA982259
731 C	9	021CAMND	ESTs	598853	IMAGE Consortium	AI592195
732 C	11	021CAMND	ESTs	598355	IMAGE Consortium	AI592115
733 E	1	021CAMND	ESTs, Weakly similar to VILLIN [Gallus gallus]	1278245	IMAGE Consortium	AA880830
734 E	3	021CAMND	Mus musculus DNA cytosine methyltransferase mRNA	1279181	IMAGE Consortium	AA981355
735 E	5	021CAMND	myeloid differentiation primary response gene 118	1278753	IMAGE Consortium	AA895566
736 E	7	021CAMND	CCR4-NOT transcription complex, subunit 7	1350205	IMAGE Consortium	AA967217
737 E	9	021CAMND	ESTs	597143	IMAGE Consortium	AA139038
738 E	11	021CAMND	ESTs	598359	IMAGE Consortium	AI592116
739 G	1	021CAMND	hippocalcin-like 1	1278289	IMAGE Consortium	AA882018
740 G	3	021CAMND	endosulfine alpha	1278511	IMAGE Consortium	AA895259
741 G	5	021CAMND	glycogen synthase 3, brain	1278854	IMAGE Consortium	AA895620
742 G	7	021CAMND	gelsolin	1350327	IMAGE Consortium	AA980357
743 G	9	021CAMND	ESTs	598099	IMAGE Consortium	AA265247
744 G	11	021CAMND	ESTs, Weakly similar to endocrine regulator [H.sapiens]	597657	IMAGE Consortium	AI592036
745 A	1	021JAMNE	ESTs, Weakly similar to PROTEIN A55 [Vaccinia virus (strain copenhagen)]	597726	IMAGE Consortium	AI386424
746 A	3	021JAMNE	ESTs	597916	IMAGE Consortium	AA163793
747 A	5	021JAMNE	integral membrane protein 2	775197	IMAGE Consortium	AA387218
748 A	7	021JAMNE				
749 A	9	021JAMNE	DNA segment, Chr 9, Wayne State University 10, expressed	775479	IMAGE Consortium	AA388215
750 A	11	021JAMNE	zinc finger protein 37	790148	IMAGE Consortium	AA388202
751 C	1	021JAMNE	lipase, hormone sensitive	597754	IMAGE Consortium	AA154113
752 C	3	021JAMNE	ESTs, Weakly similar to ADT2_MOUSE ADP,ATP CARRIER PROTEIN, FIBROBLAST ISOFORM [M.musculus]	775068	IMAGE Consortium	AA387364
753 C	5	021JAMNE	ESTs	775207	IMAGE Consortium	AA386845
754 C	7	021JAMNE	ESTs, Weakly similar to G protein beta1 subunit [R.norvegicus]	775390	IMAGE Consortium	AA386911
755 C	9	021JAMNE	ESTs, Moderately similar to KIAA1313 protein [H.sapiens]	775518	IMAGE Consortium	AA388290
756 C	11	021JAMNE	ESTs, Weakly similar to CD63_MOUSE CD63 ANTIGEN [M.musculus]	805008	IMAGE Consortium	AA397283
757 E	1	021JAMNE	ESTs	597774	IMAGE Consortium	AA164064
758 E	3	021JAMNE	Myg1 protein	775077	IMAGE Consortium	AA387191
759 E	5	021JAMNE	ESTs	775218	IMAGE Consortium	AA387219
760 E	7	021JAMNE	ESTs	775398	IMAGE Consortium	AA387252

761 E	9 021JAMNE	procollagen-proline, 2-oxoglutarate 4-dioxygenase (proline 4-hydroxylase), alpha II polypeptide	775550 IMAGE Consortium	AA388310
762 E	11 021JAMNE	ESTs	805070 IMAGE Consortium	AA397221
763 G	1 021JAMNE	ESTs	598521 IMAGE Consortium	AA161853
764 G	3 021JAMNE	v-ral simian leukemia viral oncogene homolog B (ras related)	775114 IMAGE Consortium	AA386860
765 G	5 021JAMNE	calsequestrin 2	775226 IMAGE Consortium	AA386846
766 G	7 021JAMNE	ESTs	775401 IMAGE Consortium	AA387338
767 G	9 021JAMNE	glial cell line derived neurotrophic factor family receptor alpha 1	774901 IMAGE Consortium	AA387098
768 G	11 021JAMNE	sterol O-acyltransferase 1	805142 IMAGE Consortium	AI551701
769 A	1 021QAMNF	profilin 2	804519 IMAGE Consortium	AA474438
770 A	3 021QAMNF	fos-like antigen 2	805493 IMAGE Consortium	AI551662
771 A	5 021QAMNF	regucalcin	790065 IMAGE Consortium	AA388066
772 A	7 021QAMNF	ESTs	819869 IMAGE Consortium	AA437952
773 A	9 021QAMNF	ESTs, Weakly similar to ZK1058.5 [C.elegans]	820034 IMAGE Consortium	AI551578
774 A	11 021QAMNF	ESTs	805658 IMAGE Consortium	AA399839
775 C	1 021QAMNF	ESTs, Highly similar to KIAA1435 protein [H.sapiens]	804535 IMAGE Consortium	AA474360
776 C	3 021QAMNF	ESTs, Moderately similar to RS20_HUMAN 40S RIBOSOMAL PROTEIN S20 [R.norvegicus]	804950 IMAGE Consortium	AA397345
777 C	5 021QAMNF	ESTs, Moderately similar to KIAA0966 protein [H.sapiens]	805834 IMAGE Consortium	AI604703
778 C	7 021QAMNF	ESTs	819892 IMAGE Consortium	AI551546
779 C	9 021QAMNF			
780 C	11 021QAMNF	ESTs, Moderately similar to PEXD_HUMAN PEROXISOMAL MEMBRANE PROTEIN PEX13 [H.sapiens]	805739 IMAGE Consortium	AA437649
781 E	1 021QAMNF	ESTs	804608 IMAGE Consortium	AA474723
782 E	3 021QAMNF	ESTs	804982 IMAGE Consortium	AA397170
783 E	5 021QAMNF	ESTs	805868 IMAGE Consortium	AA399859
784 E	7 021QAMNF	ESTs	819896 IMAGE Consortium	AA437961
785 E	9 021QAMNF	ESTs, Weakly similar to CALB_MOUSE CALCINEURIN B SUBUNIT ISOFORM 1 [M.musculus]	805145 IMAGE Consortium	AA473114
786 E	11 021QAMNF	ESTs	805745 IMAGE Consortium	AA437739
787 G	1 021QAMNF	ESTs	805369 IMAGE Consortium	AA473386
788 G	3 021QAMNF	wingless-related MMTV integration site 5A	789868 IMAGE Consortium	AA388110
789 G	5 021QAMNF	faciogenital dysplasia homolog	819721 IMAGE Consortium	AA437519
790 G	7 021QAMNF	proprotein convertase subtilisin/kexin type 7	819916 IMAGE Consortium	AA437757
791 G	9 021QAMNF	Kruppel-like factor 9	805390 IMAGE Consortium	AI551651
792 G	11 021QAMNF	Huntington disease gene homolog	805796 IMAGE Consortium	AA437749
793 A	1 021XAMNG			
794 A	3 021XAMNG	cysteine rich protein 61	834600 IMAGE Consortium	AA466852
795 A	5 021XAMNG	glutathione S-transferase, pi 2	819851 IMAGE Consortium	AA437941
796 A	7 021XAMNG	Dlxin-1	819975 IMAGE Consortium	AA437797

797 A	9 021XAMNG	ESTs	894288 IMAGE Consortium	AA517339
798 A	11 021XAMNG	ESTs, Moderately similar to serine/threonine kinase [M.musculus]	894414 IMAGE Consortium	AA517489
799 C	1 021XAMNG	ESTs	833592 IMAGE Consortium	AI596493
800 C	3 021XAMNG	ESTs	834637 IMAGE Consortium	AA466871
801 C	5 021XAMNG	ESTs	819854 IMAGE Consortium	AA437939
802 C	7 021XAMNG	ESTs	820001 IMAGE Consortium	AA403754
803 C	9 021XAMNG	ESTs, Highly similar to unnamed protein product [H.sapiens]	894305 IMAGE Consortium	AA517358
804 C	11 021XAMNG	transcription factor Dp 1	894434 IMAGE Consortium	AA517509
805 E	1 021XAMNG	kidney predominant protein NCU-G1	833698 IMAGE Consortium	AA466720
806 E	3 021XAMNG	ESTs, Weakly similar to AF077034_1 HSPC010 [H.sapiens]	819737 IMAGE Consortium	AA437663
807 E	5 021XAMNG	ring finger protein 11	819907 IMAGE Consortium	AA437704
808 E	7 021XAMNG	ESTs	820026 IMAGE Consortium	AI551573
809 E	9 021XAMNG	ESTs	894310 IMAGE Consortium	AA517350
810 E	11 021XAMNG	ESTs, Highly similar to TRANSCRIPTION FACTOR BTF3 [Homo sapiens]	894451 IMAGE Consortium	AA517524
811 G	1 021XAMNG	ESTs	833799 IMAGE Consortium	AI596504
812 G	3 021XAMNG	ESTs	819760 IMAGE Consortium	AI595386
813 G	5 021XAMNG	proteasome (prosome, macropain) 26S subunit, non-ATPase, 7	819924 IMAGE Consortium	AA437761
814 G	7 021XAMNG	DNA segment, Chr 4, Wayne State University 43, expressed	820069 IMAGE Consortium	AA444219
815 G	9 021XAMNG	ESTs	894338 IMAGE Consortium	AI613728
816 G	11 021XAMNG	ESTs, Weakly similar to AF159856_1 N system amino acids transporter NAT-1 [M.musculus]	894468 IMAGE Consortium	AA517531
817 A	1 0214AMNH	high mobility group box 1	893901 IMAGE Consortium	AA517299
818 A	3 0214AMNH	ESTs	894097 IMAGE Consortium	AI613698
819 A	5 0214AMNH	heterogeneous nuclear ribonucleoprotein H1	893834 IMAGE Consortium	AA517175
820 A	7 0214AMNH	ESTs	934094 IMAGE Consortium	AA544576
821 A	9 0214AMNH	Mus musculus mRNA for microtubule associated protein 44 (Mtap44 gene)	933769 IMAGE Consortium	AA545696
822 A	11 0214AMNH	phosphatidylcholine transfer protein-like	934263 IMAGE Consortium	AA543948
823 C	1 0214AMNH	ESTs	932730 IMAGE Consortium	AI596372
824 C	3 0214AMNH	ESTs	835267 IMAGE Consortium	AI551720
825 C	5 0214AMNH	ESTs	893941 IMAGE Consortium	AA517148
826 C	7 0214AMNH	ESTs, Highly similar to UDP-Gal:betaGlcNAc beta 1,3-galactosyltransferase-III [M.musculus]	934130 IMAGE Consortium	AA543834
827 C	9 0214AMNH	Ewing sarcoma homolog	933800 IMAGE Consortium	AA545706
828 C	11 0214AMNH	ESTs	934268 IMAGE Consortium	AA543943
829 E	1 0214AMNH	ESTs, Highly similar to mm-Mago [M.musculus]	932847 IMAGE Consortium	AA538564
830 E	3 0214AMNH	ESTs	892720 IMAGE Consortium	AA501174

831 E	5 0214AMNH	ESTs, Weakly similar to possible ribosomal protein	893989 IMAGE Consortium	AA516658
832 E	7 0214AMNH	myosin Ic	934158 IMAGE Consortium	AA544749
833 E	9 0214AMNH	ribosomal protein L29	933877 IMAGE Consortium	AA544402
834 E	11 0214AMNH	ESTs, Highly similar to KIAA0186 [H.sapiens]	934291 IMAGE Consortium	AA543968
835 G	1 0214AMNH	ESTs, Moderately similar to PLATELET ENDOTHELIAL CELL ADHESION MOLECULE PRECURSOR [Mus musculus]	932976 IMAGE Consortium	AA538188
836 G	3 0214AMNH	zinc finger protein 296	894195 IMAGE Consortium	AA517061
837 G	5 0214AMNH	ESTs	836991 IMAGE Consortium	AA432829
838 G	7 0214AMNH	ESTs	934230 IMAGE Consortium	AA544681
839 G	9 0214AMNH	ESTs, Weakly similar to NUML_MOUSE NADH-UBIQUINONE OXIDOREDUCTASE MLRQ SUBUNIT [M.musculus]	933899 IMAGE Consortium	AA544416
840 G	11 0214AMNH	ESTs, Highly similar to nuclear protein SA-1 [M.musculus]	934408 IMAGE Consortium	AA545027
841 A	1 021BAMNI	ESTs, Moderately similar to unnamed protein product [H.sapiens]	934595 IMAGE Consortium	AA536772
842 A	3 021BAMNI	ESTs, Highly similar to OXRTGU L-gulonolactone oxidase [R.norvegicus]	934807 IMAGE Consortium	AA617321
843 A	5 021BAMNI	ESTs	598930 IMAGE Consortium	AA168619
844 A	7 021BAMNI	transmembrane protein 6	617435 IMAGE Consortium	AA174396
845 A	9 021BAMNI	ESTs, Highly similar to p71 [R.norvegicus]	599123 IMAGE Consortium	AA154730
846 A	11 021BAMNI	ESTs, Highly similar to KE03 protein [H.sapiens]	947464 IMAGE Consortium	AI558180
847 C	1 021BAMNI	ESTs	934627 IMAGE Consortium	AA536789
848 C	3 021BAMNI	programmed cell death 4	934824 IMAGE Consortium	AA617346
849 C	5 021BAMNI	ESTs, Weakly similar to KIAA0403 [H.sapiens]	598954 IMAGE Consortium	AI592210
850 C	7 021BAMNI	hematopoietically expressed homeobox	618210 IMAGE Consortium	AA170521
851 C	9 021BAMNI	ESTs	599134 IMAGE Consortium	AI592238
852 C	11 021BAMNI	ESTs, Weakly similar to hypothetical protein [H.sapiens]	962810 IMAGE Consortium	AI504260
853 E	1 021BAMNI	tyrosine 3-monooxygenase/tryptophan 5-monooxygenase activation protein, zeta polypeptide	934645 IMAGE Consortium	AA536841
854 E	3 021BAMNI	ESTs	934833 IMAGE Consortium	AA617355
855 E	5 021BAMNI	ESTs	617165 IMAGE Consortium	AI640116
856 E	7 021BAMNI	ESTs	618283 IMAGE Consortium	AA170596
857 E	9 021BAMNI	ESTs	946331 IMAGE Consortium	AI550961
858 E	11 021BAMNI	ESTs, Highly similar to I54388 LZTR-1 - human [H.sapiens]	947550 IMAGE Consortium	AA589209
859 G	1 021BAMNI			
860 G	3 021BAMNI	beta-galactosidase complex	934838 IMAGE Consortium	AA617351
861 G	5 021BAMNI	ESTs	617234 IMAGE Consortium	AA174675
862 G	7 021BAMNI	selected mouse cDNA on the Y	617564 IMAGE Consortium	AA170509
863 G	9 021BAMNI	ESTs	947236 IMAGE Consortium	AA592010
864 G	11 021BAMNI	ESTs	962905 IMAGE Consortium	AI597204
865 A	1 021IAMNJ	ESTs, Weakly similar to Yil083cp [S.cerevisiae]	946255 IMAGE Consortium	AA591751

866 A	3 021IAMNJ	ESTs	935134 IMAGE Consortium	AA616984
867 A	5 021IAMNJ	ESTs, Weakly similar to C1q-related factor [M.musculus]	945034 IMAGE Consortium	AA537246
868 A	7 021IAMNJ	ESTs, Weakly similar to ZINC FINGER PROTEIN 91 [Homo sapiens]	945493 IMAGE Consortium	AI585415
869 A	9 021IAMNJ	Tnf receptor-associated factor 2	962923 IMAGE Consortium	AA619651
870 A	11 021IAMNJ	ceroid-lipofuscinosis, neuronal 8	976795 IMAGE Consortium	AA563326
871 C	1 021IAMNJ	poly (ADP-ribose) glycohydrolase	945431 IMAGE Consortium	AA546790
872 C	3 021IAMNJ	tubulin alpha 2	935171 IMAGE Consortium	AA617612
873 C	5 021IAMNJ	ESTs	945073 IMAGE Consortium	AA537336
874 C	7 021IAMNJ	acid sphingomyelinase-like phosphodiesterase 3a	945529 IMAGE Consortium	AA545949
875 C	9 021IAMNJ	ESTs, Weakly similar to AF151800_1 CGI-41 protein [H.sapiens]	962947 IMAGE Consortium	AA619658
876 C	11 021IAMNJ	RAN binding protein 9	976830 IMAGE Consortium	AA563437
877 E	1 021IAMNJ	ESTs	945610 IMAGE Consortium	AI585420
878 E	3 021IAMNJ	ESTs, Moderately similar to /prediction	935180 IMAGE Consortium	AA616995
879 E	5 021IAMNJ	ESTs	934941 IMAGE Consortium	AA617547
880 E	7 021IAMNJ	DNA segment, Chr 6, Wayne State University 163, expressed	945548 IMAGE Consortium	AI585416
881 E	9 021IAMNJ	calpain 5	977024 IMAGE Consortium	AI645410
882 E	11 021IAMNJ	ribosomal protein S17	976859 IMAGE Consortium	AA563445
883 G	1 021IAMNJ	mitogen activated protein kinase kinase kinase 12	945650 IMAGE Consortium	AA546328
884 G	3 021IAMNJ	ESTs	935197 IMAGE Consortium	AI558252
885 G	5 021IAMNJ	EST	945114 IMAGE Consortium	AI643567
886 G	7 021IAMNJ	ESTs, Highly similar to PYRROLINE-5-CARBOXYLATE REDUCTASE [Homo sapiens]	962678 IMAGE Consortium	AA562429
887 G	9 021IAMNJ	mini chromosome maintenance deficient (S. cerevisiae)	977156 IMAGE Consortium	AI645415
888 G	11 021IAMNJ	Rab acceptor 1 (prenylated)	976876 IMAGE Consortium	AA562543
889 A	1 021PAMNK	ESTs, Weakly similar to JAK3_MOUSE TYROSINE-PROTEIN KINASE JAK3 [M.musculus]	976924 IMAGE Consortium	AA562690
890 A	3 021PAMNK	histone deacetylase 2	977792 IMAGE Consortium	AA562726
891 A	5 021PAMNK	ESTs, Highly similar to HSPC003 [H.sapiens]	988633 IMAGE Consortium	AA638280
892 A	7 021PAMNK	ESTs	988944 IMAGE Consortium	AA591079
893 A	9 021PAMNK	neuraminidase 1	988383 IMAGE Consortium	AA571324
894 A	11 021PAMNK	v-ral simian leukemia viral oncogene homolog A (ras related)	992291 IMAGE Consortium	AA589328
895 C	1 021PAMNK	malate dehydrogenase, mitochondrial	976934 IMAGE Consortium	AA562693
896 C	3 021PAMNK	ESTs, Moderately similar to AF161401_1 HSPC283 [H.sapiens]	988415 IMAGE Consortium	AA571746
897 C	5 021PAMNK	HGF-regulated tyrosine kinase substrate	989384 IMAGE Consortium	AA571461
898 C	7 021PAMNK	ESTs, Moderately similar to KIAA0947 protein [H.sapiens]	988962 IMAGE Consortium	AA591085
899 C	9 021PAMNK	DNA segment, Chr 17, Wayne State University 76, expressed	991029 IMAGE Consortium	AA590197
900 C	11 021PAMNK	ESTs, Highly similar to PRE-MRNA SPLICING FACTOR	992355 IMAGE Consortium	AI588075

901 E	1 021PAMNK	PRP9 [ <i>Saccharomyces cerevisiae</i> ] ESTs, Moderately similar to ER29_RAT ENDOPLASMIC RETICULUM PROTEIN ERP29 PRECURSOR [ <i>R.norvegicus</i> ]	976936 IMAGE Consortium	AA562694
902 E	3 021PAMNK	ESTs, Highly similar to Similar to Plasmodium falciparum glutamic acid-rich protein precursor [ <i>H.sapiens</i> ]	988438 IMAGE Consortium	AA571753
903 E	5 021PAMNK	ESTs	988698 IMAGE Consortium	AA638311
904 E	7 021PAMNK	calcium modulating ligand	978061 IMAGE Consortium	AA563401
905 E	9 021PAMNK	DNA segment, Chr 13, Wayne State University 156, expressed	992031 IMAGE Consortium	AA574954
906 E	11 021PAMNK	ESTs, Moderately similar to no similarities to reported gene products [ <i>H.sapiens</i> ]	992385 IMAGE Consortium	AI588077
907 G	1 021PAMNK	5-azacytidine induced gene 2	976964 IMAGE Consortium	AA562713
908 G	3 021PAMNK	ESTs, Weakly similar to putative RNA helicase [ <i>M.musculus</i> ]	988484 IMAGE Consortium	AA571860
909 G	5 021PAMNK	cytokine inducible SH2-containing protein 3	988726 IMAGE Consortium	AA638328
910 G	7 021PAMNK	ESTs	978133 IMAGE Consortium	AA619988
911 G	9 021PAMNK	ESTs	992105 IMAGE Consortium	AA575047
912 G	11 021PAMNK	DNA segment, Chr 11, ERATO Doi 707, expressed	992391 IMAGE Consortium	AA574968
913 A	1 021WAMNL	early development regulator1 (homolog of polyhomeotic 1)	989179 IMAGE Consortium	AA571780
914 A	3 021WAMNL	ubiquitin-conjugating enzyme 4	989977 IMAGE Consortium	AA591129
915 A	5 021WAMNL	calpain 7	990919 IMAGE Consortium	AA591282
916 A	7 021WAMNL	ESTs	1004226 IMAGE Consortium	AA600629
917 A	9 021WAMNL	activating transcription factor 2	1004418 IMAGE Consortium	AA600465
918 A	11 021WAMNL	ESTs, Weakly similar to STRI_RAT STRIATIN□ [ <i>R.norvegicus</i> ]	1004713 IMAGE Consortium	AA606562
919 C	1 021WAMNL	DNA segment, Chr 5, ERATO Doi 249, expressed	989577 IMAGE Consortium	AA571562
920 C	3 021WAMNL	ESTs	990121 IMAGE Consortium	AA591207
921 C	5 021WAMNL	ESTs, Highly similar to Nop10p [ <i>H.sapiens</i> ]	1004045 IMAGE Consortium	AI606375
922 C	7 021WAMNL	expressed in non-metastatic cells 1, protein (NM23A) (nucleoside diphosphate kinase)	1004275 IMAGE Consortium	AA600663
923 C	9 021WAMNL	ESTs	1004453 IMAGE Consortium	AA791979
924 C	11 021WAMNL	ESTs, Highly similar to VACUOLAR ATP SYNTHASE 14 KD SUBUNIT [ <i>Manduca sexta</i> ]	1004727 IMAGE Consortium	AA606567
925 E	1 021WAMNL	ESTs	989642 IMAGE Consortium	AA571638
926 E	3 021WAMNL	fatty acid binding protein 5, epidermal	990152 IMAGE Consortium	AA591215
927 E	5 021WAMNL	pale ear	1004085 IMAGE Consortium	AA600571
928 E	7 021WAMNL	heterogeneous nuclear ribonucleoprotein D-like	1004291 IMAGE Consortium	AA600672
929 E	9 021WAMNL	mini chromosome maintenance deficient 2 ( <i>S. cerevisiae</i> )	1004502 IMAGE Consortium	AA792000
930 E	11 021WAMNL	ESTs	1004748 IMAGE Consortium	AI594234
931 G	1 021WAMNL	ESTs, Moderately similar to S12207 hypothetical protein [ <i>M.musculus</i> ]	989703 IMAGE Consortium	AA571674
932 G	3 021WAMNL	ESTs	990200 IMAGE Consortium	AI604501

933 G	5 021WAMNL	ESTs, Highly similar to NADH-UBIQUINONE OXIDOREDUCTASE B15 SUBUNIT [Bos taurus]	1004099 IMAGE Consortium	AA600575
934 G	7 021WAMNL	myeloblastosis oncogene-like 2	1004330 IMAGE Consortium	AA600430
935 G	9 021WAMNL	phosphoenolpyruvate carboxykinase 1, cytosolic	1004618 IMAGE Consortium	AI606392
936 G	11 021WAMNL	ESTs	992423 IMAGE Consortium	AA574986
937 A	1 0213AMNM	silica-induced gene 81	1005022 IMAGE Consortium	AA608173
938 A	3 0213AMNM	DNA segment, Chr 17, ERATO Doi 441, expressed	1005109 IMAGE Consortium	AI594263
939 A	5 0213AMNM	ESTs	1004755 IMAGE Consortium	AA606582
940 A	7 0213AMNM	CTP synthetase homolog	1004870 IMAGE Consortium	AA606974
941 A	9 0213AMNM	proteasome (prosome, macropain) subunit, beta type 4	1004953 IMAGE Consortium	AA607199
942 A	11 0213AMNM	poly A binding protein, cytoplasmic 1	1006233 IMAGE Consortium	AI594308
943 C	1 0213AMNM	ESTs	1005031 IMAGE Consortium	AA608191
944 C	3 0213AMNM	ESTs	1005128 IMAGE Consortium	AA606601
945 C	5 0213AMNM	DNA segment, Chr 11, ERATO Doi 153, expressed	1004783 IMAGE Consortium	AA606707
946 C	7 0213AMNM	ESTs, Highly similar to AF176072_1 SMN-interacting protein 1 [R.norvegicus]	1004885 IMAGE Consortium	AI594245
947 C	9 0213AMNM	Y box protein 2	1004957 IMAGE Consortium	AA607201
948 C	11 0213AMNM	calcium binding protein, intestinal	1006236 IMAGE Consortium	AA624165
949 E	1 0213AMNM	ESTs	1005048 IMAGE Consortium	AA608314
950 E	3 0213AMNM	ESTs	1005130 IMAGE Consortium	AA606602
951 E	5 0213AMNM	DNA segment, Chr 14, University of California at Los Angeles 3	1004804 IMAGE Consortium	AA606714
952 E	7 0213AMNM	BTB and CNC homology 2	1004889 IMAGE Consortium	AA606992
953 E	9 0213AMNM	ESTs, Weakly similar to I49642 estrogen-responsive finger protein - mouse [M.musculus]	1004968 IMAGE Consortium	AA607195
954 E	11 0213AMNM	dodecenoyl-Coenzyme A delta isomerase (3,2 trans-enoyl-Coenzyme A isomerase)	1005599 IMAGE Consortium	AA606914
955 G	1 0213AMNM	ESTs, Highly similar to M4 protein homolog [R.norvegicus]	1005058 IMAGE Consortium	AA608319
956 G	3 0213AMNM	solute carrier family 31, member 1	1005135 IMAGE Consortium	AA606611
957 G	5 0213AMNM	ribosomal protein, mitochondrial, S12	1004819 IMAGE Consortium	AA606858
958 G	7 0213AMNM	nuclear distribution gene C homolog (Aspergillus)	1004907 IMAGE Consortium	AA607088
959 G	9 0213AMNM	cadherin 1	1004978 IMAGE Consortium	AA607208
960 G	11 0213AMNM	ESTs	1005619 IMAGE Consortium	AA606931
961 A	1 021AAMNN	ESTs, Highly similar to RNA polymerase III subunit [H.sapiens]	1005715 IMAGE Consortium	AA607140
962 A	3 021AAMNN	melanoma X-actin	1005837 IMAGE Consortium	AI594289
963 A	5 021AAMNN	ESTs, Weakly similar to CENE_HUMAN CENTROMERIC PROTEIN E [H.sapiens]	1005934 IMAGE Consortium	AI594295
964 A	7 021AAMNN	growth factor receptor bound protein 7	1005390 IMAGE Consortium	AA608199
965 A	9 021AAMNN	ESTs, Weakly similar to AF069992_1 LIM domain interacting RING finger protein [M.musculus]	1005605 IMAGE Consortium	AA606925
966 A	11 021AAMNN			

967 C	1	021AAMNN	ESTs	1005718	IMAGE Consortium	AA607132
968 C	3	021AAMNN	myosin light chain, alkali, nonmuscle	1005850	IMAGE Consortium	AA606686
969 C	5	021AAMNN	ESTs, Highly similar to CITRATE SYNTHASE, MITOCHONDRIAL PRECURSOR [Sus scrofa]	1005953	IMAGE Consortium	AA606966
970 C	7	021AAMNN	protein phosphatase 1D magnesium-dependent, delta isoform	1005395	IMAGE Consortium	AA608209
971 C	9	021AAMNN	ESTs, Moderately similar to I53908 major vault protein - rat [R.norvegicus]	1005607	IMAGE Consortium	AA606926
972 C	11	021AAMNN	ESTs	620047	IMAGE Consortium	AI604171
973 E	1	021AAMNN	ephrin B2	1005758	IMAGE Consortium	AA607261
974 E	3	021AAMNN	ESTs	1005854	IMAGE Consortium	AA606688
975 E	5	021AAMNN	DNA segment, Chr 10, ERATO Doi 749, expressed	1005963	IMAGE Consortium	AA607033
976 E	7	021AAMNN	ESTs, Highly similar to 60S RIBOSOMAL PROTEIN L22 [Tripneustes gratilla]	1005405	IMAGE Consortium	AA608212
977 E	9	021AAMNN	histone cell cycle regulation defective homolog A (S. cerevisiae)	1005649	IMAGE Consortium	AA607020
978 E	11	021AAMNN	ESTs	620058	IMAGE Consortium	AA172343
979 G	1	021AAMNN	Finkel-Biskis-Reilly murine sarcoma virus (FBR-MuSV) ubiquitously expressed (fox derived)	1005802	IMAGE Consortium	AA606665
980 G	3	021AAMNN	prosaposin	1005857	IMAGE Consortium	AA606949
981 G	5	021AAMNN	RAB11a, member RAS oncogene family	1005969	IMAGE Consortium	AA607036
982 G	7	021AAMNN	leucine rich repeat (in FLII) interacting protein 1	1005417	IMAGE Consortium	AA608223
983 G	9	021AAMNN	ESTs	1005652	IMAGE Consortium	AA607013
984 G	11	021AAMNN	ESTs	620125	IMAGE Consortium	AI604183
985 A	1	021HAMNO	ESTs	621105	IMAGE Consortium	AI592271
986 A	3	021HAMNO	ESTs	620754	IMAGE Consortium	AI604262
987 A	5	021HAMNO	ESTs	1037669	IMAGE Consortium	AA638790
988 A	7	021HAMNO	ESTs	1007093	IMAGE Consortium	AI594140
989 A	9	021HAMNO	ESTs	1021883	IMAGE Consortium	AI048075
990 A	11	021HAMNO	dimethylarginine dimethylaminohydrolase 2	1038013	IMAGE Consortium	AA636951
991 C	1	021HAMNO	ESTs, Weakly similar to pancortin-1 [M.musculus]	621115	IMAGE Consortium	AA177491
992 C	3	021HAMNO	ESTs	620772	IMAGE Consortium	AI604266
993 C	5	021HAMNO	ESTs	1037742	IMAGE Consortium	AA638858
994 C	7	021HAMNO	ESTs, Highly similar to GAK_RAT CYCLIN G-ASSOCIATED KINASE [R.norvegicus]	1006385	IMAGE Consortium	AA624307
995 C	9	021HAMNO	ESTs	1006667	IMAGE Consortium	AI594121
996 C	11	021HAMNO	LIM and SH3 protein 1	1038029	IMAGE Consortium	AA624426
997 E	1	021HAMNO	ESTs	621131	IMAGE Consortium	AI592279
998 E	3	021HAMNO	small inducible cytokine A12	620800	IMAGE Consortium	AA174647
999 E	5	021HAMNO	Rab6, kinesin-like	1022045	IMAGE Consortium	AI048103
1000 E	7	021HAMNO	ESTs	1006443	IMAGE Consortium	AA624263
1001 E	9	021HAMNO	ESTs, Weakly similar to KIAA0872 protein [H.sapiens]	1006707	IMAGE Consortium	AA624411
1002 E	11	021HAMNO	mannosidase 1, beta	1038035	IMAGE Consortium	AA624429



1003 G	1	021HAMNO	ESTs	620445	IMAGE Consortium	AI604225
1004 G	3	021HAMNO	ESTs	621546	IMAGE Consortium	AI592338
1005 G	5	021HAMNO	gene for 101F6 protein	1006071	IMAGE Consortium	AA607153
1006 G	7	021HAMNO	ESTs, Highly similar to NEDD-4 PROTEIN [Homo sapiens]	1006446	IMAGE Consortium	AA624259
1007 G	9	021HAMNO	ESTs, Weakly similar to phospholipase [H.sapiens]	1022874	IMAGE Consortium	AA623384
1008 G	11	021HAMNO	adipose differentiation related protein	1038042	IMAGE Consortium	AA624422
1009 A	1	021OAMNP	ESTs, Highly similar to HYPOTHETICAL 17.0 KD PROTEIN F56D1.3 IN CHROMOSOME II [Caenorhabditis elegans]	1038128	IMAGE Consortium	AA624679
1010 A	3	021OAMNP	ESTs, Highly similar to PROTEIN TRANSPORT PROTEIN SEC61 BETA SUBUNIT [Homo sapiens; Canis familiaris]	1037686	IMAGE Consortium	AA638788
1011 A	5	021OAMNP	ESTs	1037819	IMAGE Consortium	AA636920
1012 A	7	021OAMNP	nucleophosmin 1	1038071	IMAGE Consortium	AA624447
1013 A	9	021OAMNP	actin-like 6	1038302	IMAGE Consortium	AA671221
1014 A	11	021OAMNP	ESTs, Moderately similar to KIAA0978 protein [H.sapiens]	1038413	IMAGE Consortium	AA624493
1015 C	1	021OAMNP	ESTs, Weakly similar to AF129738_1 cullin 3 [M.musculus]	1038149	IMAGE Consortium	AA624696
1016 C	3	021OAMNP	survival motor neuron	1037706	IMAGE Consortium	AA638804
1017 C	5	021OAMNP	ESTs, Weakly similar to CG7725 gene product [D.melanogaster]	1037839	IMAGE Consortium	AA638922
1018 C	7	021OAMNP	ubiquitin c-terminal hydrolase related polypeptide	1038232	IMAGE Consortium	AA671103
1019 C	9	021OAMNP	heat shock protein, 74 kDa, A	1038304	IMAGE Consortium	AA671222
1020 C	11	021OAMNP	nuclear receptor binding factor 2	1038417	IMAGE Consortium	AA624495
1021 E	1	021OAMNP	tyrosine 3-monooxygenase/tryptophan 5-monooxygenase activation protein, theta polypeptide	1038193	IMAGE Consortium	AA624828
1022 E	3	021OAMNP	enhancer of rudimentary homolog (Drosophila)	1037720	IMAGE Consortium	AA638845
1023 E	5	021OAMNP	mannose-6-phosphate receptor, cation dependent	1037848	IMAGE Consortium	AA638917
1024 E	7	021OAMNP	ribosomal protein, mitochondrial, S14	1038249	IMAGE Consortium	AA671125
1025 E	9	021OAMNP	ESTs	1038310	IMAGE Consortium	AA671225
1026 E	11	021OAMNP	suppressor of Ty 4 homolog (S. cerevisiae)	1038423	IMAGE Consortium	AA624498
1027 G	1	021OAMNP	ESTs, Moderately similar to unnamed protein product [H.sapiens]	1037562	IMAGE Consortium	AA623567
1028 G	3	021OAMNP	ESTs	1037752	IMAGE Consortium	AA638862
1029 G	5	021OAMNP	ESTs	1037864	IMAGE Consortium	AA638929
1030 G	7	021OAMNP	Sid394p	1038265	IMAGE Consortium	AA671202
1031 G	9	021OAMNP	ESTs	1038322	IMAGE Consortium	AA671297
1032 G	11	021OAMNP	eukaryotic translation initiation factor 2B	1038429	IMAGE Consortium	AA624500
1033 A	1	021VAMNQ	ESTs, Weakly similar to AF151865_1 CGI-107 protein [H.sapiens]	1038484	IMAGE Consortium	AA624597
1034 A	3	021VAMNQ	ESTs, Weakly similar to DUS5_RAT DUAL SPECIFICITY PROTEIN PHOSPHATASE 5 [R.norvegicus]	1038599	IMAGE Consortium	AA624901

1035 A	5 021VAMNQ	ESTs, Highly similar to HYPOTHETICAL 64.3 KD GTP-BINDING PROTEIN C02F5.3 IN CHROMOSOME III [Caenorhabditis elegans]	1109656 IMAGE Consortium	AA822787
1036 A	7 021VAMNQ	DNA segment, Chr 17, human D6S56E 2	1110264 IMAGE Consortium	AA795588
1037 A	9 021VAMNQ	ESTs, Weakly similar to truncated form platelet-activating factor acetylhydrolase 45kD subunit [M.musculus]	1111133 IMAGE Consortium	AA688784
1038 A	11 021VAMNQ	protein phosphatase 1, regulatory (inhibitor) subunit 1A	2102241 IMAGE Consortium	AW209345
1039 C	1 021VAMNQ	WSB-1	1038486 IMAGE Consortium	AA624598
1040 C	3 021VAMNQ	ESTs	1037975 IMAGE Consortium	AA636906
1041 C	5 021VAMNQ	cleavage and polyadenylation specific factor 2, 100kD subunit	1109660 IMAGE Consortium	AA822789
1042 C	7 021VAMNQ	folate receptor 1 (adult)	1110386 IMAGE Consortium	AA674821
1043 C	9 021VAMNQ	ESTs, Weakly similar to MDR1_MOUSE MULTIDRUG RESISTANCE PROTEIN 1 [M.musculus]	1111143 IMAGE Consortium	AA688787
1044 C	11 021VAMNQ	ESTs	2101559 IMAGE Consortium	AW209892
1045 E	1 021VAMNQ	ESTs	1038496 IMAGE Consortium	AA624602
1046 E	3 021VAMNQ	ESTs	1082064 IMAGE Consortium	AI593652
1047 E	5 021VAMNQ			
1048 E	7 021VAMNQ	sorting nexin 3	1110090 IMAGE Consortium	AA795488
1049 E	9 021VAMNQ	ESTs, Weakly similar to zinc finger protein 95 [M.musculus]	1110437 IMAGE Consortium	AA675190
1050 E	11 021VAMNQ	DNA segment, Chr 15, Kozak 1	2101846 IMAGE Consortium	AW210138
1051 G	1 021VAMNQ	jumonji	1038498 IMAGE Consortium	AA624603
1052 G	3 021VAMNQ	membrane protein, palmitoylated 6 (MAGUK p55 subfamily member 6)	1109276 IMAGE Consortium	AA666910
1053 G	5 021VAMNQ	ESTs	1109733 IMAGE Consortium	AI591726
1054 G	7 021VAMNQ	Ttk protein kinase	1110210 IMAGE Consortium	AA795558
1055 G	9 021VAMNQ	ESTs, Highly similar to AF161371_1 HSPC253 [H.sapiens]	1110475 IMAGE Consortium	AA675250
1056 G	11 021VAMNQ	mini chromosome maintenance deficient 4 homolog (S. cerevisiae)	2101942 IMAGE Consortium	AW210222
1057 A	1 0212AMNR	ESTs, Moderately similar to AF156878_1 opioid growth factor receptor [R.norvegicus]	2102060 IMAGE Consortium	AW210322
1058 A	3 0212AMNR	purine-nucleoside phosphorylase	2101536 IMAGE Consortium	AW209875
1059 A	5 0212AMNR	ESTs	2101885 IMAGE Consortium	AW210188
1060 A	7 0212AMNR	ESTs, Weakly similar to contains strong similarity to a DNAJ-like domain [C.elegans] {IMAGE:834891}	834891 IMAGE Consortium	AA467252
1061 A	9 0212AMNR	ESTs, Weakly similar to C13F10.2 [C.elegans] {IMAGE:934431}	934431 IMAGE Consortium	AA545049
1062 A	11 0212AMNR	ESTs, Highly similar to Cdc5-like protein [R.norvegicus] {IMAGE:833661}	833661 IMAGE Consortium	AA466838
1063 C	1 0212AMNR	ESTs	2102076 IMAGE Consortium	AW210341
1064 C	3 0212AMNR	lethal giant larvae homolog	2101541 IMAGE Consortium	AW209884
1065 C	5 0212AMNR	wee 1 homolog (S. pombe)	2101164 IMAGE Consortium	AW209546

1066 C	7 0212AMNR	Mus musculus aromatic-L-amino-acid decarboxylase mRNA, complete cds {IMAGE:804722}	804722 IMAGE Consortium	AA474389
1067 C	9 0212AMNR	ESTs, Weakly similar to unknown [S.cerevisiae] {IMAGE:805547}	805547 IMAGE Consortium	AA397252
1068 C	11 0212AMNR	IMAGE EST {IMAGE:834148}	834148 IMAGE Consortium	AA413090
1069 E	1 0212AMNR	cell division cycle 2 homolog A (S. pombe)	2102122 IMAGE Consortium	AW210382
1070 E	3 0212AMNR	ESTs	2101575 IMAGE Consortium	AW209912
1071 E	5 0212AMNR	aplysia ras-related homolog 9 (RhoC)	2101183 IMAGE Consortium	AW209577
1072 E	7 0212AMNR	Mus musculus putative G-protein (GP-1) mRNA, complete cds {IMAGE:760753}	760753 IMAGE Consortium	AA386616
1073 E	9 0212AMNR	ESTs {IMAGE:805579}	805579 IMAGE Consortium	AA397280
1074 E	11 0212AMNR	Mus musculus mRNA for type II phosphatidylinositolphosphate kinase-alpha, complete cds {IMAGE:932803}	932803 IMAGE Consortium	AA538499
1075 G	1 0212AMNR	Mus musculus brain cDNA, clone MNCb-0941, similar to AJ243320 (Canis familiaris)	2102134 IMAGE Consortium	AW210387
1076 G	3 0212AMNR	ESTs, Weakly similar to rbm3 [M.musculus]	2101599 IMAGE Consortium	AW209930
1077 G	5 0212AMNR	ESTs	2101205 IMAGE Consortium	AW209597
1078 G	7 0212AMNR			
1079 G	9 0212AMNR	ESTs {IMAGE:805893}	805893 IMAGE Consortium	AA399827
1080 G	11 0212AMNR	IMAGE EST {IMAGE:761302}	761302 IMAGE Consortium	AA388323
1081 A	1 0219AMNS	ESTs, Weakly similar to ZINC FINGER PROTEIN HF.12 [Homo sapiens] {IMAGE:761019}	761019 IMAGE Consortium	AA387262
1082 A	3 0219AMNS	IMAGE EST {IMAGE:533314}	533314 IMAGE Consortium	AA104976
1083 A	5 0219AMNS	ESTs, Highly similar to Gene product with similarity to Ubiquitin binding enzyme [H.sapiens] {IMAGE:790140}	790140 IMAGE Consortium	AA388099
1084 A	7 0219AMNS	ESTs	619223 IMAGE Consortium	AI449137
1085 A	9 0219AMNS	ESTs	619371 IMAGE Consortium	AA175535
1086 A	11 0219AMNS	ESTs	618941 IMAGE Consortium	AI592512
1087 C	1 0219AMNS	ESTs, Weakly similar to 36k hydrophobic nucleolar protein [M.musculus] {IMAGE:805549}	805549 IMAGE Consortium	AA397253
1088 C	3 0219AMNS	ESTs {IMAGE:523241}	523241 IMAGE Consortium	AA067119
1089 C	5 0219AMNS	ESTs {IMAGE:775484}	775484 IMAGE Consortium	AA387243
1090 C	7 0219AMNS	ESTs, Weakly similar to AF164615_1 Gag-Pro-Pol protein [H.sapiens]	619239 IMAGE Consortium	AI592558
1091 C	9 0219AMNS	ESTs	618662 IMAGE Consortium	AA174964
1092 C	11 0219AMNS	guanine nucleotide binding protein, beta 4	619683 IMAGE Consortium	AI604108
1093 E	1 0219AMNS	IMAGE EST {IMAGE:761264}	761264 IMAGE Consortium	AA388221
1094 E	3 0219AMNS			
1095 E	5 0219AMNS	ESTs	619003 IMAGE Consortium	AA170609
1096 E	7 0219AMNS	ESTs	619255 IMAGE Consortium	AI592562
1097 E	9 0219AMNS	ESTs	619394 IMAGE Consortium	AA175539
1098 E	11 0219AMNS	ESTs	619690 IMAGE Consortium	AI604105

1099 G	1 0219AMNS	ESTs {IMAGE:833720}	833720 IMAGE Consortium	AA466731
1100 G	3 0219AMNS	ESTs {IMAGE:523814}	523814 IMAGE Consortium	AA067146
1101 G	5 0219AMNS	ESTs	619042 IMAGE Consortium	AA175521
1102 G	7 0219AMNS	protein phosphatase 3, catalytic subunit, gamma isoform	619279 IMAGE Consortium	AA175592
1103 G	9 0219AMNS	ESTs	619425 IMAGE Consortium	AI592589
1104 G	11 0219AMNS	ESTs	619759 IMAGE Consortium	AA176044
1105 A	1 021GAMNT	pleckstrin	619998 IMAGE Consortium	AI604164
1106 A	3 021GAMNT	ESTs, Highly similar to NADH-CYTOCHROME B5 REDUCTASE [Homo sapiens] {IMAGE:524540}	524540 IMAGE Consortium	AA068440
1107 A	5 021GAMNT	ESTs, Highly similar to PTB-ASSOCIATED SPLICING FACTOR [Homo sapiens] {IMAGE:550916}	550916 IMAGE Consortium	AA098022
1108 A	7 021GAMNT	ESTs {IMAGE:661780}	661780 IMAGE Consortium	AA207878
1109 A	9 021GAMNT			
1110 A	11 021GAMNT			
1111 C	1 021GAMNT	ESTs, Highly similar to SRB7 [H.sapiens] {IMAGE:524274}	524274 IMAGE Consortium	AA067890
1112 C	3 021GAMNT	ESTs {IMAGE:525134}	525134 IMAGE Consortium	AA096594
1113 C	5 021GAMNT	ESTs, Highly similar to CALCIUM/CALMODULIN-DEPENDENT PROTEIN KINASE TYPE II GAMMA CHAIN [Rattus norvegicus] {IMAGE:532707}	532707 IMAGE Consortium	AA068735
1114 C	7 021GAMNT	ESTs, Moderately similar to RAF SERINE/THREONINE-PROTEIN KINASE TRANSFORMING PROTEIN [Murine sarcoma virus 3611] {IMAGE:651430}	651430 IMAGE Consortium	AA215234
1115 C	9 021GAMNT	ESTs, Moderately similar to the KIAA0197 gene is expressed ubiquitously. [H.sapiens] {IMAGE:790971}	790971 IMAGE Consortium	AA415358
1116 C	11 021GAMNT	ESTs {IMAGE:330495}	330495 IMAGE Consortium	W14898
1117 E	1 021GAMNT	Phosphoribosylglycinamide formyltransferase {IMAGE:535188}	535188 IMAGE Consortium	AA086734
1118 E	3 021GAMNT	Platelet-activating factor acetylhydrolase, isoform 1b, beta1 subunit {IMAGE:523706}	523706 IMAGE Consortium	AA067153
1119 E	5 021GAMNT	IMAGE EST {IMAGE:524338}	524338 IMAGE Consortium	AA065706
1120 E	7 021GAMNT	IMAGE EST {IMAGE:651622}	651622 IMAGE Consortium	AA212170
1121 E	9 021GAMNT	ESTs, Weakly similar to 112.3 KD PROTEIN IN PYK1-SNC1 INTERGENIC REGION [S.cerevisiae] {IMAGE:944355}	944355 IMAGE Consortium	AA547684
1122 E	11 021GAMNT	ESTs {IMAGE:330233}	330233 IMAGE Consortium	W14737
1123 G	1 021GAMNT	IMAGE EST {IMAGE:533100}	533100 IMAGE Consortium	AA068980
1124 G	3 021GAMNT	IMAGE EST {IMAGE:534269}	534269 IMAGE Consortium	AA072765
1125 G	5 021GAMNT	Mus musculus C6-I proteasome chain mRNA, complete cds {IMAGE:550898}	550898 IMAGE Consortium	AA098608
1126 G	7 021GAMNT	ESTs {IMAGE:651903}	651903 IMAGE Consortium	AA212155
1127 G	9 021GAMNT	ESTs, Weakly similar to ELL [M.musculus] {IMAGE:944891}	944891 IMAGE Consortium	AA545429
1128 G	11 021GAMNT	ESTs {IMAGE:330613}	330613 IMAGE Consortium	W14186

1129 A	1 021NAMNU	Mus musculus uterine-specific proline-rich acidic protein mRNA, complete cds {IMAGE:336053}	336053 IMAGE Consortium	W17866
1130 A	3 021NAMNU	IMAGE EST {IMAGE:385063}	385063 IMAGE Consortium	W64752
1131 A	5 021NAMNU	IMAGE EST {IMAGE:421885}	421885 IMAGE Consortium	W91416
1132 A	7 021NAMNU	ESTs, Weakly similar to glucose transport protein, hepatic [M.musculus] {IMAGE:426758}	426758 IMAGE Consortium	AA002666
1133 A	9 021NAMNU	Mus musculus mRNA for synaptotagmin III, complete cds {IMAGE:427562}	427562 IMAGE Consortium	AA002524
1134 A	11 021NAMNU	IMAGE EST {IMAGE:438604}	438604 IMAGE Consortium	AA050979
1135 C	1 021NAMNU	ESTs {IMAGE:338152}	338152 IMAGE Consortium	W29532
1136 C	3 021NAMNU	ESTs {IMAGE:402843}	402843 IMAGE Consortium	W80245
1137 C	5 021NAMNU	IMAGE EST {IMAGE:421712}	421712 IMAGE Consortium	W98962
1138 C	7 021NAMNU	ESTs {IMAGE:426033}	426033 IMAGE Consortium	AA002836
1139 C	9 021NAMNU	IMAGE EST {IMAGE:438706}	438706 IMAGE Consortium	AA051104
1140 C	11 021NAMNU	ESTs, Highly similar to CATION-TRANSPORTING ATPASE PAFL [Synechococcus pcc7942] {IMAGE:442484}	442484 IMAGE Consortium	AA015291
1141 E	1 021NAMNU	IMAGE EST {IMAGE:348006}	348006 IMAGE Consortium	W29778
1142 E	3 021NAMNU	IMAGE EST {IMAGE:403220}	403220 IMAGE Consortium	W82016
1143 E	5 021NAMNU	Protein kinase C, delta {IMAGE:421002}	421002 IMAGE Consortium	W91539
1144 E	7 021NAMNU	IMAGE EST {IMAGE:425664}	425664 IMAGE Consortium	AA000282
1145 E	9 021NAMNU	ESTs {IMAGE:438616}	438616 IMAGE Consortium	AA051009
1146 E	11 021NAMNU	ESTs {IMAGE:444162}	444162 IMAGE Consortium	AA015176
1147 G	1 021NAMNU	Laminin, alpha 2 {IMAGE:333498}	333498 IMAGE Consortium	W18311
1148 G	3 021NAMNU	IMAGE EST {IMAGE:418706}	418706 IMAGE Consortium	W99246
1149 G	5 021NAMNU	ESTs {IMAGE:421155}	421155 IMAGE Consortium	W91036
1150 G	7 021NAMNU	Mouse mRNA for KIFC1, complete cds {IMAGE:426992}	426992 IMAGE Consortium	AA003218
1151 G	9 021NAMNU	Mus musculus mitogen- and stress-activated protein kinase-2 (mMSK2) mRNA, complete cds {IMAGE:439453}	439453 IMAGE Consortium	AA004099
1152 G	11 021NAMNU	IMAGE EST {IMAGE:445988}	445988 IMAGE Consortium	AA020051
1153 A	1 021UAMNV	Mus musculus mRNA for 3'UTR of Clc1 gene {IMAGE:464457}	464457 IMAGE Consortium	AA033085
1154 A	3 021UAMNV	ESTs, Highly similar to PEPSINOGEN F PRECURSOR [Oryctolagus cuniculus] {IMAGE:472743}	472743 IMAGE Consortium	AA038137
1155 A	5 021UAMNV			
1156 A	7 021UAMNV	ESTs	622616 IMAGE Consortium	AA184357
1157 A	9 021UAMNV	ESTs	634458 IMAGE Consortium	AA185687
1158 A	11 021UAMNV	Mus musculus mRNA for CIRP, complete cds {IMAGE:475679}	475679 IMAGE Consortium	AA050300
1159 C	1 021UAMNV	IMAGE EST {IMAGE:464391}	464391 IMAGE Consortium	AA030506
1160 C	3 021UAMNV	IMAGE EST {IMAGE:474305}	474305 IMAGE Consortium	AA038862
1161 C	5 021UAMNV	ESTs	622305 IMAGE Consortium	AA183622
1162 C	7 021UAMNV	ESTs	621941 IMAGE Consortium	AI591860

1163 C	9 021UAMNV	BP-3 alloantigen	634524 IMAGE Consortium	AA185565
1164 C	11 021UAMNV	ESTs {IMAGE:478125}	478125 IMAGE Consortium	AA049592
1165 E	1 021UAMNV	IMAGE EST {IMAGE:463186}	463186 IMAGE Consortium	AA030405
1166 E	3 021UAMNV	IMAGE EST {IMAGE:472374}	472374 IMAGE Consortium	AA036255
1167 E	5 021UAMNV	ESTs, Weakly similar to unknown protein IT1 [H.sapiens]	621633 IMAGE Consortium	AI591821
1168 E	7 021UAMNV	ESTs	622728 IMAGE Consortium	AA184222
1169 E	9 021UAMNV	Cytochrome P450, 2e1, ethanol inducible {IMAGE:476836}	476836 IMAGE Consortium	AA050328
1170 E	11 021UAMNV	ESTs {IMAGE:478428}	478428 IMAGE Consortium	AA049819
1171 G	1 021UAMNV	IMAGE EST {IMAGE:463860}	463860 IMAGE Consortium	AA028539
1172 G	3 021UAMNV	myeloid ecotropic viral integration site-related gene 2	622843 IMAGE Consortium	AA177689
1173 G	5 021UAMNV	ESTs, Weakly similar to PS0136 H-2 class I histocompatibility antigen T15 - mouse [M.musculus]	621662 IMAGE Consortium	AI591822
1174 G	7 021UAMNV	ESTs	622823 IMAGE Consortium	AI662502
1175 G	9 021UAMNV	IMAGE EST {IMAGE:475381}	475381 IMAGE Consortium	AA041819
1176 G	11 021UAMNV	Mus musculus ETO/MTG8-related protein ETO-2 mRNA, complete cds {IMAGE:479774}	479774 IMAGE Consortium	AA051563
1177 A	1 0211AMNW	EST {IMAGE:481950}	481950 IMAGE Consortium	AA059950
1178 A	3 0211AMNW	Kinesin heavy chain member 4 {IMAGE:481146}	481146 IMAGE Consortium	AA060358
1179 A	5 0211AMNW	IMAGE EST {IMAGE:493212}	493212 IMAGE Consortium	AA097795
1180 A	7 0211AMNW	Mus domesticus ZNF127 (zfn127) gene, complete cds {IMAGE:574274}	574274 IMAGE Consortium	AA119257
1181 A	9 0211AMNW	IMAGE EST {IMAGE:580810}	580810 IMAGE Consortium	AA146027
1182 A	11 0211AMNW			
1183 C	1 0211AMNW	IMAGE EST {IMAGE:483753}	483753 IMAGE Consortium	AA061408
1184 C	3 0211AMNW	IMAGE EST {IMAGE:483006}	483006 IMAGE Consortium	AA060858
1185 C	5 0211AMNW	IMAGE EST {IMAGE:571219}	571219 IMAGE Consortium	AA107718
1186 C	7 0211AMNW	ESTs, Weakly similar to ORF YLL039c [S.cerevisiae] {IMAGE:575698}	575698 IMAGE Consortium	AA123007
1187 C	9 0211AMNW	ESTs, Weakly similar to VITAMIN K-DEPENDENT GAMMA-CARBOXYLASE [Homo sapiens] {IMAGE:582706}	582706 IMAGE Consortium	AA154146
1188 C	11 0211AMNW	IMAGE EST {IMAGE:583238}	583238 IMAGE Consortium	AA154812
1189 E	1 0211AMNW	ESTs, Highly similar to T-CADHERIN PRECURSOR [Gallus gallus] {IMAGE:482972}	482972 IMAGE Consortium	AA061041
1190 E	3 0211AMNW	ESTs {IMAGE:483150}	483150 IMAGE Consortium	AA060910
1191 E	5 0211AMNW	IMAGE EST {IMAGE:574119}	574119 IMAGE Consortium	AA119086
1192 E	7 0211AMNW	ESTs {IMAGE:575605}	575605 IMAGE Consortium	AA124548
1193 E	9 0211AMNW	IMAGE EST {IMAGE:582588}	582588 IMAGE Consortium	AA153968
1194 E	11 0211AMNW	ESTs {IMAGE:598870}	598870 IMAGE Consortium	AA162785
1195 G	1 0211AMNW	ATP-binding cassette 1 {IMAGE:482369}	482369 IMAGE Consortium	AA063753
1196 G	3 0211AMNW	ESTs {IMAGE:483014}	483014 IMAGE Consortium	AA060971
1197 G	5 0211AMNW	IMAGE EST {IMAGE:573770}	573770 IMAGE Consortium	AA118841

1198 G	7 0211AMNW	ESTs, Moderately similar to zinc finger protein [M.musculus] {IMAGE:574258}	574258 IMAGE Consortium	AA117644
1199 G	9 0211AMNW	ESTs {IMAGE:580935}	580935 IMAGE Consortium	AA139918
1200 G	11 0211AMNW	Oncostatin receptor {IMAGE:598146}	598146 IMAGE Consortium	AA265259
1201 A	1 0218AMNX	IMAGE EST {IMAGE:597493}	597493 IMAGE Consortium	AA154419
1202 A	3 0218AMNX	ESTs, Moderately similar to KIAA0365 [H.sapiens] {IMAGE:598206}	598206 IMAGE Consortium	AA161817
1203 A	5 0218AMNX	ESTs {IMAGE:620659}	620659 IMAGE Consortium	AA174533
1204 A	7 0218AMNX	ESTs {IMAGE:617359}	617359 IMAGE Consortium	AA174428
1205 A	9 0218AMNX	IMAGE EST {IMAGE:635240}	635240 IMAGE Consortium	AA189716
1206 A	11 0218AMNX	IMAGE EST {IMAGE:634503}	634503 IMAGE Consortium	AA185562
1207 C	1 0218AMNX	IMAGE EST {IMAGE:583809}	583809 IMAGE Consortium	AA153024
1208 C	3 0218AMNX	ESTs {IMAGE:618483}	618483 IMAGE Consortium	AA175284
1209 C	5 0218AMNX	IMAGE EST {IMAGE:619207}	619207 IMAGE Consortium	AA175695
1210 C	7 0218AMNX	Forkhead homolog 14, (Drosophila) {IMAGE:619776}	619776 IMAGE Consortium	AA176045
1211 C	9 0218AMNX	IMAGE EST {IMAGE:634466}	634466 IMAGE Consortium	AA184909
1212 C	11 0218AMNX	IMAGE EST {IMAGE:634504}	634504 IMAGE Consortium	AA185701
1213 E	1 0218AMNX	ESTs {IMAGE:596543}	596543 IMAGE Consortium	AA138313
1214 E	3 0218AMNX	Mus musculus classII MHC transactivator CIITA mRNA, complete cds {IMAGE:620236}	620236 IMAGE Consortium	AA178010
1215 E	5 0218AMNX	IMAGE EST {IMAGE:619958}	619958 IMAGE Consortium	AA178037
1216 E	7 0218AMNX	IMAGE EST {IMAGE:617437}	617437 IMAGE Consortium	AA174397
1217 E	9 0218AMNX	IMAGE EST {IMAGE:622693}	622693 IMAGE Consortium	AA184189
1218 E	11 0218AMNX	ESTs {IMAGE:634580}	634580 IMAGE Consortium	AA189624
1219 G	1 0218AMNX	ESTs, Highly similar to cyclic nucleotide phosphodiesterase [M.musculus] {IMAGE:597524}	597524 IMAGE Consortium	AA144377
1220 G	3 0218AMNX	IMAGE EST {IMAGE:617193}	617193 IMAGE Consortium	AA174774
1221 G	5 0218AMNX	IMAGE EST {IMAGE:619141}	619141 IMAGE Consortium	AA175618
1222 G	7 0218AMNX	Nitric oxide synthase 3, endothelial cell {IMAGE:620940}	620940 IMAGE Consortium	AA177240
1223 G	9 0218AMNX	ESTs {IMAGE:622458}	622458 IMAGE Consortium	AA177358
1224 G	11 0218AMNX	ESTs {IMAGE:636007}	636007 IMAGE Consortium	AA183350
1225 A	1 021FAMNY	IMAGE EST {IMAGE:636193}	636193 IMAGE Consortium	AA183187
1226 A	3 021FAMNY	Immunoglobulin heavy chain 1 (serum IgG2a) {IMAGE:636695}	636695 IMAGE Consortium	AA182334
1227 A	5 021FAMNY	IMAGE EST {IMAGE:636714}	636714 IMAGE Consortium	AA182335
1228 A	7 021FAMNY	ESTs {IMAGE:642379}	642379 IMAGE Consortium	AA185202
1229 A	9 021FAMNY	ESTs {IMAGE:642205}	642205 IMAGE Consortium	AA212243
1230 A	11 021FAMNY	ESTs	635065 IMAGE Consortium	AA163260
1231 C	1 021FAMNY	ESTs {IMAGE:622253}	622253 IMAGE Consortium	AA183369
1232 C	3 021FAMNY	ESTs {IMAGE:638350}	638350 IMAGE Consortium	AA200703
1233 C	5 021FAMNY	IMAGE EST {IMAGE:640501}	640501 IMAGE Consortium	AA204066
1234 C	7 021FAMNY	Kinesin family member 3b {IMAGE:641650}	641650 IMAGE Consortium	AA209952
1235 C	9 021FAMNY			

1236 C	11 021FAMNY	ESTs	635124 IMAGE Consortium	AA162409
1237 E	1 021FAMNY	IMAGE EST {IMAGE:636361}	636361 IMAGE Consortium	AA183016
1238 E	3 021FAMNY	ESTs {IMAGE:637291}	637291 IMAGE Consortium	AA185873
1239 E	5 021FAMNY	IMAGE EST {IMAGE:638419}	638419 IMAGE Consortium	AA200293
1240 E	7 021FAMNY	ESTs, Weakly similar to QUINONE OXIDOREDUCTASE [M.musculus] {IMAGE:653324}	653324 IMAGE Consortium	AA241313
1241 E	9 021FAMNY	ESTs {IMAGE:652983}	652983 IMAGE Consortium	AA217415
1242 E	11 021FAMNY	ESTs	635185 IMAGE Consortium	AA178741
1243 G	1 021FAMNY	ESTs {IMAGE:634790}	634790 IMAGE Consortium	AA163180
1244 G	3 021FAMNY	IMAGE EST {IMAGE:639008}	639008 IMAGE Consortium	AA200172
1245 G	5 021FAMNY	ESTs {IMAGE:637233}	637233 IMAGE Consortium	AA189900
1246 G	7 021FAMNY	IMAGE EST {IMAGE:642847}	642847 IMAGE Consortium	AA185648
1247 G	9 021FAMNY	ESTs {IMAGE:642394}	642394 IMAGE Consortium	AA184806
1248 G	11 021FAMNY	ESTs	635231 IMAGE Consortium	AA189761
1249 A	1 021MAMNZ	DNA segment, Chr 16, ERATO Doi 536, expressed	634633 IMAGE Consortium	AA154594
1250 A	3 021MAMNZ	ESTs	635624 IMAGE Consortium	AA166304
1251 A	5 021MAMNZ	ESTs	635962 IMAGE Consortium	AA183287
1252 A	7 021MAMNZ	forkhead box K1	317953 IMAGE Consortium	W11538
1253 A	9 021MAMNZ	mucin 1, transmembrane	333411 IMAGE Consortium	W16358
1254 A	11 021MAMNZ	guanine nucleotide binding protein, alpha o	354068 IMAGE Consortium	W43961
1255 C	1 021MAMNZ	ESTs	634688 IMAGE Consortium	AI592740
1256 C	3 021MAMNZ	ESTs	635649 IMAGE Consortium	AA166546
1257 C	5 021MAMNZ	T-box brain gene 1	303483 IMAGE Consortium	W09676
1258 C	7 021MAMNZ	mitogen activated protein kinase kinase kinase kinase 1	318867 IMAGE Consortium	W34229
1259 C	9 021MAMNZ	phosphodiesterase 1A, calmodulin-dependent	333755 IMAGE Consortium	W16027
1260 C	11 021MAMNZ	forkhead box D1	367062 IMAGE Consortium	AI893923
1261 E	1 021MAMNZ	ESTs	634705 IMAGE Consortium	AA154795
1262 E	3 021MAMNZ	ESTs, Weakly similar to putative REX-2 [M.musculus]	635676 IMAGE Consortium	AA165750
1263 E	5 021MAMNZ	androgen regulated vas deferens protein	313461 IMAGE Consortium	W10293
1264 E	7 021MAMNZ	kallikrein binding protein	330805 IMAGE Consortium	W14912
1265 E	9 021MAMNZ	potassium channel, subfamily K, member 3	334194 IMAGE Consortium	W36914
1266 E	11 021MAMNZ	annexin A11	368271 IMAGE Consortium	W54181
1267 G	1 021MAMNZ	ESTs	634716 IMAGE Consortium	AI666289
1268 G	3 021MAMNZ	ESTs	634981 IMAGE Consortium	AA164054
1269 G	5 021MAMNZ	non-catalytic region of tyrosine kinase adaptor protein 2	315152 IMAGE Consortium	W08947
1270 G	7 021MAMNZ	protein tyrosine phosphatase 4a3	331144 IMAGE Consortium	W13832
1271 G	9 021MAMNZ	properdin factor, complement	334712 IMAGE Consortium	W18268
1272 G	11 021MAMNZ	ESTs	374232 IMAGE Consortium	AI893930
1273 A	1 021MAMOO	ELAV (embryonic lethal, abnormal vision, Drosophila)-like 3 (Hu antigen C)	404503 IMAGE Consortium	W83106
1274 A	3 021MAMOO	paired related homeobox 2	422870 IMAGE Consortium	W97877
1275 A	5 021MAMOO	RAD51 like 3 (S. cerevisiae)	478233 IMAGE Consortium	AA049653



1276 A	7 021MAM00	cadherin 15	477228 IMAGE Consortium	AI323029
1277 A	9 021MAM00			
1278 A	11 021MAM00	folate hydrolase	579705 IMAGE Consortium	AA116793
1279 C	1 021MAM00	solute carrier family 9 (sodium/hydrogen exchanger), member 1	370997 IMAGE Consortium	W53273
1280 C	3 021MAM00	casein kinase II, alpha 1, related sequence 4	439256 IMAGE Consortium	AA003928
1281 C	5 021MAM00	phosphatidylinositol-4-phosphate 5-kinase, type 1 alpha	479073 IMAGE Consortium	AA048876
1282 C	7 021MAM00	solute carrier family 22, member 2	572016 IMAGE Consortium	AI587821
1283 C	9 021MAM00	paired-Ig-like receptor B	598176 IMAGE Consortium	AA161806
1284 C	11 021MAM00	immunoglobulin kappa chain variable 28 (V28)	597591 IMAGE Consortium	AA152940
1285 E	1 021MAM00	T-cell death associated gene	374607 IMAGE Consortium	W59335
1286 E	3 021MAM00	granzyme G	442922 IMAGE Consortium	AA014972
1287 E	5 021MAM00	macrophage receptor with collagenous structure	481356 IMAGE Consortium	AA060455
1288 E	7 021MAM00	ecotropic viral integration site 2	574757 IMAGE Consortium	AA124623
1289 E	9 021MAM00	interferon-induced protein with tetratricopeptide repeats 1	616653 IMAGE Consortium	AA170469
1290 E	11 021MAM00	macrophage expressed gene 1	620773 IMAGE Consortium	AA178272
1291 G	1 021MAM00	lymphocyte specific 1	388693 IMAGE Consortium	W71049
1292 G	3 021MAM00			
1293 G	5 021MAM00	protease, serine, 12 neurotrypsin, (motopsin)	484148 IMAGE Consortium	AI893676
1294 G	7 021MAM00	PCTAIRE-motif protein kinase 3	578334 IMAGE Consortium	AA123333
1295 G	9 021MAM00	ubiquitin-protein ligase e3 componen n-recognin	619438 IMAGE Consortium	AA172532
1296 G	11 021MAM00	circadian locomoter output cycles kaput	636179 IMAGE Consortium	AA183163
1297 A	1 021TAMO1	wingless-related MMTV integration site 6	670181 IMAGE Consortium	AA222138
1298 A	3 021TAMO1	alpha-L-iduronidase	694087 IMAGE Consortium	AA238872
1299 A	5 021TAMO1	oxytocin	640127 IMAGE Consortium	AA197698
1300 A	7 021TAMO1	nuclear receptor subfamily 1, group I, member 2	737758 IMAGE Consortium	AA277370
1301 A	9 021TAMO1	avian reticuloendotheliosis viral (v-rel) oncogene related B	820638 IMAGE Consortium	AA673132
1302 A	11 021TAMO1	novel ras effector 1	484206 IMAGE Consortium	AI552504
1303 C	1 021TAMO1	zinc finger protein of the cerebellum 1	671755 IMAGE Consortium	AA242256
1304 C	3 021TAMO1	procollagen, type XII, alpha 1	696896 IMAGE Consortium	AA220331
1305 C	5 021TAMO1	nescient helix loop helix 1	660623 IMAGE Consortium	AA223023
1306 C	7 021TAMO1	ELAV (embryonic lethal, abnormal vision, Drosophila)-like 4 (Hu antigen D)	746596 IMAGE Consortium	AA268007
1307 C	9 021TAMO1	tumor necrosis factor receptor superfamily, member 11b (osteoprotegerin)	851484 IMAGE Consortium	AA461886
1308 C	11 021TAMO1	small inducible cytokine subfamily A, member 22	577486 IMAGE Consortium	AA137913
1309 E	1 021TAMO1	murinoglobulin 2	677145 IMAGE Consortium	AA212714
1310 E	3 021TAMO1	contactin 3	716972 IMAGE Consortium	AA266190
1311 E	5 021TAMO1	cytidine monophospho-N-acetylneuraminic acid hydroxylase	679492 IMAGE Consortium	AA244565
1312 E	7 021TAMO1	coagulation factor IX	748308 IMAGE Consortium	AA267985
1313 E	9 021TAMO1	retinoid X receptor alpha	871479 IMAGE Consortium	AA462070
1314 E	11 021TAMO1	c-myc binding protein	583913 IMAGE Consortium	AA145831

1315 G	1 021TAMO1	mitogen activated protein kinase kinase kinase 5	678913 IMAGE Consortium	AA245466
1316 G	3 021TAMO1	ESTs, Highly similar to G10D_MOUSE PROBABLE G PROTEIN-COUPLED RECEPTOR G10D [M.musculus]	717887 IMAGE Consortium	AA250557
1317 G	5 021TAMO1	rat regenerating islet-derived, mouse homolog 1	680179 IMAGE Consortium	AA237793
1318 G	7 021TAMO1	carbonic anhydrase 5, mitochondrial	748650 IMAGE Consortium	AA270290
1319 G	9 021TAMO1	fatty acid binding protein 2, intestinal	921551 IMAGE Consortium	AA529824
1320 G	11 021TAMO1	zinc finger protein 326	640140 IMAGE Consortium	AA197562
1321 A	1 0210AMO2	ESTs	638229 IMAGE Consortium	AI646672
1322 A	3 0210AMO2	ESTs	638636 IMAGE Consortium	AA200256
1323 A	5 0210AMO2	ESTs, Weakly similar to T17453 ERG-associated protein ESET - mouse [M.musculus]	640148 IMAGE Consortium	AA197701
1324 A	7 0210AMO2	EST	636910 IMAGE Consortium	AI450169
1325 A	9 0210AMO2	ESTs, Moderately similar to MO25 PROTEIN [Mus musculus]	636362 IMAGE Consortium	AI645170
1326 A	11 0210AMO2	ganglioside-induced differentiation-associated-protein 3	636517 IMAGE Consortium	AA182242
1327 C	1 0210AMO2	Mus musculus Slc4a10 mRNA for NCBE, complete cds	638329 IMAGE Consortium	AA200780
1328 C	3 0210AMO2	ESTs	639476 IMAGE Consortium	AA200855
1329 C	5 0210AMO2	ESTs, Highly similar to RADIAL SPOKE PROTEIN 3 [Chlamydomonas reinhardtii]	640195 IMAGE Consortium	AA197708
1330 C	7 0210AMO2	Mus musculus X chromosome: L1cam locus	636941 IMAGE Consortium	AI645219
1331 C	9 0210AMO2	ESTs	637097 IMAGE Consortium	AI645242
1332 C	11 0210AMO2	ESTs	636532 IMAGE Consortium	AA182274
1333 E	1 0210AMO2	ESTs, Weakly similar to Collagen alpha1 [R.norvegicus]	639085 IMAGE Consortium	AA199569
1334 E	3 0210AMO2	kinesin family member 21B	638899 IMAGE Consortium	AI645902
1335 E	5 0210AMO2	ESTs	640317 IMAGE Consortium	AA199371
1336 E	7 0210AMO2	ESTs	636948 IMAGE Consortium	AI645217
1337 E	9 0210AMO2	ESTs	636395 IMAGE Consortium	AA183207
1338 E	11 0210AMO2	formyl peptide receptor 1	636543 IMAGE Consortium	AA182250
1339 G	1 0210AMO2	ESTs	638441 IMAGE Consortium	AA200346
1340 G	3 0210AMO2	ESTs	638906 IMAGE Consortium	AI645903
1341 G	5 0210AMO2	serine/threonine kinase 19	640504 IMAGE Consortium	AA204212
1342 G	7 0210AMO2	ESTs	636172 IMAGE Consortium	AI645148
1343 G	9 0210AMO2	insulin-like growth factor binding protein 5	636421 IMAGE Consortium	AA183239
1344 G	11 0210AMO2	Mus musculus Sep2 mRNA, partial cds	637302 IMAGE Consortium	AA189636
1345 A	1 0217AMO3	ESTs, Weakly similar to collagen type VII [H.sapiens]	640727 IMAGE Consortium	AA204410
1346 A	3 0217AMO3	ESTs	642528 IMAGE Consortium	AI606028
1347 A	5 0217AMO3	integrin beta 1 binding protein 2	336703 IMAGE Consortium	AI390966
1348 A	7 0217AMO3	ESTs, Weakly similar to choline/ethanolamine kinase [M.musculus]	336420 IMAGE Consortium	AI390183
1349 A	9 0217AMO3	ESTs	643315 IMAGE Consortium	AA189896
1350 A	11 0217AMO3	EGF-like module containing, mucin-like, hormone receptor-like sequence 1	642792 IMAGE Consortium	AA185619
1351 C	1 0217AMO3	Mus musculus proteasome beta-subunit C5 (Psmb1)	642328 IMAGE Consortium	AA185248

		mRNA, partial cds		
1352 C	3 0217AMO3	GATA-binding protein 1	642552 IMAGE Consortium	AA185505
1353 C	5 0217AMO3	ESTs	337645 IMAGE Consortium	AI391122
1354 C	7 0217AMO3	ESTs	336470 IMAGE Consortium	W18499
1355 C	9 0217AMO3	ESTs	643323 IMAGE Consortium	AA189409
1356 C	11 0217AMO3	ESTs	642826 IMAGE Consortium	AA189216
1357 E	1 0217AMO3	small inducible cytokine B subfamily (Cys-X-Cys), member 9	642354 IMAGE Consortium	AA185256
1358 E	3 0217AMO3	DNA segment, Chr 4, ERATO Doi 314, expressed	642582 IMAGE Consortium	AI606031
1359 E	5 0217AMO3	hypothetical protein, clone 1-73	348058 IMAGE Consortium	W29809
1360 E	7 0217AMO3	ESTs	644912 IMAGE Consortium	AI450469
1361 E	9 0217AMO3	ESTs	643392 IMAGE Consortium	AI390445
1362 E	11 0217AMO3	ESTs	644290 IMAGE Consortium	AA208743
1363 G	1 0217AMO3	ESTs	642391 IMAGE Consortium	AI606011
1364 G	3 0217AMO3	ESTs, Highly similar to Unknown [H.sapiens]	642605 IMAGE Consortium	AA185340
1365 G	5 0217AMO3	ESTs	348146 IMAGE Consortium	AI391037
1366 G	7 0217AMO3	ESTs	643076 IMAGE Consortium	AI627078
1367 G	9 0217AMO3	ESTs	643411 IMAGE Consortium	AA186117
1368 G	11 0217AMO3	ESTs	644297 IMAGE Consortium	AA204527
1369 A	1 021EAMO4	ESTs	643678 IMAGE Consortium	AI390467
1370 A	3 021EAMO4	ESTs	643942 IMAGE Consortium	AA203947
1371 A	5 021EAMO4	ESTs	659565 IMAGE Consortium	AA219796
1372 A	7 021EAMO4	ESTs, Moderately similar to KIAA0738 protein [H.sapiens]	660172 IMAGE Consortium	AA220355
1373 A	9 021EAMO4	ESTs, Highly similar to dynein light intermediate chain 53/55 [R.norvegicus]	669990 IMAGE Consortium	AA253855
1374 A	11 021EAMO4	ESTs	653602 IMAGE Consortium	AI390749
1375 C	1 021EAMO4	ESTs, Moderately similar to PWP1_HUMAN PERIODIC TRYPTOPHAN PROTEIN 1 HOMOLOG [H.sapiens]	642958 IMAGE Consortium	AI627071
1376 C	3 021EAMO4	ESTs, Highly similar to cordon-bleu gene product [M.musculus]	660806 IMAGE Consortium	AI644849
1377 C	5 021EAMO4	ESTs	659620 IMAGE Consortium	AA219849
1378 C	7 021EAMO4	ESTs	661000 IMAGE Consortium	AA230694
1379 C	9 021EAMO4	synaptogyrin 4	670113 IMAGE Consortium	AA231032
1380 C	11 021EAMO4	ESTs	652925 IMAGE Consortium	AI390667
1381 E	1 021EAMO4	ESTs	642972 IMAGE Consortium	AA185212
1382 E	3 021EAMO4	ESTs	670560 IMAGE Consortium	AA230809
1383 E	5 021EAMO4	ESTs	659649 IMAGE Consortium	AI390192
1384 E	7 021EAMO4	procollagen, type II, alpha 1	669703 IMAGE Consortium	AA253899
1385 E	9 021EAMO4	ESTs	660724 IMAGE Consortium	AA222067
1386 E	11 021EAMO4	ESTs	652991 IMAGE Consortium	AI390673
1387 G	1 021EAMO4	ESTs	642989 IMAGE Consortium	AA185588
1388 G	3 021EAMO4	ESTs	659032 IMAGE Consortium	AI390440
1389 G	5 021EAMO4	DNA segment, Chr 14, ERATO Doi 581, expressed	658949 IMAGE Consortium	AI390435

1390 G	7 021EAMO4	ESTs	669752 IMAGE Consortium	AA230712
1391 G	9 021EAMO4	Tnf receptor-associated factor 3	653279 IMAGE Consortium	AA241302
1392 G	11 021EAMO4	ESTs	653740 IMAGE Consortium	AI390759
1393 A	1 021LAMO5	ESTs	653181 IMAGE Consortium	AI390697
1394 A	3 021LAMO5	ESTs	652703 IMAGE Consortium	AI390602
1395 A	5 021LAMO5	complement factor H-related protein	680298 IMAGE Consortium	AA237589
1396 A	7 021LAMO5	ESTs	676816 IMAGE Consortium	AI391431
1397 A	9 021LAMO5	Mus musculus unknown mRNA, partial cds	697189 IMAGE Consortium	AA221818
1398 A	11 021LAMO5	ESTs, Weakly similar to cDNA EST EMBL:D67419 comes from this gene [C.elegans]	681381 IMAGE Consortium	AA242493
1399 C	1 021LAMO5	ESTs	652500 IMAGE Consortium	AI390560
1400 C	3 021LAMO5	ESTs	679580 IMAGE Consortium	AA245063
1401 C	5 021LAMO5	yolk sac gene 2	680326 IMAGE Consortium	AA237605
1402 C	7 021LAMO5	cyclin-dependent kinase-like 2 (CDC2-related kinase)	679056 IMAGE Consortium	AA245952
1403 C	9 021LAMO5	ESTs	696525 IMAGE Consortium	AI425433
1404 C	11 021LAMO5	peroxisomal membrane protein 2, 22 kDa	680787 IMAGE Consortium	AA242441
1405 E	1 021LAMO5	SH3 domain protein 1B	652549 IMAGE Consortium	AI390576
1406 E	3 021LAMO5	ESTs, Moderately similar to Unknown [H.sapiens]	679587 IMAGE Consortium	AI451789
1407 E	5 021LAMO5	ESTs	680390 IMAGE Consortium	AA237721
1408 E	7 021LAMO5	ESTs, Moderately similar to A37990 aldose reductase homolog - mouse [M.musculus]	676912 IMAGE Consortium	AI390242
1409 E	9 021LAMO5	ESTs	696746 IMAGE Consortium	AI466802
1410 E	11 021LAMO5	ESTs		
1411 G	1 021LAMO5	ESTs	652595 IMAGE Consortium	AI390584
1412 G	3 021LAMO5	sulfotransferase-related protein SULT-X2	679629 IMAGE Consortium	AA245097
1413 G	5 021LAMO5	ESTs	680412 IMAGE Consortium	AA237727
1414 G	7 021LAMO5	ESTs	678570 IMAGE Consortium	AI390287
1415 G	9 021LAMO5	ESTs	696920 IMAGE Consortium	AI425441
1416 G	11 021LAMO5	lecithin cholesterol acyltransferase	680857 IMAGE Consortium	AA250068
1417 A	1 021SAMO6	nuclear receptor subfamily 1, group H, member 3	681807 IMAGE Consortium	AA244389
1418 A	3 021SAMO6	ESTs	693741 IMAGE Consortium	AA265101
1419 A	5 021SAMO6	ESTs	719297 IMAGE Consortium	AA254875
1420 A	7 021SAMO6	ESTs	718913 IMAGE Consortium	AI464678
1421 A	9 021SAMO6	hypothetical protein I54	720639 IMAGE Consortium	AA261489
1422 A	11 021SAMO6	ESTs	718432 IMAGE Consortium	AI464601
1423 C	1 021SAMO6	ESTs	692300 IMAGE Consortium	AA260592
1424 C	3 021SAMO6	ESTs, Highly similar to KYNU_RAT KYNURENINASE [R.norvegicus]	694610 IMAGE Consortium	AA220156
1425 C	5 021SAMO6	ESTs	719336 IMAGE Consortium	AA255015
1426 C	7 021SAMO6	ESTs	719786 IMAGE Consortium	AA261046
1427 C	9 021SAMO6	ESTs	720673 IMAGE Consortium	AI464834
1428 C	11 021SAMO6	ESTs, Highly similar to SERINE/THREONINE PROTEIN PHOSPHATASE 2B CATALYTIC SUBUNIT, BETA	717754 IMAGE Consortium	AI607728

		ISOFORM [Homo sapiens]		
1429 E	1	021SAMO6 ESTs	693151	IMAGE Consortium AI466982
1430 E	3	021SAMO6 ESTs	719086	IMAGE Consortium AA266018
1431 E	5	021SAMO6 ESTs	719342	IMAGE Consortium AA255069
1432 E	7	021SAMO6 cytochrome P450, 7a1	719901	IMAGE Consortium AI464796
1433 E	9	021SAMO6 DNA segment, Chr 16, ERATO Doi 450, expressed	720699	IMAGE Consortium AI464839
1434 E	11	021SAMO6 ESTs, Moderately similar to unnamed protein product [M.musculus]	717823	IMAGE Consortium AA250524
1435 G	1	021SAMO6 ESTs	693305	IMAGE Consortium AA245289
1436 G	3	021SAMO6 ESTs	719106	IMAGE Consortium AA267545
1437 G	5	021SAMO6 ESTs	719406	IMAGE Consortium AA255253
1438 G	7	021SAMO6 complement component 2 (within H-2S)	720089	IMAGE Consortium AA255312
1439 G	9	021SAMO6 ESTs	718349	IMAGE Consortium AI464592
1440 G	11	021SAMO6 ESTs	717826	IMAGE Consortium AA249991
1441 A	1	021ZAMO7 ESTs	718713	IMAGE Consortium AI464653
1442 A	3	021ZAMO7 ESTs	718070	IMAGE Consortium AA266428
1443 A	5	021ZAMO7 ESTs, Weakly similar to KRAB-zinc finger protein KZF-1 [R.norvegicus]	721245	IMAGE Consortium AA267204
1444 A	7	021ZAMO7 ESTs, Moderately similar to AF151805_1 CGI-47 protein [H.sapiens]	721449	IMAGE Consortium AA266704
1445 A	9	021ZAMO7 ESTs	720888	IMAGE Consortium AI464865
1446 A	11	021ZAMO7 ESTs	721853	IMAGE Consortium AI595316
1447 C	1	021ZAMO7 ESTs	718722	IMAGE Consortium AI464649
1448 C	3	021ZAMO7 nuclear factor I/B	716653	IMAGE Consortium AA265406
1449 C	5	021ZAMO7 ESTs	721280	IMAGE Consortium AI464920
1450 C	7	021ZAMO7 ESTs	722182	IMAGE Consortium AA268537
1451 C	9	021ZAMO7 runt related transcription factor 3	722331	IMAGE Consortium AI608086
1452 C	11	021ZAMO7 Mus musculus transposon-derived Buster2 transposase-like protein gene, partial cds	721935	IMAGE Consortium AI595325
1453 E	1	021ZAMO7 ESTs	698171	IMAGE Consortium AI466749
1454 E	3	021ZAMO7 ESTs, Moderately similar to AF127481_1 non-ocogenic Rho GTPase-specific GTP exchange factor [H.sapiens]	718173	IMAGE Consortium AA266350
1455 E	5	021ZAMO7 ESTs	721312	IMAGE Consortium AI464924
1456 E	7	021ZAMO7 ESTs, Highly similar to SERINE/THREONINE PROTEIN PHOSPHATASE PPE1 [Schizosaccharomyces pombe]	720753	IMAGE Consortium AA266807
1457 E	9	021ZAMO7 ESTs, Weakly similar to AF152929_1 A-kinase anchoring protein 18 gamma [H.sapiens]	720925	IMAGE Consortium AA267224
1458 E	11	021ZAMO7 ESTs	721993	IMAGE Consortium AI608041
1459 G	1	021ZAMO7 ESTs	699189	IMAGE Consortium AI466765
1460 G	3	021ZAMO7 early growth response 1	721000	IMAGE Consortium AA266872
1461 G	5	021ZAMO7 ESTs	721329	IMAGE Consortium AI464929
1462 G	7	021ZAMO7 inhibitor of kappa light polypeptide gene enhancer in B-cells, kinase beta	720766	IMAGE Consortium AI464846

1463 G	9 021ZAMO7	ESTs	720957 IMAGE Consortium	AI464874
1464 G	11 021ZAMO7	ESTs	734374 IMAGE Consortium	AA268753
1465 A	1 0216AMO8	ESTs, Weakly similar to insulin receptor substrate protein of 53 kDa [H.sapiens]	734723 IMAGE Consortium	AA259290
1466 A	3 0216AMO8	ESTs	723159 IMAGE Consortium	AI463748
1467 A	5 0216AMO8	ESTs	723319 IMAGE Consortium	AA267882
1468 A	7 0216AMO8	hypothetical proetin MNCb-4414, similar to NM_005827 UDP-galactose transporter related (Homo sapiens)	733513 IMAGE Consortium	AA259871
1469 A	9 0216AMO8	ESTs	749272 IMAGE Consortium	AA289726
1470 A	11 0216AMO8	ESTs	749593 IMAGE Consortium	AA388698
1471 C	1 0216AMO8	SRY-box containing gene 18	734946 IMAGE Consortium	AA261240
1472 C	3 0216AMO8	ESTs	722458 IMAGE Consortium	AA254298
1473 C	5 0216AMO8	Public domain EST	722786 IMAGE Consortium	
1474 C	7 0216AMO8	hypothetical protein, clone 2-25	733537 IMAGE Consortium	AA260011
1475 C	9 0216AMO8	ESTs, Weakly similar to Y141_HUMAN HYPOTHETICAL PROTEIN KIAA0141 [H.sapiens]	748615 IMAGE Consortium	AA269733
1476 C	11 0216AMO8	ESTs	749601 IMAGE Consortium	AA288465
1477 E	1 0216AMO8	ESTs	722199 IMAGE Consortium	AI608061
1478 E	3 0216AMO8	ESTs	723201 IMAGE Consortium	AI661389
1479 E	5 0216AMO8	ESTs	723530 IMAGE Consortium	AI463771
1480 E	7 0216AMO8	ESTs	735012 IMAGE Consortium	AA260035
1481 E	9 0216AMO8	cell division cycle 45 homolog (S. cerevisiae)-like	747922 IMAGE Consortium	AA260911
1482 E	11 0216AMO8	ESTs	749694 IMAGE Consortium	AA388758
1483 G	1 0216AMO8	ESTs	722279 IMAGE Consortium	AA254084
1484 G	3 0216AMO8	ESTs	723212 IMAGE Consortium	AI463752
1485 G	5 0216AMO8	ESTs	722822 IMAGE Consortium	AI608134
1486 G	7 0216AMO8	erythrocyte protein band 4.2	749886 IMAGE Consortium	AA388664
1487 G	9 0216AMO8	ESTs	748656 IMAGE Consortium	AA269742
1488 G	11 0216AMO8	nasal embryonic LHRH factor	749711 IMAGE Consortium	AA388971
1489 A	1 021DAMO9	EST	750134 IMAGE Consortium	AI662823
1490 A	3 021DAMO9	ESTs, Highly similar to ZINC FINGER PROTEIN 42 [Homo sapiens]	335095 IMAGE Consortium	W17956
1491 A	5 021DAMO9	protein kinase inhibitor, alpha	334835 IMAGE Consortium	AI391322
1492 A	7 021DAMO9	doublecortin	736852 IMAGE Consortium	AA271274
1493 A	9 021DAMO9	ESTs, Moderately similar to AF163315_1 putative N-acetyltransferase Camello 2 [M.musculus]	750414 IMAGE Consortium	AI508455
1494 A	11 021DAMO9	ESTs	751376 IMAGE Consortium	AA395924
1495 C	1 021DAMO9	ESTs	750175 IMAGE Consortium	AI606527
1496 C	3 021DAMO9	ESTs	335139 IMAGE Consortium	AI390109
1497 C	5 021DAMO9	crystallin, gamma D	334895 IMAGE Consortium	W36490
1498 C	7 021DAMO9	latent transforming growth factor beta binding protein 1	747086 IMAGE Consortium	AI607704
1499 C	9 021DAMO9	ESTs, Highly similar to katanin p80 subunit [H.sapiens]	750444 IMAGE Consortium	AA390012
1500 C	11 021DAMO9	ESTs	750674 IMAGE Consortium	AA290113

1501 E	1	021DAMO9	ESTs	750208	IMAGE Consortium	AI606530
1502 E	3	021DAMO9	ESTs	334414	IMAGE Consortium	W18280
1503 E	5	021DAMO9	low density lipoprotein-related protein 5	334928	IMAGE Consortium	AI391331
1504 E	7	021DAMO9	ESTs, Moderately similar to TETRASPAN MEMBRANE PROTEIN IL-TMP [Homo sapiens]	737676	IMAGE Consortium	AA277642
1505 E	9	021DAMO9	ESTs	751186	IMAGE Consortium	AI508692
1506 E	11	021DAMO9	ESTs, Moderately similar to unnamed protein product [H.sapiens]	750698	IMAGE Consortium	AI508487
1507 G	1	021DAMO9	wingless-related MMTV integration site 7B	334147	IMAGE Consortium	W37044
1508 G	3	021DAMO9	ESTs	334464	IMAGE Consortium	AI391308
1509 G	5	021DAMO9	ESTs	735645	IMAGE Consortium	AA270217
1510 G	7	021DAMO9	ESTs	736977	IMAGE Consortium	AI466334
1511 G	9	021DAMO9	ESTs	750496	IMAGE Consortium	AI508468
1512 G	11	021DAMO9	ESTs	750704	IMAGE Consortium	AI508488
1513 A	1	021XAMOA	ESTs	751618	IMAGE Consortium	AI508741
1514 A	3	021XAMOA	ESTs	751856	IMAGE Consortium	AA396236
1515 A	5	021XAMOA	ESTs, Moderately similar to putative RNA binding protein KOC [H.sapiens]	750335	IMAGE Consortium	AI606545
1516 A	7	021XAMOA	ESTs	752468	IMAGE Consortium	AI508591
1517 A	9	021XAMOA	EST AF007010	762414	IMAGE Consortium	AA274135
1518 A	11	021XAMOA	ESTs, Highly similar to KIAA1008 protein [H.sapiens]	764190	IMAGE Consortium	AA285792
1519 C	1	021XAMOA	COP9 (constitutive photomorphogenic), subunit 7b (Arabidopsis)	751628	IMAGE Consortium	AA389979
1520 C	3	021XAMOA	ESTs, Moderately similar to S37032 gene LL5 protein - rat [R.norvegicus]	751976	IMAGE Consortium	AI508545
1521 C	5	021XAMOA	ESTs	750362	IMAGE Consortium	AI606549
1522 C	7	021XAMOA	ESTs	752504	IMAGE Consortium	AA290360
1523 C	9	021XAMOA	ESTs	762492	IMAGE Consortium	AI595040
1524 C	11	021XAMOA	ESTs	764196	IMAGE Consortium	AA285795
1525 E	1	021XAMOA	ESTs	751630	IMAGE Consortium	AA389980
1526 E	3	021XAMOA	ESTs	749973	IMAGE Consortium	AA388948
1527 E	5	021XAMOA	tumor necrosis factor receptor superfamily, member 18	764764	IMAGE Consortium	AA274169
1528 E	7	021XAMOA	origin recognition complex, subunit 5 homolog (S. cerevisiae)	752510	IMAGE Consortium	AA290362
1529 E	9	021XAMOA	ESTs	762568	IMAGE Consortium	AI595044
1530 E	11	021XAMOA	ESTs	764206	IMAGE Consortium	AI606565
1531 G	1	021XAMOA	ESTs	750917	IMAGE Consortium	AI508512
1532 G	3	021XAMOA	ESTs, Moderately similar to KIAA0748 protein [H.sapiens]	750239	IMAGE Consortium	AI606535
1533 G	5	021XAMOA	ESTs	752182	IMAGE Consortium	AI508555
1534 G	7	021XAMOA	ESTs, Highly similar to UBIQUITIN-CONJUGATING ENZYME E2-17 KD [Caenorhabditis elegans]	752516	IMAGE Consortium	AI508600
1535 G	9	021XAMOA	ESTs	764089	IMAGE Consortium	AA285990
1536 G	11	021XAMOA	ESTs	762870	IMAGE Consortium	AI595060

1537 A	1 0214AMOB	DNA segment, Chr 5, ERATO Doi 431, expressed	778307 IMAGE Consortium	AA413897
1538 A	3 0214AMOB	ESTs	778484 IMAGE Consortium	AA413513
1539 A	5 0214AMOB	ESTs	778696 IMAGE Consortium	AA413749
1540 A	7 0214AMOB	potassium voltage-gated channel, Isk-related subfamily, gene 4	777202 IMAGE Consortium	AI508866
1541 A	9 0214AMOB	ESTs	777553 IMAGE Consortium	AA276690
1542 A	11 0214AMOB	ESTs	765347 IMAGE Consortium	AA275171
1543 C	1 0214AMOB	protein tyrosine phosphatase, receptor type, R	778314 IMAGE Consortium	AA413865
1544 C	3 0214AMOB	synaptotagmin 11	779211 IMAGE Consortium	AA445762
1545 C	5 0214AMOB	ATP synthase, H <sup>+</sup> transporting, mitochondrial F1F0 complex, subunit e	778733 IMAGE Consortium	AA413907
1546 C	7 0214AMOB	ESTs	777271 IMAGE Consortium	AA276812
1547 C	9 0214AMOB	receptor (calcitonin) activity modifying protein 2	777568 IMAGE Consortium	AA286544
1548 C	11 0214AMOB	ESTs	765364 IMAGE Consortium	AA275174
1549 E	1 0214AMOB	ESTs	778317 IMAGE Consortium	AA413869
1550 E	3 0214AMOB	trans-acting transcription factor 1	779239 IMAGE Consortium	AA445772
1551 E	5 0214AMOB	ESTs	778777 IMAGE Consortium	AA413918
1552 E	7 0214AMOB	ESTs, Weakly similar to growth factor receptor binding protein Grb10 [M.musculus]	777313 IMAGE Consortium	AA276909
1553 E	9 0214AMOB	ESTs	777767 IMAGE Consortium	AA286306
1554 E	11 0214AMOB	ESTs, Weakly similar to DBY protein [M.musculus]	765383 IMAGE Consortium	AA275127
1555 G	1 0214AMOB	Sin3-associated protein	778342 IMAGE Consortium	AA413873
1556 G	3 0214AMOB	ESTs, Weakly similar to unnamed protein product [H.sapiens]	778524 IMAGE Consortium	AA413701
1557 G	5 0214AMOB	E2F transcription factor 5	778782 IMAGE Consortium	AA413917
1558 G	7 0214AMOB	ESTs, Highly similar to KIAA1208 protein [H.sapiens]	777340 IMAGE Consortium	AI508964
1559 G	9 0214AMOB	ESTs	777802 IMAGE Consortium	AA286378
1560 G	11 0214AMOB	ESTs	765513 IMAGE Consortium	AA274740
1561 A	1 021BAMOC	ESTs	765959 IMAGE Consortium	AA277972
1562 A	3 021BAMOC	ESTs, Highly similar to ACTIVATOR 1 36 KD SUBUNIT [Homo sapiens]	779075 IMAGE Consortium	AA414079
1563 A	5 021BAMOC	transformed mouse 3T3 cell double minute 2	779350 IMAGE Consortium	AA414811
1564 A	7 021BAMOC	Mus musculus mRNA for Deltex2, complete cds	779466 IMAGE Consortium	AA414407
1565 A	9 021BAMOC	ESTs, Moderately similar to RRE1_HUMAN RAS-RESPONSIVE ELEMENT BINDING PROTEIN 1 [H.sapiens]	779639 IMAGE Consortium	AA466934
1566 A	11 021BAMOC	tissue specific transplantation antigen P91A	791014 IMAGE Consortium	AA473949
1567 C	1 021BAMOC	lymphocyte antigen 6 complex, locus F	776357 IMAGE Consortium	AA285471
1568 C	3 021BAMOC	ESTs, Weakly similar to similar to C.elegans hypothetical protein CET01H8.1,CEC05C12.3,CEF54D1.5. similar to trp and trp-like proteins [H.sapiens]	779109 IMAGE Consortium	AA413829
1569 C	5 021BAMOC	ESTs	779366 IMAGE Consortium	AA414110
1570 C	7 021BAMOC	ESTs, Moderately similar to M2GD_RAT	779491 IMAGE Consortium	AA414490



DIMETHYLGLYCINE DEHYDROGENASE PRECURSOR  
[R.norvegicus]

1571 C	9 021BAMOC	t-complex testis-expressed 3	779807 IMAGE Consortium	AA414787
1572 C	11 021BAMOC	ESTs	791048 IMAGE Consortium	AA415216
1573 E	1 021BAMOC	ESTs	777119 IMAGE Consortium	AA276507
1574 E	3 021BAMOC	histocompatibility 2, T region locus 9	779124 IMAGE Consortium	AA445709
1575 E	5 021BAMOC	phosphatidylinositol transfer protein, beta	779371 IMAGE Consortium	AA414830
1576 E	7 021BAMOC	ESTs	779597 IMAGE Consortium	AA413698
1577 E	9 021BAMOC	NCK-associated protein 1	779833 IMAGE Consortium	AA414730
1578 E	11 021BAMOC	ESTs	791073 IMAGE Consortium	AA415182
1579 G	1 021BAMOC	ESTs, Weakly similar to fatty acid synthase [R.norvegicus]	776443 IMAGE Consortium	AA278088
1580 G	3 021BAMOC	inhibitor of growth family, member 1	779172 IMAGE Consortium	AA445724
1581 G	5 021BAMOC	ESTs	779389 IMAGE Consortium	AA414045
1582 G	7 021BAMOC	SWI/SNF related, matrix associated, actin dependent regulator of chromatin, subfamily c, member 1	779616 IMAGE Consortium	AA413817
1583 G	9 021BAMOC	ESTs	779899 IMAGE Consortium	AA414877
1584 G	11 021BAMOC	DNA segment, Chr 13, ERATO Doi 608, expressed	791094 IMAGE Consortium	AA415207
1585 A	1 021IAMOD	constitutive photomorphogenic protein 1 (Arabidopsis)	791160 IMAGE Consortium	AA415271
1586 A	3 021IAMOD	U2 small nuclear ribonucleoprotein auxiliary factor (U2AF), 35 kDa, related sequence 2	790782 IMAGE Consortium	AA388471
1587 A	5 021IAMOD	ESTs	792569 IMAGE Consortium	AA416355
1588 A	7 021IAMOD	B-cell receptor-associated protein 31	792673 IMAGE Consortium	AA416357
1589 A	9 021IAMOD	ESTs, Weakly similar to UDP-N-acetylglucosamine pyrophosphorylase [H.sapiens]	792960 IMAGE Consortium	AA415344
1590 A	11 021IAMOD	early development regulator 2 (homolog of polyhomeotic 2)	793153 IMAGE Consortium	AA438004
1591 C	1 021IAMOD	ESTs, Highly similar to arginine-tRNA-protein transferase 1-2p [M.musculus]	791166 IMAGE Consortium	AA415294
1592 C	3 021IAMOD	ESTs	790886 IMAGE Consortium	AA388538
1593 C	5 021IAMOD	RAS-related C3 botulinum substrate 1	792593 IMAGE Consortium	AA415224
1594 C	7 021IAMOD	LIM motif-containing protein kinase 2	792690 IMAGE Consortium	AA416356
1595 C	9 021IAMOD	farnesyltransferase, CAAX box, alpha	792988 IMAGE Consortium	AA415472
1596 C	11 021IAMOD	prefoldin 2	793158 IMAGE Consortium	AA437996
1597 E	1 021IAMOD	peroxisome proliferator activated receptor binding protein	791214 IMAGE Consortium	AA415615
1598 E	3 021IAMOD	ESTs	790922 IMAGE Consortium	AA415329
1599 E	5 021IAMOD	ESTs, Weakly similar to CA11_RAT COLLAGEN ALPHA 1(I) CHAIN [R.norvegicus]	792598 IMAGE Consortium	AA416468
1600 E	7 021IAMOD	ESTs	791973 IMAGE Consortium	AA414465
1601 E	9 021IAMOD	Down syndrome critical region gene c	792016 IMAGE Consortium	AA414610
1602 E	11 021IAMOD	ESTs	793180 IMAGE Consortium	AA438010
1603 G	1 021IAMOD	DNA segment, Chr 1, Lubeck 1	791228 IMAGE Consortium	AA415621
1604 G	3 021IAMOD	ESTs, Highly similar to rA9 [R.norvegicus]	790929 IMAGE Consortium	AA415317
1605 G	5 021IAMOD	DNA segment, Chr 5, ERATO Doi 363, expressed	791896 IMAGE Consortium	AA414419

1606 G	7 021IAMOD	gene rich cluster, C8 gene	792743 IMAGE Consortium	AA416441
1607 G	9 021IAMOD	farnesyl diphosphate farnesyl transferase 1	793009 IMAGE Consortium	AA416273
1608 G	11 021IAMOD	cullin 3	791758 IMAGE Consortium	AA414532
1609 A	1 021PAMOE	ESTs	791359 IMAGE Consortium	AA415428
1610 A	3 021PAMOE	ESTs	791532 IMAGE Consortium	AA437441
1611 A	5 021PAMOE	ESTs, Weakly similar to B0507.2 gene product [C.elegans]	793215 IMAGE Consortium	AA437899
1612 A	7 021PAMOE	Cdc42 GTPase-inhibiting protein	792931 IMAGE Consortium	AA415781
1613 A	9 021PAMOE	ESTs	793955 IMAGE Consortium	AA422743
1614 A	11 021PAMOE	eukaryotic translation initiation factor 4, gamma 2	793122 IMAGE Consortium	AA437871
1615 C	1 021PAMOE	heterogeneous nuclear ribonucleoprotein H1	791369 IMAGE Consortium	AA473972
1616 C	3 021PAMOE	methionine aminopeptidase 2; initiation factor 2-associated protein (p67)	791572 IMAGE Consortium	AA437457
1617 C	5 021PAMOE	dystrophin, muscular dystrophy	793296 IMAGE Consortium	AA437900
1618 C	7 021PAMOE	transforming growth factor beta regulated gene 4	792953 IMAGE Consortium	AA437710
1619 C	9 021PAMOE	ESTs, Moderately similar to KIAA1155 protein [H.sapiens]	793006 IMAGE Consortium	AA437721
1620 C	11 021PAMOE	OG9 homeobox gene	793126 IMAGE Consortium	AA437873
1621 E	1 021PAMOE	diaphanous homolog 3 (Drosophila)	791383 IMAGE Consortium	AA415402
1622 E	3 021PAMOE	inner centromere protein	791602 IMAGE Consortium	AA437572
1623 E	5 021PAMOE	ESTs, Highly similar to 140 KD NUCLEOLAR PHOSPHOPROTEIN [Rattus norvegicus]	793395 IMAGE Consortium	AA415905
1624 E	7 021PAMOE	ESTs, Highly similar to CB80_HUMAN 80 KDA NUCLEAR CAP BINDING PROTEIN [H.sapiens]	792975 IMAGE Consortium	AA437623
1625 E	9 021PAMOE	ESTs, Weakly similar to ACYL CARRIER PROTEIN, MITOCHONDRIAL PRECURSOR [Neurospora crassa]	793020 IMAGE Consortium	AA437633
1626 E	11 021PAMOE	EST AA408438	793141 IMAGE Consortium	AA437988
1627 G	1 021PAMOE	ESTs, Moderately similar to TRI3_HUMAN THYROID RECEPTOR INTERACTING PROTEIN 3 [H.sapiens]	791404 IMAGE Consortium	AA416367
1628 G	3 021PAMOE	zinc finger protein 216	791607 IMAGE Consortium	AA437578
1629 G	5 021PAMOE	ESTs, Weakly similar to cAMP-dependent Rap1 guanine-nucleotide exchange factor [M.musculus]	793552 IMAGE Consortium	AA415729
1630 G	7 021PAMOE	ADP-ribosylation-like 4	792995 IMAGE Consortium	AA437628
1631 G	9 021PAMOE	DNA segment, Chr 1, University of California at Los Angeles 2	793029 IMAGE Consortium	AA437729
1632 G	11 021PAMOE	ESTs, Highly similar to AF132969_1 CGI-35 protein [H.sapiens]	793164 IMAGE Consortium	AA437999
1633 A	1 021WAMOF	ESTs, Highly similar to putative twelve transmembrane domain protein [M.musculus]	351961 IMAGE Consortium	W40842
1634 A	3 021WAMOF	ESTs, Weakly similar to F-box protein FBX6b [M.musculus]	355415 IMAGE Consortium	AI427276
1635 A	5 021WAMOF	ESTs	350274 IMAGE Consortium	W34757
1636 A	7 021WAMOF	ESTs	350787 IMAGE Consortium	AI450514
1637 A	9 021WAMOF	ESTs	807482 IMAGE Consortium	AI597421
1638 A	11 021WAMOF	ESTs	793822 IMAGE Consortium	AA422639

1639 C	1 021WAMOF	seizure related gene 6	352864 IMAGE Consortium	W36045
1640 C	3 021WAMOF	glial cell line derived neurotrophic factor	354767 IMAGE Consortium	W45748
1641 C	5 021WAMOF	hematopoietic zinc finger	351007 IMAGE Consortium	W35084
1642 C	7 021WAMOF	ESTs	353064 IMAGE Consortium	AI430714
1643 C	9 021WAMOF	ESTs	807515 IMAGE Consortium	AI597423
1644 C	11 021WAMOF			
1645 E	1 021WAMOF	ESTs, Weakly similar to PEX7_MOUSE PEROXISOMAL TARGETING SIGNAL 2 RECEPTOR [M.musculus]	352901 IMAGE Consortium	AI430704
1646 E	3 021WAMOF	ESTs	354825 IMAGE Consortium	W44297
1647 E	5 021WAMOF	ESTs	350347 IMAGE Consortium	AI391178
1648 E	7 021WAMOF	DNA segment, Chr 15, ERATO Doi 781, expressed	350956 IMAGE Consortium	AI450520
1649 E	9 021WAMOF	ESTs	806804 IMAGE Consortium	AA403816
1650 E	11 021WAMOF	ESTs	793934 IMAGE Consortium	AA422670
1651 G	1 021WAMOF	ESTs	355136 IMAGE Consortium	W48261
1652 G	3 021WAMOF	ESTs	349758 IMAGE Consortium	AI391100
1653 G	5 021WAMOF	ESTs	350477 IMAGE Consortium	AI450490
1654 G	7 021WAMOF	ESTs	353172 IMAGE Consortium	AI430723
1655 G	9 021WAMOF	ESTs, Moderately similar to dJ1043E3.1 [H.sapiens]	806816 IMAGE Consortium	AA403657
1656 G	11 021WAMOF	ESTs	808776 IMAGE Consortium	AI596889
1657 A	1 0213AMOG	ESTs, Moderately similar to DUAL SPECIFICITY PROTEIN PHOSPHATASE PAC-1 [Homo sapiens]	804279 IMAGE Consortium	AA474661
1658 A	3 0213AMOG	ESTs	803834 IMAGE Consortium	AA422762
1659 A	5 0213AMOG	ESTs	833279 IMAGE Consortium	AA432457
1660 A	7 0213AMOG	ESTs	836058 IMAGE Consortium	AA426736
1661 A	9 0213AMOG	T-cell leukemia/lymphoma 1B, 3	835858 IMAGE Consortium	AA422206
1662 A	11 0213AMOG	ESTs, Weakly similar to IG DELTA CHAIN C REGION MEMBRANE-BOUND FORM [Mus musculus]	831907 IMAGE Consortium	AA444443
1663 C	1 0213AMOG	ESTs, Moderately similar to hypothetical protein [H.sapiens]	804338 IMAGE Consortium	AA474689
1664 C	3 0213AMOG	ESTs, Highly similar to unnamed protein product [H.sapiens]	803854 IMAGE Consortium	AA422977
1665 C	5 0213AMOG	ESTs, Weakly similar to COP1 protein [M.musculus]	833291 IMAGE Consortium	AA465851
1666 C	7 0213AMOG	ESTs, Highly similar to hypothetical protein [H.sapiens]	836077 IMAGE Consortium	AA426777
1667 C	9 0213AMOG	ESTs, Weakly similar to A48830 probable transcription regulator NT fin12 - mouse [M.musculus]	835901 IMAGE Consortium	AA422223
1668 C	11 0213AMOG	ESTs, Weakly similar to CG1172 gene product [D.melanogaster]	831954 IMAGE Consortium	AI549740
1669 E	1 0213AMOG	ESTs	803614 IMAGE Consortium	AA422758
1670 E	3 0213AMOG	ESTs, Moderately similar to A53770 growth factor-responsive protein, vascular smooth muscle - rat [R.norvegicus]	803875 IMAGE Consortium	AA427320
1671 E	5 0213AMOG	ESTs, Moderately similar to KIAA0615 protein [H.sapiens]	833327 IMAGE Consortium	AA432470
1672 E	7 0213AMOG	ESTs	836095 IMAGE Consortium	AA426810

1673 E	9 0213AMOG	rod outer segment membrane protein 1	831723 IMAGE Consortium	AA444932
1674 E	11 0213AMOG	calsequestrin 1	818194 IMAGE Consortium	AA435278
1675 G	1 0213AMOG	ESTs	803624 IMAGE Consortium	AA422793
1676 G	3 0213AMOG	ESTs	803894 IMAGE Consortium	AA435241
1677 G	5 0213AMOG	ESTs	833392 IMAGE Consortium	AA432563
1678 G	7 0213AMOG	DNA segment, Chr 8, ERATO Doi 2, expressed	836124 IMAGE Consortium	AA426892
1679 G	9 0213AMOG	Mus musculus domesticus X chromosome region similar to human DXS963E	831638 IMAGE Consortium	AI549697
1680 G	11 0213AMOG	Notch gene homolog 3, (Drosophila)	818383 IMAGE Consortium	AA434836
1681 A	1 021AAMOH	ESTs	818673 IMAGE Consortium	AI596306
1682 A	3 021AAMOH	ESTs	820479 IMAGE Consortium	AI467582
1683 A	5 021AAMOH	ESTs	836434 IMAGE Consortium	AA426902
1684 A	7 021AAMOH	ESTs	847179 IMAGE Consortium	AA420050
1685 A	9 021AAMOH	ESTs	847445 IMAGE Consortium	AI552167
1686 A	11 021AAMOH	ESTs	849179 IMAGE Consortium	AI552235
1687 C	1 021AAMOH	ESTs	818894 IMAGE Consortium	AA434887
1688 C	3 021AAMOH	ESTs	820928 IMAGE Consortium	AI552317
1689 C	5 021AAMOH	Bcl2-like 10	836445 IMAGE Consortium	AA426934
1690 C	7 021AAMOH	ESTs	846453 IMAGE Consortium	AA433503
1691 C	9 021AAMOH	ESTs	846785 IMAGE Consortium	AA422239
1692 C	11 021AAMOH	ESTs		
1693 E	1 021AAMOH	ESTs	820170 IMAGE Consortium	AI551592
1694 E	3 021AAMOH	colony stimulating factor 3 receptor (granulocyte)	849416 IMAGE Consortium	AA457864
1695 E	5 021AAMOH	ESTs	836511 IMAGE Consortium	AA426838
1696 E	7 021AAMOH	ESTs	847201 IMAGE Consortium	AA420091
1697 E	9 021AAMOH	ESTs	848256 IMAGE Consortium	AA445521
1698 E	11 021AAMOH	ESTs	848613 IMAGE Consortium	AI552207
1699 G	1 021AAMOH	C-type (calcium dependent, carbohydrate recognition domain) lectin, superfamily member 6	820216 IMAGE Consortium	AA423158
1700 G	3 021AAMOH	ESTs, Weakly similar to weak similarity to Arabidopsis thaliana ubiquitin-like protein 8 [C.elegans]	849489 IMAGE Consortium	AI552277
1701 G	5 021AAMOH	ESTs	846160 IMAGE Consortium	AA415717
1702 G	7 021AAMOH	ESTs	846570 IMAGE Consortium	AA433581
1703 G	9 021AAMOH	ESTs	846894 IMAGE Consortium	AA433598
1704 G	11 021AAMOH	reversion-inducing-cysteine-rich protein with kazal motifs	849385 IMAGE Consortium	AI552258
1705 A	1 021HAMOI	amyloid beta (A4) precursor protein-binding, family B, member 1 interacting protein	849497 IMAGE Consortium	AA457914
1706 A	3 021HAMOI	gamma-glutamyl hydrolase	850244 IMAGE Consortium	AA451458
1707 A	5 021HAMOI	ESTs	850631 IMAGE Consortium	AA450992
1708 A	7 021HAMOI	ESTs, Highly similar to dJ323M22.2 [H.sapiens]	861100 IMAGE Consortium	AA571907
1709 A	9 021HAMOI	ESTs, Weakly similar to distal intestinal serine protease [M.musculus]	873598 IMAGE Consortium	AA472767
1710 A	11 021HAMOI	ESTs	874626 IMAGE Consortium	AA472455

1711 C	1 021HAMOI	ESTs	849522 IMAGE Consortium	AI481291
1712 C	3 021HAMOI	ESTs	851024 IMAGE Consortium	AA462521
1713 C	5 021HAMOI	EST	851376 IMAGE Consortium	AI481086
1714 C	7 021HAMOI	ESTs	851718 IMAGE Consortium	AA501085
1715 C	9 021HAMOI	cell cycle progression 2 protein	875065 IMAGE Consortium	AA472141
1716 C	11 021HAMOI	ESTs	873923 IMAGE Consortium	AA472082
1717 E	1 021HAMOI	ESTs	849778 IMAGE Consortium	AA450396
1718 E	3 021HAMOI	period homolog 3 (Drosophila)	850330 IMAGE Consortium	AA451527
1719 E	5 021HAMOI	ESTs	860102 IMAGE Consortium	AA500807
1720 E	7 021HAMOI	ESTs	860528 IMAGE Consortium	AA499950
1721 E	9 021HAMOI	ESTs	872984 IMAGE Consortium	AA466234
1722 E	11 021HAMOI	ESTs	873943 IMAGE Consortium	AA475186
1723 G	1 021HAMOI	ESTs	849812 IMAGE Consortium	AI608329
1724 G	3 021HAMOI	ESTs	850400 IMAGE Consortium	AI480949
1725 G	5 021HAMOI	ectonucleoside triphosphate diphosphohydrolase 6	850714 IMAGE Consortium	AA451048
1726 G	7 021HAMOI	ESTs	851892 IMAGE Consortium	AA575550
1727 G	9 021HAMOI	Mus musculus mRNA for MLTK alpha, complete cds	875235 IMAGE Consortium	AA472779
1728 G	11 021HAMOI	ESTs, Moderately similar to T00636 hypothetical protein F21856_2 - human [H.sapiens]	875462 IMAGE Consortium	AA475365
1729 A	1 021OAMOJ	EST	876530 IMAGE Consortium	AI481639
1730 A	3 021OAMOJ	ESTs, Moderately similar to HYPOTHETICAL 60.3 KD PROTEIN R08D7.7 IN CHROMOSOME III [Caenorhabditis elegans]	871962 IMAGE Consortium	AI614440
1731 A	5 021OAMOJ	leucocyte specific transcript 1	864318 IMAGE Consortium	AA458107
1732 A	7 021OAMOJ	phospholipase A2, group IV	891392 IMAGE Consortium	AA510727
1733 A	9 021OAMOJ	ESTs	876397 IMAGE Consortium	AA475824
1734 A	11 021OAMOJ	ESTs	889512 IMAGE Consortium	AA498682
1735 C	1 021OAMOJ	ESTs	875860 IMAGE Consortium	AA575202
1736 C	3 021OAMOJ	sperm autoantigenic protein 17	864765 IMAGE Consortium	AA444790
1737 C	5 021OAMOJ	ESTs	870949 IMAGE Consortium	AA462184
1738 C	7 021OAMOJ	ESTs	891470 IMAGE Consortium	AA511077
1739 C	9 021OAMOJ	ESTs	876477 IMAGE Consortium	AA475937
1740 C	11 021OAMOJ	insulin receptor substrate 2	888884 IMAGE Consortium	AA466295
1741 E	1 021OAMOJ	ESTs, Weakly similar to KIAA0982 protein [H.sapiens]	875975 IMAGE Consortium	AA623092
1742 E	3 021OAMOJ	ESTs, Highly similar to MYOSIN HEAVY CHAIN, SMOOTH MUSCLE ISOFORM [Homo sapiens]	864819 IMAGE Consortium	AA445019
1743 E	5 021OAMOJ	G0/G1 switch gene 2	864447 IMAGE Consortium	AA458241
1744 E	7 021OAMOJ	ESTs, Moderately similar to unnamed protein product [H.sapiens]	891490 IMAGE Consortium	AA511088
1745 E	9 021OAMOJ	ESTs	876694 IMAGE Consortium	AA499283
1746 E	11 021OAMOJ	RAS-like protein expressed in many tissues	888954 IMAGE Consortium	AA466324
1747 G	1 021OAMOJ	reduced in osteosclerosis transporter	872149 IMAGE Consortium	AI614454
1748 G	3 021OAMOJ	ESTs	871023 IMAGE Consortium	AA462335

1749 G	5 021OAMOJ			
1750 G	7 021OAMOJ	ESTs	876211 IMAGE Consortium	AA475644
1751 G	9 021OAMOJ	ESTs	876859 IMAGE Consortium	AA497477
1752 G	11 021OAMOJ	ESTs	901933 IMAGE Consortium	AI643612
1753 A	1 021VAMOK	ESTs	905127 IMAGE Consortium	AA518686
1754 A	3 021VAMOK	ESTs	903983 IMAGE Consortium	AA522090
1755 A	5 021VAMOK	ESTs	904995 IMAGE Consortium	AA518257
1756 A	7 021VAMOK	matrilin 2	901991 IMAGE Consortium	AA518595
1757 A	9 021VAMOK	ESTs, Moderately similar to SELP_MOUSE SELENOPROTEIN P PRECURSOR [M.musculus]	948910 IMAGE Consortium	AA544955
1758 A	11 021VAMOK	Mus musculus mRNA for putative nuclear factor	959680 IMAGE Consortium	AA546745
1759 C	1 021VAMOK			
1760 C	3 021VAMOK	ESTs	905440 IMAGE Consortium	AA529131
1761 C	5 021VAMOK	ESTs, Weakly similar to FIBRONECTIN RECEPTOR ALPHA SUBUNIT PRECURSOR [Mus musculus]	903107 IMAGE Consortium	AA521693
1762 C	7 021VAMOK	lymphocyte antigen 64	904160 IMAGE Consortium	AA518009
1763 C	9 021VAMOK	ESTs	948928 IMAGE Consortium	AA544972
1764 C	11 021VAMOK	ESTs, Highly similar to unnamed protein product [H.sapiens]	959727 IMAGE Consortium	AA546866
1765 E	1 021VAMOK	ESTs	903735 IMAGE Consortium	AI553195
1766 E	3 021VAMOK	ESTs	905515 IMAGE Consortium	AA529107
1767 E	5 021VAMOK	lysyl oxidase-like 2	903231 IMAGE Consortium	AA522066
1768 E	7 021VAMOK	procollagen-proline, 2-oxoglutarate 4-dioxygenase (proline 4-hydroxylase), alpha 1 polypeptide	902719 IMAGE Consortium	AA518751
1769 E	9 021VAMOK	ESTs, Moderately similar to AF151866_1 CGI-108 protein [H.sapiens]	949801 IMAGE Consortium	AA537516
1770 E	11 021VAMOK	ATP-binding cassette, sub-family A (ABC1), member 3	959738 IMAGE Consortium	AA546905
1771 G	1 021VAMOK	ESTs	904472 IMAGE Consortium	AI553238
1772 G	3 021VAMOK	ESTs	904867 IMAGE Consortium	AA522153
1773 G	5 021VAMOK	ESTs, Highly similar to unnamed protein product [H.sapiens]	902529 IMAGE Consortium	AA518515
1774 G	7 021VAMOK	ESTs	949054 IMAGE Consortium	AA543902
1775 G	9 021VAMOK	ESTs	959539 IMAGE Consortium	AA546498
1776 G	11 021VAMOK	ESTs, Highly similar to Y174_HUMAN HYPOTHETICAL PROTEIN KIAA0174 [H.sapiens]	959756 IMAGE Consortium	AA546912
1777 A	1 0212AMOL	ESTs	959909 IMAGE Consortium	AA547137
1778 A	3 0212AMOL	homeo box C9	367538 IMAGE Consortium	W54619
1779 A	5 0212AMOL	peripheral myelin protein, 22 kDa	368778 IMAGE Consortium	W56983
1780 A	7 0212AMOL	ESTs	921930 IMAGE Consortium	AA512148
1781 A	9 0212AMOL			
1782 A	11 0212AMOL	ESTs	944882 IMAGE Consortium	AA545415
1783 C	1 0212AMOL	ESTs, Weakly similar to HuEMAP [H.sapiens]	960026 IMAGE Consortium	AA547257
1784 C	3 0212AMOL	ESTs, Highly similar to TRANSCRIPTION FACTOR IIIA	368286 IMAGE Consortium	W54185

		[ <i>Xenopus laevis</i> ]		
1785 C	5 0212AMOL	ESTs, Highly similar to PiUS [ <i>R.norvegicus</i> ]	368799 IMAGE Consortium	W53344
1786 C	7 0212AMOL	interleukin 6 signal transducer	935452 IMAGE Consortium	AA543149
1787 C	9 0212AMOL	ESTs, Highly similar to KIAA0398 [ <i>H.sapiens</i> ]	944451 IMAGE Consortium	AA545082
1788 C	11 0212AMOL	ESTs	960902 IMAGE Consortium	AA549444
1789 E	1 0212AMOL	integrin alpha V (Cd51)	960079 IMAGE Consortium	AA547343
1790 E	3 0212AMOL	ESTs, Highly similar to rA1 [ <i>R.norvegicus</i> ]	367698 IMAGE Consortium	W54651
1791 E	5 0212AMOL	ESTs	922185 IMAGE Consortium	AI550977
1792 E	7 0212AMOL	isovaleryl dehydrogenase precursor	935514 IMAGE Consortium	AA543454
1793 E	9 0212AMOL	ESTs	944514 IMAGE Consortium	AA545228
1794 E	11 0212AMOL	Mus musculus nuclear body associated kinase 1a (Nbak1) mRNA, alternatively spliced, complete cds	960951 IMAGE Consortium	AA549549
1795 G	1 0212AMOL	hect (homologous to the E6-AP (UBE3A) carboxyl terminus) domain and RCC1 (CHC1)-like domain (RLD) 2	960128 IMAGE Consortium	AA547373
1796 G	3 0212AMOL	minK-like protein	368452 IMAGE Consortium	W53688
1797 G	5 0212AMOL	nitric oxide synthase 2, inducible, macrophage	922250 IMAGE Consortium	AA512708
1798 G	7 0212AMOL	ESTs, Moderately similar to OB-RGRP [ <i>M.musculus</i> ]	935548 IMAGE Consortium	AA543557
1799 G	9 0212AMOL	receptor for MCF virus 1	944538 IMAGE Consortium	AA545117
1800 G	11 0212AMOL	Mus musculus Bat-4 gene, complete cds	959898 IMAGE Consortium	AA547111
1801 A	1 0219AMOM	ESTs	960066 IMAGE Consortium	AA547481
1802 A	3 0219AMOM	ESTs, Highly similar to hypothetical protein [ <i>H.sapiens</i> ]	961033 IMAGE Consortium	AA547169
1803 A	5 0219AMOM	ESTs	961320 IMAGE Consortium	AA547628
1804 A	7 0219AMOM	ESTs	960883 IMAGE Consortium	AA549368
1805 A	9 0219AMOM	ESTs	961975 IMAGE Consortium	AA672334
1806 A	11 0219AMOM	ESTs	961466 IMAGE Consortium	AA549265
1807 C	1 0219AMOM	ESTs, Weakly similar to envelope polyprotein [ <i>M.musculus</i> ]	960105 IMAGE Consortium	AA547361
1808 C	3 0219AMOM	ESTs	961110 IMAGE Consortium	AA547314
1809 C	5 0219AMOM	ESTs, Weakly similar to wizL [ <i>M.musculus</i> ]	960604 IMAGE Consortium	AA549590
1810 C	7 0219AMOM	CD2-associated proten	961694 IMAGE Consortium	AA549691
1811 C	9 0219AMOM	ESTs, Highly similar to exonuclease 1 homologue [ <i>M.musculus</i> ]	961991 IMAGE Consortium	AA672342
1812 C	11 0219AMOM	DNA segment, Chr 1, Lubeck 1	961503 IMAGE Consortium	AA549296
1813 E	1 0219AMOM	ESTs, Weakly similar to AF133093_9 protein with characteristics of a rhoGAP [ <i>M.musculus</i> ]	960117 IMAGE Consortium	AA547367
1814 E	3 0219AMOM	ESTs	960427 IMAGE Consortium	AA549221
1815 E	5 0219AMOM	ESTs	960630 IMAGE Consortium	AA549612
1816 E	7 0219AMOM	ring finger protein 12	961788 IMAGE Consortium	AA672290
1817 E	9 0219AMOM	cullin 3	961070 IMAGE Consortium	AA547180
1818 E	11 0219AMOM	ESTs, Weakly similar to strong similarity to the SNF2/RAD54 family of helicases [ <i>C.elegans</i> ]	961538 IMAGE Consortium	AA549392
1819 G	1 0219AMOM	ESTs, Highly similar to Cdc5-like protein [ <i>R.norvegicus</i> ]	960201 IMAGE Consortium	AA547602
1820 G	3 0219AMOM	ESTs, Weakly similar to putative WHSC1 protein	961178 IMAGE Consortium	AA547432

		[H.sapiens]		
1821 G	5	0219AMOM	ESTs	961374 IMAGE Consortium AA547667
1822 G	7	0219AMOM	ESTs	963217 IMAGE Consortium AA671581
1823 G	9	0219AMOM	ESTs	961102 IMAGE Consortium AA547206
1824 G	11	0219AMOM	thiamin pyrophosphokinase	1003004 IMAGE Consortium AA684442
1825 A	1	021GAMON	ligase I, DNA, ATP-dependent	1003055 IMAGE Consortium AA684487
1826 A	3	021GAMON	ESTs	1003378 IMAGE Consortium AA684161
1827 A	5	021GAMON	ESTs, Moderately similar to HYPOTHETICAL 27.1 KD PROTEIN CCE1-CAP1 INTERGENIC REGION [Saccharomyces cerevisiae]	1003488 IMAGE Consortium AA684338
1828 A	7	021GAMON	next to the Brca1	976400 IMAGE Consortium AA619774
1829 A	9	021GAMON	ESTs, Weakly similar to SCF complex protein cul-1 [M.musculus]	976490 IMAGE Consortium AA619940
1830 A	11	021GAMON	ESTs, Moderately similar to T00380 KIAA0637 protein - human [H.sapiens]	976637 IMAGE Consortium AA619752
1831 C	1	021GAMON	nuclear receptor-binding SET-domain protein 1	1003086 IMAGE Consortium AA684004
1832 C	3	021GAMON	ESTs, Moderately similar to ubiquitin protein ligase [M.musculus]	1003383 IMAGE Consortium AA684173
1833 C	5	021GAMON	ESTs	1003512 IMAGE Consortium AA684400
1834 C	7	021GAMON	methylenetetrahydrofolate dehydrogenase (NAD+ dependent), methenyltetrahydrofolate cyclohydrolase	976418 IMAGE Consortium AA619789
1835 C	9	021GAMON	spindlin	976497 IMAGE Consortium AA619954
1836 C	11	021GAMON	DNA segment, Chr 2, ERATO Doi 198, expressed	976648 IMAGE Consortium AA619901
1837 E	1	021GAMON	SEC23B (S. cerevisiae)	1003102 IMAGE Consortium AA684008
1838 E	3	021GAMON	ESTs, Weakly similar to SKIP_HUMAN NUCLEAR PROTEIN SKIP [H.sapiens]	1003410 IMAGE Consortium AA684185
1839 E	5	021GAMON	ESTs	1003523 IMAGE Consortium AA684413
1840 E	7	021GAMON	unc13 homolog (C. elegans) 1	976427 IMAGE Consortium AA619801
1841 E	9	021GAMON	nuclear receptor coactivator 4	976520 IMAGE Consortium AA619964
1842 E	11	021GAMON	SH3-domain GRB2-like B1 (endophilin)	976656 IMAGE Consortium AA619756
1843 G	1	021GAMON	ESTs, Weakly similar to hypothetical protein [H.sapiens]	1003206 IMAGE Consortium AA684090
1844 G	3	021GAMON	ESTs, Moderately similar to HYPOTHETICAL TRP-ASP REPEATS CONTAINING PROTEIN IN PGI1-KTR4 INTERGENIC REGION [Saccharomyces cerevisiae]	1003412 IMAGE Consortium AA684186
1845 G	5	021GAMON	ESTs	1003554 IMAGE Consortium AA684425
1846 G	7	021GAMON	ESTs	976443 IMAGE Consortium AA619863
1847 G	9	021GAMON	ESTs	976522 IMAGE Consortium AA619965
1848 G	11	021GAMON	cathepsin L	976659 IMAGE Consortium AA619763
1849 A	1	021NAMOO	ESTs	976705 IMAGE Consortium AA619821
1850 A	3	021NAMOO	ESTs	1026237 IMAGE Consortium AA616671
1851 A	5	021NAMOO	DNA segment, Chr 7, ERATO Doi 373, expressed	1003838 IMAGE Consortium AA675376
1852 A	7	021NAMOO	ESTs	1003995 IMAGE Consortium AA674781
1853 A	9	021NAMOO	ESTs	1051006 IMAGE Consortium AA596291



1854 A	11 021NAMOO			
1855 C	1 021NAMOO	EST AA536654	976716 IMAGE Consortium	AA619820
1856 C	3 021NAMOO	ESTs	1025562 IMAGE Consortium	AA608228
1857 C	5 021NAMOO	ESTs, Highly similar to SMALL NUCLEAR RIBONUCLEOPROTEIN SM D2 [Homo sapiens]	1003858 IMAGE Consortium	AA674492
1858 C	7 021NAMOO	nuclear factor I/C	1049407 IMAGE Consortium	AI594339
1859 C	9 021NAMOO	ESTs	1051132 IMAGE Consortium	AA597087
1860 C	11 021NAMOO	ESTs	1052192 IMAGE Consortium	AA596243
1861 E	1 021NAMOO	ubiquitination factor E4B (homologous to yeast UFD2)	976721 IMAGE Consortium	AA619922
1862 E	3 021NAMOO	ESTs	1026377 IMAGE Consortium	AA616841
1863 E	5 021NAMOO	phosphoribosyl pyrophosphate synthetase 1	1003885 IMAGE Consortium	AA674516
1864 E	7 021NAMOO	ESTs	1025032 IMAGE Consortium	AA793588
1865 E	9 021NAMOO	ESTs	1051191 IMAGE Consortium	AI429014
1866 E	11 021NAMOO	DNA segment, Chr 15, ERATO Doi 586, expressed	1052206 IMAGE Consortium	AI594455
1867 G	1 021NAMOO	ESTs	976729 IMAGE Consortium	AA619836
1868 G	3 021NAMOO	ESTs	1026393 IMAGE Consortium	AA648027
1869 G	5 021NAMOO	high mobility group box 3	1003901 IMAGE Consortium	AA674577
1870 G	7 021NAMOO	ESTs	1026000 IMAGE Consortium	AA616549
1871 G	9 021NAMOO	ESTs	1051237 IMAGE Consortium	AI429019
1872 G	11 021NAMOO	nuclear factor I/C	1050872 IMAGE Consortium	AA592566
1873 A	1 021UAMOP			
1874 A	3 021UAMOP	DNA polymerase epsilon	1053504 IMAGE Consortium	AA608457
1875 A	5 021UAMOP	ESTs, Weakly similar to brk kinase substrate [H.sapiens]	1069250 IMAGE Consortium	AA793059
1876 A	7 021UAMOP	vascular endothelial growth factor	1068845 IMAGE Consortium	AA793036
1877 A	9 021UAMOP	ESTs, Moderately similar to KIAA0830 protein [H.sapiens]	1078999 IMAGE Consortium	AA734890
1878 A	11 021UAMOP	ESTs	1054933 IMAGE Consortium	AA619052
1879 C	1 021UAMOP	ESTs	1051741 IMAGE Consortium	AI594422
1880 C	3 021UAMOP	ESTs	1052863 IMAGE Consortium	AA930522
1881 C	5 021UAMOP	ESTs	1068544 IMAGE Consortium	AI585466
1882 C	7 021UAMOP	ESTs	1079027 IMAGE Consortium	AI593606
1883 C	9 021UAMOP	ESTs, Weakly similar to BcDNA:LD21720 gene product [D.melanogaster]	1068048 IMAGE Consortium	AA764450
1884 C	11 021UAMOP	EST	1054956 IMAGE Consortium	AI551241
1885 E	1 021UAMOP	ESTs	1052582 IMAGE Consortium	AI593934
1886 E	3 021UAMOP	ESTs	1053650 IMAGE Consortium	AA638480
1887 E	5 021UAMOP	ESTs, Weakly similar to AAKG_MOUSE 5'-AMP-ACTIVATED PROTEIN KINASE, GAMMA-1 SUBUNIT [M.musculus]	1067834 IMAGE Consortium	AI536510
1888 E	7 021UAMOP	folate receptor 2 (fetal)	1068952 IMAGE Consortium	AA796142
1889 E	9 021UAMOP	ESTs, Highly similar to unnamed protein product [H.sapiens]	1068060 IMAGE Consortium	AA764455
1890 E	11 021UAMOP	ESTs	1064416 IMAGE Consortium	AI603997
1891 G	1 021UAMOP	DNA segment, Chr 11, Wayne State University 47,	1051884 IMAGE Consortium	AI594431

		expressed		
1892 G	3 021UAMOP	ESTs, Moderately similar to syntrophin 5 [H.sapiens]	1052987 IMAGE Consortium	AI594004
1893 G	5 021UAMOP	arachidonate 12-lipoxygenase	1068613 IMAGE Consortium	AA796544
1894 G	7 021UAMOP	ESTs, Highly similar to S43495 protein-tyrosine kinase [M.musculus]	1077771 IMAGE Consortium	AA796461
1895 G	9 021UAMOP	ESTs	1068076 IMAGE Consortium	AA764545
1896 G	11 021UAMOP	ESTs, Weakly similar to UOG1_MOUSE UOG-1 PROTEIN [M.musculus]	1066034 IMAGE Consortium	AA673894
1897 A	1 0211AMOP	pancreatitis-associated protein	1066145 IMAGE Consortium	AA619407
1898 A	3 0211AMOP	ESTs, Highly similar to dedicator of cyto-kinesis 1 [H.sapiens]	1066484 IMAGE Consortium	AA615918
1899 A	5 0211AMOP	ESTs	1066710 IMAGE Consortium	AA611815
1900 A	7 0211AMOP	thrombomodulin	1095181 IMAGE Consortium	AA870655
1901 A	9 0211AMOP	ESTs, Moderately similar to PLASMA-CELL MEMBRANE GLYCOPROTEIN PC-1 [Mus musculus]	1096179 IMAGE Consortium	AA871395
1902 A	11 0211AMOP	defensin related cryptdin, related sequence 10	1097475 IMAGE Consortium	AA869166
1903 C	1 0211AMOP	Hrs binding protein	1066282 IMAGE Consortium	AA611511
1904 C	3 0211AMOP	ESTs	1067290 IMAGE Consortium	AA763781
1905 C	5 0211AMOP	Mus musculus NORPEG-like protein mRNA, complete cds	1066898 IMAGE Consortium	AA616006
1906 C	7 0211AMOP	ESTs	1079170 IMAGE Consortium	AA823475
1907 C	9 0211AMOP	myosin, heavy polypeptide 2, skeletal muscle, adult	1094784 IMAGE Consortium	AA869609
1908 C	11 0211AMOP	phospholipase A2, group IIA (platelets, synovial fluid)	1096831 IMAGE Consortium	AA871838
1909 E	1 0211AMOP	receptor-associated protein of the synapse, 43 kDa	1065569 IMAGE Consortium	AA611702
1910 E	3 0211AMOP	ESTs	1066582 IMAGE Consortium	AA616089
1911 E	5 0211AMOP	ESTs, Moderately similar to rabphilin-3a related protein [R.norvegicus]	1066916 IMAGE Consortium	AA616014
1912 E	7 0211AMOP	ESTs	1079183 IMAGE Consortium	AA823488
1913 E	9 0211AMOP	ESTs	1096258 IMAGE Consortium	AA871451
1914 E	11 0211AMOP	T-cell expressing clone j6	389774 IMAGE Consortium	W71468
1915 G	1 0211AMOP	ESTs	1067041 IMAGE Consortium	AA764296
1916 G	3 0211AMOP	ESTs, Highly similar to hGCN5 [H.sapiens]	1067314 IMAGE Consortium	AA672656
1917 G	5 0211AMOP	ESTs, Highly similar to stanniocalcin-2 [M.musculus]	1094315 IMAGE Consortium	AI592910
1918 G	7 0211AMOP	ESTs	1079242 IMAGE Consortium	AA823511
1919 G	9 0211AMOP	ESTs, Weakly similar to ankyrin 3 [M.musculus]	1095730 IMAGE Consortium	AA869040
1920 G	11 0211AMOP	dickkopf homolog 1 (Xenopus laevis)	402616 IMAGE Consortium	W79975
1921 A	1 0218AMOR			
1922 A	3 0218AMOR	ESTs, Moderately similar to DIAMINE ACETYLTRANSFERASE [Mus musculus]	402298 IMAGE Consortium	AI425770
1923 A	5 0218AMOR	N-terminal Asn amidase	390147 IMAGE Consortium	W70679
1924 A	7 0218AMOR	low density lipoprotein-related protein 6	1107396 IMAGE Consortium	AA647231
1925 A	9 0218AMOR	Public domain EST	1107868 IMAGE Consortium	
1926 A	11 0218AMOR	ESTs, Weakly similar to AF089896_1 topoisomerase-	1125009 IMAGE Consortium	AA692960

		related function protein [H.sapiens]		
1927 C	1 0218AMOR	Public domain EST	401306 IMAGE Consortium	
1928 C	3 0218AMOR			
1929 C	5 0218AMOR	ESTs	391084 IMAGE Consortium	AI430193
1930 C	7 0218AMOR	ESTs	1107454 IMAGE Consortium	AA623620
1931 C	9 0218AMOR	B-cell CLL/lymphoma 7C	1107920 IMAGE Consortium	AA623781
1932 C	11 0218AMOR	AF1q protein	1125035 IMAGE Consortium	AA692978
1933 E	1 0218AMOR	EST	401350 IMAGE Consortium	AI413446
1934 E	3 0218AMOR	ESTs	402319 IMAGE Consortium	W78484
1935 E	5 0218AMOR	mitogen-activated protein kinase 7	391129 IMAGE Consortium	AI430199
1936 E	7 0218AMOR	ESTs	1106802 IMAGE Consortium	AI641979
1937 E	9 0218AMOR	ubiquitously transcribed tetratricopeptide repeat gene, X chromosome	1107961 IMAGE Consortium	AA623853
1938 E	11 0218AMOR	DNA segment, Chr 1, DA Underhill 7, expressed	1125106 IMAGE Consortium	AA693056
1939 G	1 0218AMOR	ESTs	400641 IMAGE Consortium	AI425717
1940 G	3 0218AMOR	mitogen activated protein kinase 14	401641 IMAGE Consortium	W78296
1941 G	5 0218AMOR	vertebrate homolog of C. elegans Lin-7 type 2	391220 IMAGE Consortium	AI430202
1942 G	7 0218AMOR	procollagen, type V, alpha 2	1107566 IMAGE Consortium	AA623765
1943 G	9 0218AMOR	defensin related cryptdin 3	1097514 IMAGE Consortium	AA869181
1944 G	11 0218AMOR	ESTs	1125270 IMAGE Consortium	AA693297
1945 A	1 021FAMOS	ESTs, Highly similar to PKL2_RAT PROTEIN KINASE C-LIKE 2 [R.norvegicus]	1126051 IMAGE Consortium	AA691164
1946 A	3 021FAMOS	ESTs	1134315 IMAGE Consortium	AA675602
1947 A	5 021FAMOS	bone morphogenic protein receptor, type II (serine/threonine kinase)	1134592 IMAGE Consortium	AA681180
1948 A	7 021FAMOS	ESTs, Highly similar to AF161554_1 HSPC069 [H.sapiens]	1137160 IMAGE Consortium	AA689735
1949 A	9 021FAMOS	ESTs	1137420 IMAGE Consortium	AI585782
1950 A	11 021FAMOS	HS1 binding protein 3	1139235 IMAGE Consortium	AA655508
1951 C	1 021FAMOS	ESTs, Weakly similar to envelope polyprotein [M.musculus]	1125485 IMAGE Consortium	AA675207
1952 C	3 021FAMOS	ESTs, Highly similar to 0-44 PROTEIN [Rattus norvegicus]	1125615 IMAGE Consortium	AA824095
1953 C	5 021FAMOS	ESTs	1134624 IMAGE Consortium	AA681211
1954 C	7 021FAMOS	ESTs	1137247 IMAGE Consortium	AA688933
1955 C	9 021FAMOS	ESTs	1137485 IMAGE Consortium	AA689704
1956 C	11 021FAMOS	ESTs	1138552 IMAGE Consortium	AA688673
1957 E	1 021FAMOS	ESTs	1134217 IMAGE Consortium	AA675455
1958 E	3 021FAMOS	f-box and WD-40 domain protein 2	1125652 IMAGE Consortium	AA824159
1959 E	5 021FAMOS	ESTs, Weakly similar to Edp1 protein [M.musculus]	1125977 IMAGE Consortium	AA691011
1960 E	7 021FAMOS	ESTs	1138036 IMAGE Consortium	AA691584
1961 E	9 021FAMOS	protein tyrosine phosphatase, receptor type, O	1137535 IMAGE Consortium	AA689770
1962 E	11 021FAMOS	ESTs	1137835 IMAGE Consortium	AA795130

1963 G	1	021FAMOS	ESTs	1134245	IMAGE Consortium	AA675478
1964 G	3	021FAMOS	ESTs	1134450	IMAGE Consortium	AA681258
1965 G	5	021FAMOS	ESTs	1136085	IMAGE Consortium	AA755175
1966 G	7	021FAMOS	semaF cytoplasmic domain associated protein 2	1138054	IMAGE Consortium	AA691592
1967 G	9	021FAMOS	ESTs	1152859	IMAGE Consortium	AI592441
1968 G	11	021FAMOS	mucin 3, intestinal	1138580	IMAGE Consortium	AA688695
1969 A	1	021MAMOT	hyaluronidase 1	1138827	IMAGE Consortium	AA688635
1970 A	3	021MAMOT	ESTs, Weakly similar to beta,beta-carotene 15,15'-dioxygenase [M.musculus]	1165901	IMAGE Consortium	AA710758
1971 A	5	021MAMOT	ESTs	1162397	IMAGE Consortium	AA692034
1972 A	7	021MAMOT	ESTs	1164195	IMAGE Consortium	AA734249
1973 A	9	021MAMOT	engrailed 1	1182007	IMAGE Consortium	AA711901
1974 A	11	021MAMOT	ESTs, Weakly similar to A27353 collagen alpha 1(III) chain precursor - mouse [M.musculus]	1195287	IMAGE Consortium	AA754886
1975 C	1	021MAMOT	ESTs	1166005	IMAGE Consortium	AA710144
1976 C	3	021MAMOT	ESTs	1166728	IMAGE Consortium	AA710981
1977 C	5	021MAMOT	ESTs	1163148	IMAGE Consortium	AI587686
1978 C	7	021MAMOT	ESTs	1164241	IMAGE Consortium	AA734324
1979 C	9	021MAMOT	ESTs	1180648	IMAGE Consortium	AA718224
1980 C	11	021MAMOT	ESTs, Highly similar to organic cation transporter-like protein 2 [M.musculus]	1195308	IMAGE Consortium	AA754896
1981 E	1	021MAMOT	ESTs	1164601	IMAGE Consortium	AA929439
1982 E	3	021MAMOT	ESTs	1166764	IMAGE Consortium	AA710290
1983 E	5	021MAMOT	ESTs	1154461	IMAGE Consortium	AA796923
1984 E	7	021MAMOT	CUG triplet repeat, RNA binding protein 1	1162848	IMAGE Consortium	AA770801
1985 E	9	021MAMOT	small inducible cytokine A7	1181707	IMAGE Consortium	AA711435
1986 E	11	021MAMOT	WNT1 inducible signaling pathway protein 2	1195358	IMAGE Consortium	AA754979
1987 G	1	021MAMOT	DNA segment, Chr 3, ERATO Doi 330, expressed	1165343	IMAGE Consortium	AA709728
1988 G	3	021MAMOT	ESTs	1166946	IMAGE Consortium	AA711038
1989 G	5	021MAMOT	ESTs	1164044	IMAGE Consortium	AA690608
1990 G	7	021MAMOT	ESTs	1162898	IMAGE Consortium	AI592477
1991 G	9	021MAMOT	follicistatin	1194016	IMAGE Consortium	AA726638
1992 G	11	021MAMOT	ESTs, Weakly similar to CG13089 gene product [D.melanogaster]	1193957	IMAGE Consortium	AA726565
1993 A	1	021TAMOU	ESTs	1194975	IMAGE Consortium	AA754967
1994 A	3	021TAMOU	Mus musculus strain 129/Ola papillary renal cell carcinoma-associated protein (Prcc) mRNA, complete cds	387764	IMAGE Consortium	W66780
1995 A	5	021TAMOU	ESTs	375167	IMAGE Consortium	W62706
1996 A	7	021TAMOU	disabled homolog 1 (Drosophila)	385192	IMAGE Consortium	AI893214
1997 A	9	021TAMOU	just another zinc finger protein	387319	IMAGE Consortium	AI430631
1998 A	11	021TAMOU	ESTs	1177898	IMAGE Consortium	AA718770
1999 C	1	021TAMOU	found in inflammatory zone 1	1195776	IMAGE Consortium	AA712003
2000 C	3	021TAMOU	peroxiredoxin 2	387840	IMAGE Consortium	W66866

2001 C	5 021TAMOU	ESTs, Moderately similar to CYTOCHROME P450 IVF3 [Homo sapiens]	373761 IMAGE Consortium	W64560
2002 C	7 021TAMOU	ESTs, Weakly similar to G9A [M.musculus]	385483 IMAGE Consortium	W64167
2003 C	9 021TAMOU	ESTs	1177250 IMAGE Consortium	AA789723
2004 C	11 021TAMOU	neurotrophin 3	1177923 IMAGE Consortium	AA718802
2005 E	1 021TAMOU	ESTs	1181048 IMAGE Consortium	AA688607
2006 E	3 021TAMOU	ESTs, Highly similar to GLUTATHIONE S-TRANSFERASE P [Homo sapiens]	387940 IMAGE Consortium	W70628
2007 E	5 021TAMOU	ESTs	374525 IMAGE Consortium	W64080
2008 E	7 021TAMOU	ESTs, Weakly similar to cytosolic sorting protein PACS-1a [R.norvegicus]	387012 IMAGE Consortium	AI413499
2009 E	9 021TAMOU	ESTs	1178073 IMAGE Consortium	AA718656
2010 E	11 021TAMOU	ESTs	1177995 IMAGE Consortium	AA718600
2011 G	1 021TAMOU	ESTs	1180565 IMAGE Consortium	AA681081
2012 G	3 021TAMOU	protein kinase C and casein kinase substrate in neurons 2	373221 IMAGE Consortium	W57231
2013 G	5 021TAMOU	ESTs	374566 IMAGE Consortium	W63829
2014 G	7 021TAMOU	vesicular inhibitory amino acid transporter	385606 IMAGE Consortium	AI893217
2015 G	9 021TAMOU	a disintegrin and metalloprotease domain 23	1178116 IMAGE Consortium	AA718688
2016 G	11 021TAMOU	interleukin 15	1225014 IMAGE Consortium	AA709787
2017 A	1 0210AMOV	ESTs, Moderately similar to tyrosine phosphatase [M.musculus]	1225160 IMAGE Consortium	AI428655
2018 A	3 0210AMOV	putative homeodomain transcription factor	1225550 IMAGE Consortium	AA762925
2019 A	5 0210AMOV	ESTs	1225780 IMAGE Consortium	AI428753
2020 A	7 0210AMOV	ESTs	1195889 IMAGE Consortium	AA717023
2021 A	9 0210AMOV	galactosamine (N-acetyl)-6-sulfate sulfatase	1230520 IMAGE Consortium	AA879799
2022 A	11 0210AMOV	CD22 antigen	1230835 IMAGE Consortium	AA880028
2023 C	1 0210AMOV	testis expressed gene 10	1225179 IMAGE Consortium	AA763832
2024 C	3 0210AMOV	transducin-like enhancer of split 2, homolog of Drosophila E(spl)	1226313 IMAGE Consortium	AI466119
2025 C	5 0210AMOV	ESTs	1195747 IMAGE Consortium	AA711943
2026 C	7 0210AMOV	embryonal Fyn-associated substrate	1195895 IMAGE Consortium	AA717026
2027 C	9 0210AMOV	ESTs, Weakly similar to unnamed protein product [H.sapiens]	1230550 IMAGE Consortium	AA879822
2028 C	11 0210AMOV	ribonuclease 1, pancreatic	1230895 IMAGE Consortium	AA790398
2029 E	1 0210AMOV	ESTs	1225217 IMAGE Consortium	AI428665
2030 E	3 0210AMOV	ESTs	1224925 IMAGE Consortium	AI428626
2031 E	5 0210AMOV	ESTs, Highly similar to I52257 episialin - mouse [M.musculus]	1195760 IMAGE Consortium	AA711984
2032 E	7 0210AMOV	ATP synthase, H <sup>+</sup> transporting, mitochondrial F1 complex, alpha subunit, isoform 1	1225990 IMAGE Consortium	AA739359
2033 E	9 0210AMOV	ESTs, Weakly similar to hypothetical protein [H.sapiens]	1230583 IMAGE Consortium	AA880277
2034 E	11 0210AMOV			
2035 G	1 0210AMOV	ESTs	1225317 IMAGE Consortium	AA764060

2036 G	3 0210AMOV	ESTs	1224990 IMAGE Consortium	AA709641
2037 G	5 0210AMOV	regulator of G-protein signaling 11	1195767 IMAGE Consortium	AA711999
2038 G	7 0210AMOV	synaptophysin	1195924 IMAGE Consortium	AA716807
2039 G	9 0210AMOV	ESTs, Moderately similar to ISOLEUCYL-TRNA SYNTHETASE [Staphylococcus aureus]	1230664 IMAGE Consortium	AA879905
2040 G	11 0210AMOV	killer cell lectin-like receptor, subfamily A, member 3	1226206 IMAGE Consortium	AA739464
2041 A	1 0217AMOW	RPB5-mediating protein	1226574 IMAGE Consortium	AA756609
2042 A	3 0217AMOW	ESTs	1243653 IMAGE Consortium	AA789850
2043 A	5 0217AMOW			
2044 A	7 0217AMOW	Mus musculus Stat5a (Stat5a) gene, partial cds; and Stat3 gene, 3' untranslated region	1247511 IMAGE Consortium	AA763337
2045 A	9 0217AMOW	ESTs	1248601 IMAGE Consortium	AA759439
2046 A	11 0217AMOW	ESTs	1246405 IMAGE Consortium	AA823179
2047 C	1 0217AMOW	ESTs, Weakly similar to put. gag and pol gene product [M.musculus]	1244088 IMAGE Consortium	AA790515
2048 C	3 0217AMOW	ESTs, Moderately similar to phosphoserine aminotransferase [H.sapiens]	1226947 IMAGE Consortium	AA738820
2049 C	5 0217AMOW	formin-like	1263246 IMAGE Consortium	AI509558
2050 C	7 0217AMOW	ESTs	1247516 IMAGE Consortium	AA763328
2051 C	9 0217AMOW	ESTs	1247907 IMAGE Consortium	AA759777
2052 C	11 0217AMOW	Mus musculus brain cDNA, clone MNCb-1073	1246431 IMAGE Consortium	AA823200
2053 E	1 0217AMOW	GTP binding protein (gene overexpressed in skeletal muscle)	1226653 IMAGE Consortium	AA756727
2054 E	3 0217AMOW	ESTs	1226968 IMAGE Consortium	AA738872
2055 E	5 0217AMOW	ESTs, Highly similar to unnamed protein product [H.sapiens]	1263364 IMAGE Consortium	AI509487
2056 E	7 0217AMOW	ESTs, Moderately similar to I49130 reverse transcriptase - mouse [M.musculus]	1248283 IMAGE Consortium	AA789551
2057 E	9 0217AMOW	ESTs	1247968 IMAGE Consortium	AA759948
2058 E	11 0217AMOW	ESTs	1247170 IMAGE Consortium	AI465833
2059 G	1 0217AMOW	ESTs	1226703 IMAGE Consortium	AI425682
2060 G	3 0217AMOW	ESTs	1243878 IMAGE Consortium	AI465872
2061 G	5 0217AMOW	SHP2 interacting transmembrane adaptor	1263397 IMAGE Consortium	AI509504
2062 G	7 0217AMOW	ESTs, Weakly similar to muscle glycogen synthase [M.musculus]	1262875 IMAGE Consortium	AI465768
2063 G	9 0217AMOW	ESTs	1246251 IMAGE Consortium	AA823055
2064 G	11 0217AMOW	vasoactive intestinal peptide receptor 1	1245764 IMAGE Consortium	AA798841
2065 A	1 021EAMOX	solute carrier family 2 (facilitated glucose transporter), member 5	1245874 IMAGE Consortium	AA822271
2066 A	3 021EAMOX	ESTs, Moderately similar to KIAA0776 protein [H.sapiens]	1244466 IMAGE Consortium	AA790927
2067 A	5 021EAMOX	recombination activating gene 1	1263354 IMAGE Consortium	AA874497
2068 A	7 021EAMOX	lymphocyte antigen 75	1264311 IMAGE Consortium	AA867690
2069 A	9 021EAMOX	CD68 antigen	1263770 IMAGE Consortium	AA863715

2070 A	11 021EAMOX	ESTs, Moderately similar to GCF_HUMAN GC-RICH SEQUENCE DNA-BINDING FACTOR [H.sapiens]	1264671 IMAGE Consortium	AA881394
2071 C	1 021EAMOX	ESTs	1247334 IMAGE Consortium	AI464959
2072 C	3 021EAMOX	ESTs, Highly similar to Y24K_MOUSE HYPOTHETICAL 24.1 KD PROTEIN ON CHROMOSOME 1 □ [M.musculus]	1245262 IMAGE Consortium	AA821567
2073 C	5 021EAMOX	ESTs	1263380 IMAGE Consortium	AA874520
2074 C	7 021EAMOX	ESTs	1264331 IMAGE Consortium	AI509599
2075 C	9 021EAMOX	ESTs	1265204 IMAGE Consortium	AA867882
2076 C	11 021EAMOX	ESTs	1264674 IMAGE Consortium	AA881385
2077 E	1 021EAMOX	ESTs	1245960 IMAGE Consortium	AA822370
2078 E	3 021EAMOX	D site albumin promoter binding protein	1246022 IMAGE Consortium	AA822420
2079 E	5 021EAMOX	GTF2I repeat domain-containing 1	1263425 IMAGE Consortium	AA874222
2080 E	7 021EAMOX	ESTs	1264347 IMAGE Consortium	AI509605
2081 E	9 021EAMOX	adenylate kinase 3 alpha like	1264526 IMAGE Consortium	AA863874
2082 E	11 021EAMOX	ubiquitin hydrolyzing enzyme 1	1265412 IMAGE Consortium	AA867196
2083 G	1 021EAMOX	ESTs, Weakly similar to unknown [R.norvegicus]	1247426 IMAGE Consortium	AA839813
2084 G	3 021EAMOX	thromboxane A synthase 1, platelet	1245329 IMAGE Consortium	AA821616
2085 G	5 021EAMOX	ESTs	1264201 IMAGE Consortium	AA867621
2086 G	7 021EAMOX	ESTs	1265093 IMAGE Consortium	AA881666
2087 G	9 021EAMOX	ESTs, Moderately similar to ERYTHROID KRUEPPEL-LIKE TRANSCRIPTION FACTOR [Mus musculus]	1265319 IMAGE Consortium	AA867126
2088 G	11 021EAMOX	ESTs, Weakly similar to rjs [M.musculus]	1264768 IMAGE Consortium	AI509651
2089 A	1 021LAMOY	serum amyloid A 3	1264951 IMAGE Consortium	AA881525
2090 A	3 021LAMOY	glycosylation dependent cell adhesion molecule 1	1281416 IMAGE Consortium	AA874245
2091 A	5 021LAMOY	ESTs, Weakly similar to ORF YKR087c [S.cerevisiae]	1379283 IMAGE Consortium	AI048700
2092 A	7 021LAMOY	ESTs, Moderately similar to mutY homolog □ [H.sapiens]	1380122 IMAGE Consortium	AI049068
2093 A	9 021LAMOY	ESTs	1364866 IMAGE Consortium	AI019625
2094 A	11 021LAMOY	ESTs	1345449 IMAGE Consortium	AI158708
2095 C	1 021LAMOY	ESTs	1265839 IMAGE Consortium	AA881885
2096 C	3 021LAMOY	ESTs	1328960 IMAGE Consortium	AA929289
2097 C	5 021LAMOY	ESTs, Highly similar to AMP1_HUMAN PUTATIVE METHIONINE AMINOPEPTIDASE 1 [H.sapiens]	1378722 IMAGE Consortium	AI020548
2098 C	7 021LAMOY	ESTs, Weakly similar to OZF_RAT ZINC FINGER PROTEIN OZF [R.norvegicus]	1365026 IMAGE Consortium	AI019803
2099 C	9 021LAMOY	downstream of tyrosine kinase 1	1380966 IMAGE Consortium	AI035509
2100 C	11 021LAMOY	ESTs	1361499 IMAGE Consortium	AA990041
2101 E	1 021LAMOY	ESTs, Weakly similar to leucine-rich glioma-inactivated protein precursor [H.sapiens]	1265907 IMAGE Consortium	AI509840
2102 E	3 021LAMOY	ESTs	1281519 IMAGE Consortium	AA874421
2103 E	5 021LAMOY	granzyme B	1378751 IMAGE Consortium	AI020130
2104 E	7 021LAMOY	ESTs, Weakly similar to collagen alpha 1 type X [R.norvegicus]	1365088 IMAGE Consortium	AI019893
2105 E	9 021LAMOY	demethyl-Q 7	1380996 IMAGE Consortium	AI035551

2106 E	11 021LAMOY	T-cell receptor alpha, variable 8	1344798 IMAGE Consortium	AA939602
2107 G	1 021LAMOY	homeo box A2	1329029 IMAGE Consortium	AA929366
2108 G	3 021LAMOY	ESTs, Weakly similar to NADH-UBIQUINONE OXIDOREDUCTASE 20 KD SUBUNIT PRECURSOR [Bos taurus]	1282320 IMAGE Consortium	AA864083
2109 G	5 021LAMOY	ESTs, Moderately similar to KIAA1292 protein [H.sapiens]	1378811 IMAGE Consortium	AI020447
2110 G	7 021LAMOY	solute carrier family 27 (fatty acid transporter), member 3	1381127 IMAGE Consortium	AI037339
2111 G	9 021LAMOY	ESTs	1366423 IMAGE Consortium	AI021485
2112 G	11 021LAMOY	Mus musculus glucocorticoid-induced gene 1 mRNA, complete cds	1344841 IMAGE Consortium	AA939689
2113 A	1 021SAMOZ	ESTs, Weakly similar to lost on transformation 1 [R.norvegicus]	1345830 IMAGE Consortium	AA967394
2114 A	3 021SAMOZ	inositol 1,4,5-triphosphate receptor 5	1349854 IMAGE Consortium	AA980905
2115 A	5 021SAMOZ	ESTs	1382266 IMAGE Consortium	AA798385
2116 A	7 021SAMOZ	ESTs, Weakly similar to CR16 [R.norvegicus]	1382549 IMAGE Consortium	AI036165
2117 A	9 021SAMOZ	ESTs, Weakly similar to KF1A_MOUSE KINESIN-LIKE PROTEIN KIF1A [M.musculus]	1382883 IMAGE Consortium	AI120267
2118 A	11 021SAMOZ	ESTs, Weakly similar to JC4871 phospholipase C [M.musculus]	403623 IMAGE Consortium	AI425845
2119 C	1 021SAMOZ	ESTs, Weakly similar to unnamed protein product [H.sapiens]	1345873 IMAGE Consortium	AA982082
2120 C	3 021SAMOZ	ESTs, Weakly similar to CR16 [R.norvegicus]	1349926 IMAGE Consortium	AA980959
2121 C	5 021SAMOZ	ESTs	1382313 IMAGE Consortium	AI037524
2122 C	7 021SAMOZ	ESTs	1382589 IMAGE Consortium	AI036374
2123 C	9 021SAMOZ	Rhesus blood group-associated A glycoprotein	403128 IMAGE Consortium	W81811
2124 C	11 021SAMOZ	ESTs	403654 IMAGE Consortium	W82097
2125 E	1 021SAMOZ	ESTs	1362636 IMAGE Consortium	AI006281
2126 E	3 021SAMOZ	UDP-N-acetylglucosamine-2-epimerase/N-acetylmannosamine kinase	1382016 IMAGE Consortium	AA983004
2127 E	5 021SAMOZ	nuclear DNA-binding protein	1382325 IMAGE Consortium	AI037530
2128 E	7 021SAMOZ	ESTs, Weakly similar to INVO_RAT INVOLUCRIN [R.norvegicus]	1382608 IMAGE Consortium	AI036383
2129 E	9 021SAMOZ	Indian hedgehog homolog, (Drosophila)	403861 IMAGE Consortium	W82225
2130 E	11 021SAMOZ	ESTs	404395 IMAGE Consortium	AI425316
2131 G	1 021SAMOZ	ESTs	1345962 IMAGE Consortium	AA967486
2132 G	3 021SAMOZ	ESTs, Highly similar to AF170284_1 connexin 30 [R.norvegicus]	1382084 IMAGE Consortium	AA982871
2133 G	5 021SAMOZ	ESTs	1383127 IMAGE Consortium	AI120784
2134 G	7 021SAMOZ	ESTs	1382634 IMAGE Consortium	AI036406
2135 G	9 021SAMOZ	ESTs	407708 IMAGE Consortium	W85664
2136 G	11 021SAMOZ	ESTs	404440 IMAGE Consortium	AI425318
2137 A	1 021SAMPO	inter-alpha trypsin inhibitor, heavy chain 2	407079 IMAGE Consortium	W83984
2138 A	3 021SAMPO	stathmin-like-protein RB3	405910 IMAGE Consortium	W83476



2139 A	5 021SAMP0	ESTs	408670 IMAGE Consortium	AI430295
2140 A	7 021SAMP0	ESTs	1396353 IMAGE Consortium	AI286798
2141 A	9 021SAMP0	SH3 domain protein 4	1383423 IMAGE Consortium	AI182689
2142 A	11 021SAMP0	Kirsten rat sarcoma oncogene 2, expressed	1383721 IMAGE Consortium	AI180903
2143 C	1 021SAMP0	ESTs	407119 IMAGE Consortium	W84305
2144 C	3 021SAMP0	hyaluronidase 2	405956 IMAGE Consortium	W83533
2145 C	5 021SAMP0	family with sequence similarity 4, subfamily A, member 2	407987 IMAGE Consortium	W84258
2146 C	7 021SAMP0	transmembrane protein with EGF-like and two follistatin-like domains 2	1397175 IMAGE Consortium	AI098476
2147 C	9 021SAMP0	matrix metalloproteinase 11	1383434 IMAGE Consortium	AI120903
2148 C	11 021SAMP0	guanylate nucleotide binding protein 2	1383749 IMAGE Consortium	AI180927
2149 E	1 021SAMP0	ESTs		
2150 E	3 021SAMP0	ESTs	407410 IMAGE Consortium	W89971
2151 E	5 021SAMP0	ESTs	1383835 IMAGE Consortium	AI181696
2152 E	7 021SAMP0	ESTs, Weakly similar to ACTB_HUMAN ACTIN, CYTOPLASMIC 1 [M.musculus]	1396579 IMAGE Consortium	AI882327
2153 E	9 021SAMP0	ESTs	1383538 IMAGE Consortium	AI120945
2154 E	11 021SAMP0	ESTs, Moderately similar to unnamed protein product [H.sapiens]	1383772 IMAGE Consortium	AI180939
2155 G	1 021SAMP0	potassium voltage-gated channel, subfamily H (eag-related), member 2	407176 IMAGE Consortium	W84157
2156 G	3 021SAMP0	dante	407471 IMAGE Consortium	W89290
2157 G	5 021SAMP0	cAMP responsive element modulator	1383847 IMAGE Consortium	AI181714
2158 G	7 021SAMP0	histocompatibility 2, T region locus 23	1396665 IMAGE Consortium	AI098976
2159 G	9 021SAMP0	ESTs	1383553 IMAGE Consortium	AI120971
2160 G	11 021SAMP0	ESTs	1383790 IMAGE Consortium	AI180948
2161 A	1 021AMSI	ESTs	1433746 IMAGE Consortium	AA839535
2162 A	3 021AMSI	ESTs	1446196 IMAGE Consortium	AI157253
2163 A	5 021AMSI	three prime repair exonuclease 1	1430424 IMAGE Consortium	AI182180
2164 A	7 021AMSI	ESTs	1400701 IMAGE Consortium	AI121805
2165 A	9 021AMSI	colony stimulating factor 2 (granulocyte-macrophage)	1428783 IMAGE Consortium	AI180669
2166 A	11 021AMSI	ESTs		
2167 C	1 021AMSI	ESTs	1433871 IMAGE Consortium	AA840125
2168 C	3 021AMSI	ESTs	1446244 IMAGE Consortium	AI151994
2169 C	5 021AMSI	aquaporin 5	1430484 IMAGE Consortium	AI182792
2170 C	7 021AMSI	X-linked myotubular myopathy gene 1	1400777 IMAGE Consortium	AI121593
2171 C	9 021AMSI	ESTs	1399564 IMAGE Consortium	AI121411
2172 C	11 021AMSI	tumor necrosis factor (ligand) superfamily, member 9	1449661 IMAGE Consortium	AI157872
2173 E	1 021AMSI	ESTs	1434621 IMAGE Consortium	AI877342
2174 E	3 021AMSI	ESTs	1445927 IMAGE Consortium	AI122193
2175 E	5 021AMSI	ESTs	1400079 IMAGE Consortium	AI132005
2176 E	7 021AMSI	Gag...env {provirus} [Mus musculus, MrV, Evi-2, murine AIDS virus-related provirus, Genomic Mutant, 3 genes,	1400848 IMAGE Consortium	AI121690

		4765 nt]		
2177 E	9 0215AMSI	H2.0-like homeo box gene	1399735 IMAGE Consortium	AI121405
2178 E	11 0215AMSI			
2179 G	1 0215AMSI	ESTs, Moderately similar to I68673 gene X123 protein - human [H.sapiens]	1433926 IMAGE Consortium	AA840163
2180 G	3 0215AMSI	growth differentiation factor 8	1445968 IMAGE Consortium	AI122124
2181 G	5 0215AMSI	ESTs	1400360 IMAGE Consortium	AI117739
2182 G	7 0215AMSI	heterogeneous nuclear ribonucleoprotein A1	1429125 IMAGE Consortium	AI183202
2183 G	9 0215AMSI	erythrocyte protein band 4.1-like 1	1399824 IMAGE Consortium	AI131926
2184 G	11 0215AMSI	ESTs	1449791 IMAGE Consortium	AI153146
2185 A	1 0216AMP2	ESTs, Weakly similar to similar to hydrolase [C.elegans]	1450220 IMAGE Consortium	AI155129
2186 A	3 0216AMP2	casein gamma	1448004 IMAGE Consortium	AI159406
2187 A	5 0216AMP2	transcript expressed during hematopoiesis 2	1448313 IMAGE Consortium	AI173087
2188 A	7 0216AMP2	ESTs, Weakly similar to alpha glucosidase II, alpha subunit [M.musculus]	1478043 IMAGE Consortium	AI152618
2189 A	9 0216AMP2	DNA segment, Chr 7, ERATO Doi 304, expressed	1495107 IMAGE Consortium	AI157475
2190 A	11 0216AMP2	epidermal growth factor receptor	1493993 IMAGE Consortium	AI156625
2191 C	1 0216AMP2	ESTs	1446280 IMAGE Consortium	AI152021
2192 C	3 0216AMP2	ESTs	1448077 IMAGE Consortium	AI154671
2193 C	5 0216AMP2			
2194 C	7 0216AMP2	ESTs	1478120 IMAGE Consortium	AI152709
2195 C	9 0216AMP2	ESTs	1478434 IMAGE Consortium	AI152670
2196 C	11 0216AMP2	ESTs	1478745 IMAGE Consortium	AI157703
2197 E	1 0216AMP2	nuclear transcription factor-Y beta	1447053 IMAGE Consortium	AI158083
2198 E	3 0216AMP2	lymphoid-restricted membrane protein	1446702 IMAGE Consortium	AI158088
2199 E	5 0216AMP2	ESTs	1446958 IMAGE Consortium	AI158547
2200 E	7 0216AMP2	ESTs	1478220 IMAGE Consortium	AI152800
2201 E	9 0216AMP2	cytochrome P450, 7b1	1493787 IMAGE Consortium	AI156457
2202 E	11 0216AMP2	osteomodulin	1494961 IMAGE Consortium	AI157520
2203 G	1 0216AMP2	T-box 6	1446422 IMAGE Consortium	AI152111
2204 G	3 0216AMP2	ESTs	1447447 IMAGE Consortium	AI154499
2205 G	5 0216AMP2			
2206 G	7 0216AMP2	adenylate cyclase 8	1478303 IMAGE Consortium	AI152525
2207 G	9 0216AMP2	mitogen-activated protein kinase kinase kinase 6	1477831 IMAGE Consortium	AI155236
2208 G	11 0216AMP2	ESTs	1479844 IMAGE Consortium	AI156310
2209 A	1 021DAMP3	epidermal growth factor-containing fibulin-like extracellular matrix protein 1	1480170 IMAGE Consortium	AI156278
2210 A	3 021DAMP3	ESTs, Weakly similar to AF153906_1 erythroid membrane-associated protein ERMAP [M.musculus]	1746076 IMAGE Consortium	AI036797
2211 A	5 021DAMP3	striatin	1746470 IMAGE Consortium	AI037693
2212 A	7 021DAMP3			
2213 A	9 021DAMP3	fos-like antigen 1	1498182 IMAGE Consortium	AI225491
2214 A	11 021DAMP3	protein tyrosine phosphatase, receptor type, B	1497631 IMAGE Consortium	AI664182

2215 C	1 021DAMP3	golgi SNAP receptor complex member 2	1749586 IMAGE Consortium	AI047095
2216 C	3 021DAMP3	ESTs	1746109 IMAGE Consortium	AI036909
2217 C	5 021DAMP3	ESTs, Moderately similar to testatin [M.musculus]	1746548 IMAGE Consortium	AI037715
2218 C	7 021DAMP3	ESTs	1749184 IMAGE Consortium	AI047912
2219 C	9 021DAMP3	ESTs, Highly similar to hypothetical protein [H.sapiens]	1617172 IMAGE Consortium	AA967999
2220 C	11 021DAMP3	SH3-domain binding protein 2	1511041 IMAGE Consortium	AI786129
2221 E	1 021DAMP3			
2222 E	3 021DAMP3	ESTs	1746218 IMAGE Consortium	AI036935
2223 E	5 021DAMP3	ESTs	1745871 IMAGE Consortium	AI789636
2224 E	7 021DAMP3	ESTs, Highly similar to ICE3_MOUSE APOPAIN PRECURSOR [M.musculus]	1749558 IMAGE Consortium	AI046775
2225 E	9 021DAMP3	kallikrein 4 (prostase, enamel matrix, prostate)	1498211 IMAGE Consortium	AI226226
2226 E	11 021DAMP3	Mus musculus mRNA, complete cds, similar to L-amino acid oxidase	1511158 IMAGE Consortium	AI527496
2227 G	1 021DAMP3	ESTs	1749662 IMAGE Consortium	AI046796
2228 G	3 021DAMP3	ESTs	1745580 IMAGE Consortium	AI173044
2229 G	5 021DAMP3	ESTs, Highly similar to TLBP_MOUSE TESTIS LIPID BINDING PROTEIN [M.musculus]	1746654 IMAGE Consortium	AI046575
2230 G	7 021DAMP3	lysosomal trafficking regulator	1510870 IMAGE Consortium	AI227406
2231 G	9 021DAMP3	gap junction membrane channel protein beta 4	1498230 IMAGE Consortium	AI226235
2232 G	11 021DAMP3	ESTs	1497839 IMAGE Consortium	AI664331
2233 A	1 021KAMP4	whey acidic protein	1511486 IMAGE Consortium	AI786507
2234 A	3 021KAMP4	basonuclin	1764286 IMAGE Consortium	AI047752
2235 A	5 021KAMP4			
2236 A	7 021KAMP4	ESTs	537212 IMAGE Consortium	AA117864
2237 A	9 021KAMP4			
2238 A	11 021KAMP4	protein kinase, DNA activated, catalytic polypeptide	537728 IMAGE Consortium	AI324450
2239 C	1 021KAMP4	ESTs	1851189 IMAGE Consortium	AI197423
2240 C	3 021KAMP4	signal transducing adaptor molecule (SH3 domain and ITAM motif) 1	1764536 IMAGE Consortium	AI115506
2241 C	5 021KAMP4	casein kinase 1, epsilon	537059 IMAGE Consortium	AI550083
2242 C	7 021KAMP4			
2243 C	9 021KAMP4	ESTs, Highly similar to SERINE/THREONINE PROTEIN PHOSPHATASE 2B CATALYTIC SUBUNIT, BETA ISOFORM [Homo sapiens]	537509 IMAGE Consortium	AI892461
2244 C	11 021KAMP4	ESTs, Weakly similar to AF144731_1 putative splicing factor YT521-B [R.norvegicus]	537797 IMAGE Consortium	AA120565
2245 E	1 021KAMP4	hepatocyte growth factor	1852022 IMAGE Consortium	AI225823
2246 E	3 021KAMP4	deleted in azoospermia-like	1852783 IMAGE Consortium	AI256688
2247 E	5 021KAMP4			
2248 E	7 021KAMP4	ESTs	537256 IMAGE Consortium	AI550098
2249 E	9 021KAMP4	ESTs, Moderately similar to 2204390A synaptojanin [R.norvegicus]	537568 IMAGE Consortium	AA118143

2250 E	11 021KAMP4	testis expressed gene 9	538015 IMAGE Consortium	AA119696
2251 G	1 021KAMP4	Nik related kinase	1852105 IMAGE Consortium	AI225895
2252 G	3 021KAMP4	ESTs	1764676 IMAGE Consortium	AI115571
2253 G	5 021KAMP4	epidermal growth factor receptor pathway substrate 15	537100 IMAGE Consortium	AA117832
2254 G	7 021KAMP4	growth arrest specific 6	537284 IMAGE Consortium	AI324446
2255 G	9 021KAMP4	growth hormone releasing hormone	536865 IMAGE Consortium	AA114706
2256 G	11 021KAMP4	M.musculus mRNA for e1 protein	538077 IMAGE Consortium	AA114799
2257 A	1 021RAMP5	DNA segment, Chr 18, Wayne State University 70, expressed	538208 IMAGE Consortium	AA119562
2258 A	3 021RAMP5	DNA segment, Chr 8, ERATO Doi 118, expressed	538454 IMAGE Consortium	AI550172
2259 A	5 021RAMP5	ESTs	538637 IMAGE Consortium	AI550186
2260 A	7 021RAMP5	phosphodiesterase 10A	760844 IMAGE Consortium	AI508642
2261 A	9 021RAMP5			
2262 A	11 021RAMP5	sarcospan	406138 IMAGE Consortium	W83284
2263 C	1 021RAMP5	ESTs, Moderately similar to unnamed protein product [H.sapiens]	538218 IMAGE Consortium	AI550156
2264 C	3 021RAMP5	teratocarcinoma-derived growth factor	538474 IMAGE Consortium	AA116676
2265 C	5 021RAMP5	praja1	537943 IMAGE Consortium	AA119752
2266 C	7 021RAMP5	ajuba	2101967 IMAGE Consortium	AW210250
2267 C	9 021RAMP5	huntingtin-associated protein 1	403259 IMAGE Consortium	W81952
2268 C	11 021RAMP5	apolipoprotein CIV	404046 IMAGE Consortium	W82417
2269 E	1 021RAMP5	LIM only 4	538287 IMAGE Consortium	AI325787
2270 E	3 021RAMP5	synaptobrevin like 1	537811 IMAGE Consortium	AI324452
2271 E	5 021RAMP5	ESTs, Highly similar to EUKARYOTIC INITIATION FACTOR 1A [Homo sapiens]	538685 IMAGE Consortium	AI550189
2272 E	7 021RAMP5	Spi-C transcription factor (Spi-1/PU.1 related)	988963 IMAGE Consortium	AA591091
2273 E	9 021RAMP5	ribosomal protein L7	403969 IMAGE Consortium	W82240
2274 E	11 021RAMP5	M.musculus mRNA for ventricular alkali myosin light chain	404103 IMAGE Consortium	W82497
2275 G	1 021RAMP5	double cortin and calcium/calmodulin-dependent protein kinase-like 1	538298 IMAGE Consortium	AA116787
2276 G	3 021RAMP5	ESTs, Highly similar to TRANSLATIONAL INITIATION FACTOR 2 ALPHA SUBUNIT [Rattus norvegicus; Bos taurus]	538553 IMAGE Consortium	AI550176
2277 G	5 021RAMP5	aryl hydrocarbon receptor nuclear translocator-like	537975 IMAGE Consortium	AI550145
2278 G	7 021RAMP5	histidine decarboxylase cluster	2102077 IMAGE Consortium	AW210352
2279 G	9 021RAMP5	protein that interacts with C kinase 1	406016 IMAGE Consortium	AI893973
2280 G	11 021RAMP5	Max dimerization protein 4	402006 IMAGE Consortium	AI425758
2281 A	1 021YAMP6	Mus musculus S6 kinase 2 {IMAGE:421114}	421114 IMAGE Consortium	W91436
2282 A	3 021YAMP6	tubulin alpha 2	407117 IMAGE Consortium	W84304
2283 A	5 021YAMP6	IMAGE EST {IMAGE:438376}	438376 IMAGE Consortium	AA008228
2284 A	7 021YAMP6	small nuclear ribonucleoprotein E	426671 IMAGE Consortium	AA002725
2285 A	9 021YAMP6	Mouse mRNA for mMIS5, complete cds {IMAGE:444918}	444918 IMAGE Consortium	AA016759
2286 A	11 021YAMP6	Nef-associated factor 1	444725 IMAGE Consortium	AA016332

2287 C	1 021YAMP6	myelin basic protein expression factor 2, repressor	407001 IMAGE Consortium	W83960
2288 C	3 021YAMP6	3 monooxygenase/tryptophan 5 monooxygenase activation protein, beta polypeptide	407180 IMAGE Consortium	AA052594
2289 C	5 021YAMP6	IMAGE EST {IMAGE:438506}	438506 IMAGE Consortium	AA050778
2290 C	7 021YAMP6	perlecan (heparan sulfate proteoglycan 2)	438329 IMAGE Consortium	AA008180
2291 C	9 021YAMP6	glutamate-cysteine ligase , modifier subunit	440961 IMAGE Consortium	AA008956
2292 C	11 021YAMP6	proteasome (prosome, macropain) subunit, alpha type 4	441164 IMAGE Consortium	AI322459
2293 E	1 021YAMP6	zinc finger protein 144	406658 IMAGE Consortium	W83643
2294 E	3 021YAMP6	ESTs	407985 IMAGE Consortium	AA052693
2295 E	5 021YAMP6			
2296 E	7 021YAMP6	cyclin D3	427529 IMAGE Consortium	AA002467
2297 E	9 021YAMP6	gap junction membrane channel protein beta 5	445670 IMAGE Consortium	AI894067
2298 E	11 021YAMP6	vesicle-associated membrane protein 2	439726 IMAGE Consortium	AA014530
2299 G	1 021YAMP6	ATPase inhibitor	406863 IMAGE Consortium	W83805
2300 G	3 021YAMP6	IMAGE EST {IMAGE:423177}	423177 IMAGE Consortium	W97303
2301 G	5 021YAMP6	DNA segment, Chr 11, ERATO Doi 416, expressed	437969 IMAGE Consortium	AA007789
2302 G	7 021YAMP6	procollagen, type XIV, alpha 1	426173 IMAGE Consortium	AA002997
2303 G	9 021YAMP6	proprotein convertase subtilisin/kexin type 5	439378 IMAGE Consortium	AI894035
2304 G	11 021YAMP6	nel-like 2 homolog (chicken)	439202 IMAGE Consortium	AA003877
2305 A	1 0215AMP7	Mus musculus mRNA for vascular cadherin-2 {IMAGE:459569}	459569 IMAGE Consortium	AA027396
2306 A	3 0215AMP7	keratin complex 2, basic, gene 4	464229 IMAGE Consortium	AA028578
2307 A	5 0215AMP7	ribosomal protein, mitochondrial, L26	477770 IMAGE Consortium	AA048585
2308 A	7 0215AMP7	inter alpha-trypsin inhibitor, heavy chain 4	479543 IMAGE Consortium	AA049060
2309 A	9 0215AMP7	Mouse housekeeping DXS253E (P3) and DXS254E (GdX) genes, 3' end, and complete cds {IMAGE:483262}	483262 IMAGE Consortium	AA061673
2310 A	11 0215AMP7	MAD homolog 2 (Drosophila)	483280 IMAGE Consortium	AA061479
2311 C	1 0215AMP7			
2312 C	3 0215AMP7	WNT1 inducible signaling pathway protein 1	465093 IMAGE Consortium	AA032662
2313 C	5 0215AMP7	IMAGE EST {IMAGE:478710}	478710 IMAGE Consortium	AA051380
2314 C	7 0215AMP7	SH3 domain protein 2B	479638 IMAGE Consortium	AA051447
2315 C	9 0215AMP7	ESTs {IMAGE:483636}	483636 IMAGE Consortium	AA061332
2316 C	11 0215AMP7	stromal cell derived factor 5	482206 IMAGE Consortium	AA061047
2317 E	1 0215AMP7	ESTs, Highly similar to CART PROTEIN PRECURSOR [Rattus norvegicus] {IMAGE:466874}	466874 IMAGE Consortium	AA031207
2318 E	3 0215AMP7	septin 5	445831 IMAGE Consortium	AA020101
2319 E	5 0215AMP7	Solute carrier family 4 (anion exchanger), member 2 {IMAGE:479476}	479476 IMAGE Consortium	AA048952
2320 E	7 0215AMP7	ribosomal protein L18	479685 IMAGE Consortium	AA051462
2321 E	9 0215AMP7	Crystallin, gamma S {IMAGE:483498}	483498 IMAGE Consortium	AA060041
2322 E	11 0215AMP7	nuclear RNA helicase Bat1	483714 IMAGE Consortium	AA061218
2323 G	1 0215AMP7	superoxide dismutase 3, extracellular	459497 IMAGE Consortium	AA027504
2324 G	3 0215AMP7			

2325 G	5 0215AMP7	Bcl2-like {IMAGE:478723}	478723 IMAGE Consortium	AA051441
2326 G	7 0215AMP7	NADH dehydrogenase flavoprotein 1	479775 IMAGE Consortium	AA051567
2327 G	9 0215AMP7	Mouse mRNA for dbpA murine homologue, complete cds {IMAGE:481949}	481949 IMAGE Consortium	AA059953
2328 G	11 0215AMP7	nuclear receptor co-repressor 2	481825 IMAGE Consortium	AA059713
2329 A	1 021CAMP8	IMAGE EST {IMAGE:574658}	574658 IMAGE Consortium	AA125255
2330 A	3 021CAMP8	Tissue inhibitor of metalloproteinase 3 {IMAGE:580753}	580753 IMAGE Consortium	AA145784
2331 A	5 021CAMP8	nitrogen fixation gene 1 (S. cerevisiae)	313534 IMAGE Consortium	AI322977
2332 A	7 021CAMP8	serine/threonine kinase 6	635521 IMAGE Consortium	AI893749
2333 A	9 021CAMP8	hexokinase 1	635778 IMAGE Consortium	AA163537
2334 A	11 021CAMP8	matrix metalloproteinase 23	653582 IMAGE Consortium	AA217682
2335 C	1 021CAMP8	Transporter 1, ABC (ATP binding cassette) {IMAGE:576775}	576775 IMAGE Consortium	AA123837
2336 C	3 021CAMP8	focal adhesion kinase	577544 IMAGE Consortium	AA139443
2337 C	5 021CAMP8	mannosidase 1, alpha	598192 IMAGE Consortium	AA163718
2338 C	7 021CAMP8	DNA segment, Chr 18, Wayne State University 70, expressed	619509 IMAGE Consortium	AA172608
2339 C	9 021CAMP8	complement component 1, q subcomponent binding protein	642511 IMAGE Consortium	AA185481
2340 C	11 021CAMP8	leukocyte cell-derived chemotaxin 2	658696 IMAGE Consortium	AA221986
2341 E	1 021CAMP8	H2-K region expressed gene 6	570567 IMAGE Consortium	AA108481
2342 E	3 021CAMP8	integrin alpha 4 (Cd49d)	581712 IMAGE Consortium	AA152636
2343 E	5 021CAMP8	cathepsin S	596534 IMAGE Consortium	AA146437
2344 E	7 021CAMP8	nuclear receptor subfamily 2, group F, member 2	618835 IMAGE Consortium	AA170421
2345 E	9 021CAMP8	Guanidinoacetate methyltransferase {IMAGE:671133}	671133 IMAGE Consortium	AA222404
2346 E	11 021CAMP8	regulator of G-protein signaling 2	658741 IMAGE Consortium	AA221794
2347 G	1 021CAMP8	inositol polyphosphate-5-phosphatase, 145 kDa	574851 IMAGE Consortium	AA125192
2348 G	3 021CAMP8			
2349 G	5 021CAMP8	IL2-inducible T-cell kinase	596235 IMAGE Consortium	AA137960
2350 G	7 021CAMP8	glycine N-methyltransferase	634934 IMAGE Consortium	AI893740
2351 G	9 021CAMP8	ESTs, Highly similar to VACUOLAR ATP SYNTHASE SUBUNIT AC45 PRECURSOR [Bos taurus] {IMAGE:671660}	671660 IMAGE Consortium	AA241780
2352 G	11 021CAMP8	Wnt inhibitory factor 1	658891 IMAGE Consortium	AA217902
2353 A	1 021JAMP9	ESTs {IMAGE:678748}	678748 IMAGE Consortium	AA239479
2354 A	3 021JAMP9	ESTs, Weakly similar to apolipoprotein F [H.sapiens]	679114 IMAGE Consortium	AI390319
2355 A	5 021JAMP9	Mus musculus MEK binding partner 1 (Mp1) mRNA, complete cds {IMAGE:718194}	718194 IMAGE Consortium	AA266352
2356 A	7 021JAMP9	Mus musculus putative G protein-coupled receptor TDAG8 (TDAG8) mRNA, complete cds {IMAGE:723208}	723208 IMAGE Consortium	AA254500
2357 A	9 021JAMP9	cell division cycle 71 homolog (S. cerevisiae)-like 1	734744 IMAGE Consortium	AA261750
2358 A	11 021JAMP9	Zinc finger protein 30 {IMAGE:747957}	747957 IMAGE Consortium	AA260956
2359 C	1 021JAMP9	ESTs, Moderately similar to RETINOL	681236 IMAGE Consortium	AA244820

		DEHYDROGENASE TYPE II [R.norvegicus] {IMAGE:681236}		
2360 C	3 021JAMP9	proteasome (prosome, macropain) 26S subunit, non-ATPase, 4	681449 IMAGE Consortium	AA237807
2361 C	5 021JAMP9	dihydrolipoamide dehydrogenase	681768 IMAGE Consortium	AA245976
2362 C	7 021JAMP9	ESTs, Highly similar to PROTEIN PHOSPHATASE PP2A, 65 KD REGULATORY SUBUNIT, ALPHA ISOFORM [Homo sapiens] {IMAGE:733882}	733882 IMAGE Consortium	AA268171
2363 C	9 021JAMP9	parvalbumin	720348 IMAGE Consortium	AA270885
2364 C	11 021JAMP9	Mus musculus methyl-CpG binding protein MBD1 (Mbd1) mRNA, complete cds {IMAGE:750171}	750171 IMAGE Consortium	AA396451
2365 E	1 021JAMP9	Proprotein convertase subtilisin/kexin type 3 {IMAGE:680455}	680455 IMAGE Consortium	AA237179
2366 E	3 021JAMP9	mannose binding lectin, liver (A)	680110 IMAGE Consortium	AA244667
2367 E	5 021JAMP9	serine protease inhibitor, Kazal type 3	697396 IMAGE Consortium	AA242201
2368 E	7 021JAMP9	Mus musculus maf10 (mAF10) mRNA, complete cds {IMAGE:736069}	736069 IMAGE Consortium	AA270281
2369 E	9 021JAMP9	protein kinase, cAMP dependent, catalytic, alpha	720387 IMAGE Consortium	AA261433
2370 E	11 021JAMP9	ornithine transcarbamylase	748359 IMAGE Consortium	AA268022
2371 G	1 021JAMP9	Phosphatidylethanolamine N-methyltransferase {IMAGE:680973}	680973 IMAGE Consortium	AA242031
2372 G	3 021JAMP9	hydroxysteroid dehydrogenase-3, delta<5>-3-beta	681708 IMAGE Consortium	AA245848
2373 G	5 021JAMP9	caspase 1	718334 IMAGE Consortium	AA261094
2374 G	7 021JAMP9	karyopherin (importin) alpha 1	720974 IMAGE Consortium	AA266741
2375 G	9 021JAMP9	ribosomal protein S6	720445 IMAGE Consortium	AA261524
2376 G	11 021JAMP9	small inducible cytokine A3	751833 IMAGE Consortium	AA396229
2377 A	1 0213AMPA	non-catalytic region of tyrosine kinase adaptor protein 1	751332 IMAGE Consortium	AA289435
2378 A	3 0213AMPA	Mus musculus cell death activator CIDE-B (Cide-b) mRNA, complete cds {IMAGE:790532}	790532 IMAGE Consortium	AA290390
2379 A	5 0213AMPA	histocompatibility 2, class II antigen A, beta 1	765285 IMAGE Consortium	AA273267
2380 A	7 0213AMPA	homer, neuronal immediate early gene, 2	818542 IMAGE Consortium	AA434982
2381 A	9 0213AMPA	etoposide induced 2.4 mRNA	316478 IMAGE Consortium	W11184
2382 A	11 0213AMPA	phosphate regulating neutral endopeptidases on the X chromosome	875117 IMAGE Consortium	AA472625
2383 C	1 0213AMPA	cytochrome c oxidase, subunit XVII assembly protein homolog (yeast)	748463 IMAGE Consortium	AA272869
2384 C	3 0213AMPA	ESTs, Weakly similar to Zic2 protein [M.musculus] {IMAGE:791455}	791455 IMAGE Consortium	AA416308
2385 C	5 0213AMPA	maternal inhibition of differentiation	807700 IMAGE Consortium	AA432678
2386 C	7 0213AMPA	CD36 antigen	846947 IMAGE Consortium	AI553078
2387 C	9 0213AMPA	small nuclear ribonucleoprotein N	317222 IMAGE Consortium	W13699
2388 C	11 0213AMPA	succinate-Coenzyme A ligase, ADP-forming, beta subunit	870855 IMAGE Consortium	AA462048
2389 E	1 0213AMPA	esterase 31	747188 IMAGE Consortium	AA260296
2390 E	3 0213AMPA	Selenophosphate synthetase 2 {IMAGE:791719}	791719 IMAGE Consortium	AA414662

2391 E	5 0213AMPA	CD48 antigen	765555 IMAGE Consortium	AA275139
2392 E	7 0213AMPA	glycogenin 1	818620 IMAGE Consortium	AA444326
2393 E	9 0213AMPA	mannosidase 2, alpha B2	316205 IMAGE Consortium	W14484
2394 E	11 0213AMPA			
2395 G	1 0213AMPA			
2396 G	3 0213AMPA	transforming growth factor alpha	777327 IMAGE Consortium	AA276915
2397 G	5 0213AMPA	ESTs, Weakly similar to slit protein 2 precursor [D.melanogaster] {IMAGE:820354}	820354 IMAGE Consortium	AA423126
2398 G	7 0213AMPA	sphingomyelin phosphodiesterase 2, neutral	807844 IMAGE Consortium	AA432637
2399 G	9 0213AMPA	loricrin	316909 IMAGE Consortium	AI536281
2400 G	11 0213AMPA	RAB3D, member RAS oncogene family	864389 IMAGE Consortium	AA458159
2401 A	1 021AAMPB	ESTs	871971 IMAGE Consortium	AI614443
2402 A	3 021AAMPB	ATPase, class II, type 9A	891343 IMAGE Consortium	AA510705
2403 A	5 021AAMPB	CD3 antigen, gamma polypeptide	1054728 IMAGE Consortium	AA796027
2404 A	7 021AAMPB	heat shock protein 40	1121890 IMAGE Consortium	AA657278
2405 A	9 021AAMPB	discs, large homolog 3 (Drosophila)	1225100 IMAGE Consortium	AA756386
2406 A	11 021AAMPB	matrix metalloproteinase 15	1193999 IMAGE Consortium	AA726630
2407 C	1 021AAMPB	ESTs, Highly similar to homogentisate 1,2-dioxygenase [M.musculus]	871999 IMAGE Consortium	AA473465
2408 C	3 021AAMPB	kallikrein 3, plasma	891359 IMAGE Consortium	AA510906
2409 C	5 021AAMPB	H2-K region expressed gene 2	935662 IMAGE Consortium	AA543980
2410 C	7 021AAMPB	DNA segment, Chr 4, ERATO Doi 659, expressed	1134385 IMAGE Consortium	AA675670
2411 C	9 021AAMPB	huntingtin interacting protein 1 related	1225184 IMAGE Consortium	AI428659
2412 C	11 021AAMPB	SH3 domain protein 5	1177991 IMAGE Consortium	AA718587
2413 E	1 021AAMPB	Selectin, platelet (p-selectin) ligand {IMAGE:876033}	876033 IMAGE Consortium	AA475311
2414 E	3 021AAMPB	secreted acidic cysteine rich glycoprotein	903245 IMAGE Consortium	AI553166
2415 E	5 021AAMPB	Jun oncogene	949554 IMAGE Consortium	AA537136
2416 E	7 021AAMPB	cysteine rich protein	1122847 IMAGE Consortium	AA794055
2417 E	9 021AAMPB	elastase 1, pancreatic	1195893 IMAGE Consortium	AA717025
2418 E	11 021AAMPB	cystatin C	1162792 IMAGE Consortium	AA770768
2419 G	1 021AAMPB	ESTs {IMAGE:890726}	890726 IMAGE Consortium	AA510013
2420 G	3 021AAMPB	prostaglandin-endoperoxide synthase 1	903272 IMAGE Consortium	AA518993
2421 G	5 021AAMPB	aldo-keto reductase AKR1C13	1052178 IMAGE Consortium	AA596393
2422 G	7 021AAMPB	prolyl 4-hydroxylase, beta polypeptide	1162896 IMAGE Consortium	AA770866
2423 G	9 021AAMPB	RAD30 (S. cerevisiae) homolog B	1178409 IMAGE Consortium	AA718031
2424 G	11 021AAMPB	IG MU CHAIN C REGION	1179412 IMAGE Consortium	AA754696
2425 A	1 021HAMPC	angiopoietin related protein 2	1180320 IMAGE Consortium	AA755981
2426 A	3 021HAMPC	Ia-associated invariant chain	1247764 IMAGE Consortium	AA759679
2427 A	5 021HAMPC	DEAD/H (Asp-Glu-Ala-Asp/His) box binding protein 1	1264473 IMAGE Consortium	AA881370
2428 A	7 021HAMPC	tetratricopeptide repeat domain	1264754 IMAGE Consortium	AA881399
2429 A	9 021HAMPC	zinc finger protein 185	1332959 IMAGE Consortium	AI159217
2430 A	11 021HAMPC	immediate early response 3	1381903 IMAGE Consortium	AA833402



2431 C	1	021HAMPC	huntingtin interacting protein 2	1224917	IMAGE Consortium	AA756321
2432 C	3	021HAMPC	lysosomal-associated protein transmembrane 5	1226115	IMAGE Consortium	AA739425
2433 C	5	021HAMPC	CD79a antigen	1332396	IMAGE Consortium	AI876995
2434 C	7	021HAMPC	zinc finger protein 38	1332649	IMAGE Consortium	AI159431
2435 C	9	021HAMPC	serine protease inhibitor 7	1382373	IMAGE Consortium	AI037614
2436 C	11	021HAMPC	COP9 (constitutive photomorphogenic, subunit 2 (Arabidopsis))	1382651	IMAGE Consortium	AI036503
2437 E	1	021HAMPC	actin, alpha 1, skeletal muscle	1162935	IMAGE Consortium	AA770902
2438 E	3	021HAMPC	endomucin	1245854	IMAGE Consortium	AA822842
2439 E	5	021HAMPC	dolichol-phosphate (beta-D) mannosyltransferase 2	1264507	IMAGE Consortium	AA863865
2440 E	7	021HAMPC	regulatory factor (trans-acting) 2	1263341	IMAGE Consortium	AA874501
2441 E	9	021HAMPC	jumping translocation breakpoint	1378778	IMAGE Consortium	AI020148
2442 E	11	021HAMPC	Mus musculus (strain C57Bl/6) mRNA sequence	1382669	IMAGE Consortium	AI036073
2443 G	1	021HAMPC	nascent polypeptide-associated complex alpha polypeptide	1162959	IMAGE Consortium	AA770923
2444 G	3	021HAMPC	adaptor protein complex AP-2, alpha 2 subunit	1230717	IMAGE Consortium	AA879938
2445 G	5	021HAMPC	prolactin receptor	1333150	IMAGE Consortium	AI174135
2446 G	7	021HAMPC	dipeptidylpeptidase 4	1345705	IMAGE Consortium	AI154541
2447 G	9	021HAMPC	solute carrier family 16 (monocarboxylic acid transporters), member 1	1382413	IMAGE Consortium	AI046529
2448 G	11	021HAMPC	eukaryotic translation initiation factor 3, subunit 2 (beta, 36kD)	1382673	IMAGE Consortium	AI036075
2449 A	1	021OAMPD	solute carrier family 16 (monocarboxylic acid transporters), member 2	1382906	IMAGE Consortium	AI120278
2450 A	3	021OAMPD	ESTs, Highly similar to B47328 natural killer cell tumor-recognition protein - mouse [M.musculus]	1383381	IMAGE Consortium	AI120880
2451 A	5	021OAMPD	male enhanced antigen 1	1383741	IMAGE Consortium	AI180912
2452 A	7	021OAMPD	dual-specificity tyrosine-(Y)-phosphorylation regulated kinase 1a	1434361	IMAGE Consortium	AA866629
2453 A	9	021OAMPD				
2454 A	11	021OAMPD	CD52 antigen	330427	IMAGE Consortium	W15001
2455 C	1	021OAMPD	thyroid autoantigen 70 kDa	1382212	IMAGE Consortium	AA798305
2456 C	3	021OAMPD	proline rich protein expressed in brain	1384125	IMAGE Consortium	AI117268
2457 C	5	021OAMPD	myeloid differentiation primary response gene 88	1383757	IMAGE Consortium	AI180931
2458 C	7	021OAMPD	TYRO3 protein tyrosine kinase 3	1477926	IMAGE Consortium	AI155344
2459 C	9	021OAMPD	ESTs	1447839	IMAGE Consortium	AI158775
2460 C	11	021OAMPD	serine protease inhibitor 1-2	331065	IMAGE Consortium	W13979
2461 E	1	021OAMPD	transgelin	1382311	IMAGE Consortium	AI037523
2462 E	3	021OAMPD	flavin containing monooxygenase 1	1384169	IMAGE Consortium	AI117310
2463 E	5	021OAMPD	CD14 antigen	1383770	IMAGE Consortium	AI180938
2464 E	7	021OAMPD	Fas-associating protein with death domain	1433691	IMAGE Consortium	AA839509
2465 E	9	021OAMPD	fetal globin inducing factor	1447894	IMAGE Consortium	AI159330
2466 E	11	021OAMPD	AE-binding protein 1	331512	IMAGE Consortium	W13866

2467 G	1 021OAMPD	apolipoprotein D	1383150 IMAGE Consortium	AI120641
2468 G	3 021OAMPD	dihydrolipoamide branched chain transacylase E2	1382780 IMAGE Consortium	AI120185
2469 G	5 021OAMPD	alpha glucosidase 2, alpha neutral subunit	1383789 IMAGE Consortium	AI180958
2470 G	7 021OAMPD	poliovirus sensitivity	1447410 IMAGE Consortium	AI154217
2471 G	9 021OAMPD	arginase type II	1476909 IMAGE Consortium	AI154946
2472 G	11 021OAMPD	inhibitor of DNA binding 1	331996 IMAGE Consortium	W08321
2473 A	1 021VAMPE	ATPase, aminophospholipid transporter (APLT), class I, type 8A, member 1	2748986 IMAGE Consortium	AW495924
2474 A	3 021VAMPE	forkhead box M1	2699038 IMAGE Consortium	AW519587
2475 A	5 021VAMPE	Mus musculus mRNA for synaptogyrin 2 {IMAGE:333102}	333102 IMAGE Consortium	W15720
2476 A	7 021VAMPE	tropomyosin 2, beta	334540 IMAGE Consortium	W18330
2477 A	9 021VAMPE	RAD54 like (S. cerevisiae)	334046 IMAGE Consortium	W16123
2478 A	11 021VAMPE			
2479 C	1 021VAMPE	Burkitt lymphoma receptor 1	2749012 IMAGE Consortium	AW495941
2480 C	3 021VAMPE	heparan sulfate (glucosamine) 3-O-sulfotransferase 1	1617699 IMAGE Consortium	AA982702
2481 C	5 021VAMPE	dystroglycan 1	332866 IMAGE Consortium	W15608
2482 C	7 021VAMPE	annexin A1	334925 IMAGE Consortium	W17847
2483 C	9 021VAMPE	zinc finger protein 100	334887 IMAGE Consortium	W17826
2484 C	11 021VAMPE	Fibroblast inducible secreted protein {IMAGE:336325}	336325 IMAGE Consortium	W36541
2485 E	1 021VAMPE	procollagen, type IV, alpha 5	1749840 IMAGE Consortium	AI047053
2486 E	3 021VAMPE	lactate dehydrogenase 3, C chain, sperm specific	1746613 IMAGE Consortium	AI046562
2487 E	5 021VAMPE	desmin	332941 IMAGE Consortium	W15812
2488 E	7 021VAMPE	spinocerebellar ataxia 10 homolog (human)	335040 IMAGE Consortium	W17786
2489 E	9 021VAMPE			
2490 E	11 021VAMPE	ribosomal protein S14	335748 IMAGE Consortium	W20957
2491 G	1 021VAMPE	CD83 antigen	1617481 IMAGE Consortium	AA982557
2492 G	3 021VAMPE	arginine vasopressin	1617059 IMAGE Consortium	AA967824
2493 G	5 021VAMPE	paladin	333062 IMAGE Consortium	W15564
2494 G	7 021VAMPE	uteroglobin	333727 IMAGE Consortium	W16012
2495 G	9 021VAMPE	serine protease inhibitor 1-5	334942 IMAGE Consortium	W37032
2496 G	11 021VAMPE	ATP synthase, H <sup>+</sup> transporting mitochondrial F1 complex, beta subunit	335924 IMAGE Consortium	W36948
2497 A	1 0212AMPF	low density lipoprotein B	349681 IMAGE Consortium	W30115
2498 A	3 0212AMPF	vimentin	335768 IMAGE Consortium	AI390144
2499 A	5 0212AMPF	eukaryotic translation initiation factor 4E binding protein 1	349478 IMAGE Consortium	W30045
2500 A	7 0212AMPF	Mouse chromosome 6 BAC-284H12 (Research Genetics mouse BAC library) complete sequence {IMAGE:355935}	355935 IMAGE Consortium	W48119
2501 A	9 0212AMPF	nitrilase 1	367889 IMAGE Consortium	W54489
2502 A	11 0212AMPF	Mus musculus putative histone deacetylase (HD1) mRNA, partial cds {IMAGE:374877}	374877 IMAGE Consortium	W62765
2503 C	1 0212AMPF	ESTs, Highly similar to I58311 HMG-box containing protein 1 - rat [R.norvegicus]	348237 IMAGE Consortium	AI391046
2504 C	3 0212AMPF	calponin 2	335808 IMAGE Consortium	W34124

2505 C	5 0212AMPF	platelet-activating factor acetylhydrolase, isoform 1b, alpha1 subunit	349842 IMAGE Consortium	W33651
2506 C	7 0212AMPF	glutathione S-transferase, mu 2	367026 IMAGE Consortium	W54383
2507 C	9 0212AMPF	RAB3A, member RAS oncogene family	367954 IMAGE Consortium	W53795
2508 C	11 0212AMPF	thiosulfate sulfurtransferase, mitochondrial	373989 IMAGE Consortium	W62520
2509 E	1 0212AMPF	mitogen activated protein kinase 3	335228 IMAGE Consortium	W36966
2510 E	3 0212AMPF	heat shock protein, 25 kDa	335828 IMAGE Consortium	W34262
2511 E	5 0212AMPF	microtubule-associated protein 5	350186 IMAGE Consortium	W35860
2512 E	7 0212AMPF	vascular endothelial growth factor B	367377 IMAGE Consortium	W50682
2513 E	9 0212AMPF	serine protease inhibitor 1-2	367239 IMAGE Consortium	W54403
2514 E	11 0212AMPF	protein-tyrosine sulfotransferase 1	385445 IMAGE Consortium	W61385
2515 G	1 0212AMPF	amelogenin	335326 IMAGE Consortium	W36498
2516 G	3 0212AMPF	phospholipase c neighboring	335869 IMAGE Consortium	W34154
2517 G	5 0212AMPF	histidine ammonia lyase	350321 IMAGE Consortium	W40721
2518 G	7 0212AMPF	vasodilator-stimulated phosphoprotein	367694 IMAGE Consortium	W55649
2519 G	9 0212AMPF	mini chromosome maintenance deficient 7 (S. cerevisiae)	367244 IMAGE Consortium	W54667
2520 G	11 0212AMPF	avian reticuloendotheliosis viral (v-rel) oncogene homolog A	389904 IMAGE Consortium	W70696
2521 A	1 0211AJ7N	Internal_Control_E	Internal_Control_E	Controls
2522 A	3 0211AJ7N	Internal_Control_F	Internal_Control_F	Controls
2523 A	5 0211AJ7N	Internal_Control_G	Internal_Control_G	Controls
2524 A	7 0211AJ7N	Internal_Control_H	Internal_Control_H	Controls
2525 A	9 0211AJ7N	Internal_Control_I	Internal_Control_I	Controls
2526 A	11 0211AJ7N	Internal_Control_J	Internal_Control_J	Controls
2527 C	1 0211AJ7N	Internal_Control_K	Internal_Control_K	Controls
2528 C	3 0211AJ7N	Internal_Control_L	Internal_Control_L	Controls
2529 C	5 0211AJ7N	Internal_Control_M	Internal_Control_M	Controls
2530 C	7 0211AJ7N	Internal_Control_N	Internal_Control_N	Controls
2531 C	9 0211AJ7N	Internal_Control_O	Internal_Control_O	Controls
2532 C	11 0211AJ7N	Internal_Control_P	Internal_Control_P	Controls
2533 E	1 0211AJ7N	Internal_Control_Q	Internal_Control_Q	Controls
2534 E	3 0211AJ7N	Internal_Control_R	Internal_Control_R	Controls
2535 E	5 0211AJ7N	Internal_Control_S	Internal_Control_S	Controls
2536 E	7 0211AJ7N	Internal_Control_T	Internal_Control_T	Controls
2537 E	9 0211AJ7N	Internal_Control_U	Internal_Control_U	Controls
2538 E	11 0211AJ7N	Internal_Control_V	Internal_Control_V	Controls
2539 G	1 0211AJ7N	Internal_Control_W	Internal_Control_W	Controls
2540 G	3 0211AJ7N	Internal_Control_X	Internal_Control_X	Controls
2541 G	5 0211AJ7N	Internal_Control_Y	Internal_Control_Y	Controls
2542 G	7 0211AJ7N	Internal_Control_Z	Internal_Control_Z	Controls
2543 G	9 0211AJ7N	Internal_Control_AA	Internal_Control_AA	Controls
2544 G	11 0211AJ7N	Internal_Control_AB	Internal_Control_AB	Controls

2551 A	2 0217AJ7T	Control: Ratio 1:3 (Cy3:Cy5)	Ratio_A	Controls
2552 A	4 0217AJ7T	Control: Ratio 3:1 (Cy3:Cy5)	Ratio_B	Controls
2553 A	6 0217AJ7T	Control: Ratio 1:10 (Cy3:Cy5)	Ratio_C	Controls
2554 A	8 0217AJ7T	Control: Ratio 10:1 (Cy3:Cy5)	Ratio_D	Controls
2555 A	10 0217AJ7T	Control: Ratio 1:25 (Cy3:Cy5)	Ratio_E	Controls
2556 A	12 0217AJ7T	Control: Ratio 25:1 (Cy3:Cy5)	Ratio_F	Controls
2557 C	2 0217AJ7T	Control: Sensitivity 2pg	Sensitivity_A	Controls
2558 C	4 0217AJ7T	Control: Sensitivity 20pg	Sensitivity_B	Controls
2559 C	6 0217AJ7T	Control: Sensitivity 200pg	Sensitivity_C	Controls
2560 C	8 0217AJ7T	Control: Sensitivity 2000pg	Sensitivity_D	Controls
2561 C	10 0217AJ7T	Control: Buffer Only	Buffer_Only_Control	Controls
2562 C	12 0217AJ7T	Control: Buffer Only	Buffer_Only_Control	Controls
2563 E	2 0217AJ7T	Control: Internal	Internal_Control_A	Controls
2564 E	4 0217AJ7T	Control: Buffer Only	Buffer_Only_Control	Controls
2565 E	6 0217AJ7T	Control: Internal	Internal_Control_B	Controls
2566 E	8 0217AJ7T	Control: Buffer Only	Buffer_Only_Control	Controls
2567 E	10 0217AJ7T	Control: Internal	Internal_Control_C	Controls
2568 E	12 0217AJ7T	Control: Internal	Internal_Control_D	Controls
2569 G	2 0217AJ7T	Control: Complex Target (Mus musculus)	RNA_Quality_MM	Controls
2570 G	4 0217AJ7T	Control: Complex Target (Mus musculus)	RNA_Quality_MM	Controls
2571 G	6 0217AJ7T	Control: Complex Target (Mus musculus)	RNA_Quality_MM	Controls
2572 G	8 0217AJ7T	Control: Housekeeping (Alpha Tubulin)	Housekeeping_B	Controls
2573 G	10 0217AJ7T	Control: Housekeeping (Ribosomal S9)	Housekeeping_A	Controls
2574 G	12 0217AJ7T	Control: Housekeeping (23kD HBP)	Housekeeping_C	Controls
2575 A	2 021JAMMK	ESTs	313156 IMAGE Consortium	W10473
2576 A	4 021JAMMK	ESTs	315069 IMAGE Consortium	AI605702
2577 A	6 021JAMMK	ESTs	314814 IMAGE Consortium	AI605689
2578 A	8 021JAMMK	dystrobrevin, beta	438498 IMAGE Consortium	AA008091
2579 A	10 021JAMMK	adaptor-related protein complex AP-3, sigma 2 subunit	438308 IMAGE Consortium	AA008426
2580 A	12 021JAMMK	ESTs	426113 IMAGE Consortium	AA002746
2581 C	2 021JAMMK	ESTs, Weakly similar to CG10283 gene product [D.melanogaster]	303783 IMAGE Consortium	AI605553
2582 C	4 021JAMMK	ESTs	315169 IMAGE Consortium	AI605708
2583 C	6 021JAMMK	ESTs	315765 IMAGE Consortium	AI605753
2584 C	8 021JAMMK	cerebellin 1 precursor protein	425651 IMAGE Consortium	AA000130
2585 C	10 021JAMMK	Public domain EST	437777 IMAGE Consortium	
2586 C	12 021JAMMK	liver-specific bHLH-Zip transcription factor	539802 IMAGE Consortium	AA119984
2587 E	2 021JAMMK	myocyte enhancer factor 2C	303910 IMAGE Consortium	AI325323
2588 E	4 021JAMMK	ESTs, Highly similar to nucleoporin p58 [R.norvegicus]	315282 IMAGE Consortium	W12360
2589 E	6 021JAMMK	ESTs, Moderately similar to unnamed protein product [H.sapiens]	317292 IMAGE Consortium	AI536299
2590 E	8 021JAMMK	ESTs	425706 IMAGE Consortium	AA000370

2591 E	10 021JAMMK	ESTs	420493 IMAGE Consortium	AI430438
2592 E	12 021JAMMK	ESTs, Highly similar to CH-TOG PROTEIN [H.sapiens]	539822 IMAGE Consortium	AA123314
2593 G	2 021JAMMK	arachidonate 12-lipoxygenase, 12R type	314171 IMAGE Consortium	W09859
2594 G	4 021JAMMK	ESTs	313842 IMAGE Consortium	AI605635
2595 G	6 021JAMMK	ESTs	317328 IMAGE Consortium	AI536301
2596 G	8 021JAMMK	Max dimerization protein 3	427414 IMAGE Consortium	AA002410
2597 G	10 021JAMMK	ESTs, Moderately similar to unnamed protein product [H.sapiens]	438679 IMAGE Consortium	AI466598
2598 G	12 021JAMMK	ESTs	539834 IMAGE Consortium	AA123484
2599 A	2 021QAMML	Mus musculus mRNA for VEGF induced aminopeptidase, complete cds	540258 IMAGE Consortium	AA155451
2600 A	4 021QAMML	ESTs	540673 IMAGE Consortium	AA162746
2601 A	6 021QAMML	ubiquitin C-terminal hydrolase 37	539758 IMAGE Consortium	AA120166
2602 A	8 021QAMML	ESTs, Moderately similar to PROBABLE PEPTIDYL-PROLYL CIS-TRANS ISOMERASE C21E11.05C [Schizosaccharomyces pombe]	539212 IMAGE Consortium	AI892504
2603 A	10 021QAMML	DNA segment, Chr 7, ERATO Doi 805, expressed	538698 IMAGE Consortium	AA117468
2604 A	12 021QAMML	ESTs, Moderately similar to rA4 [R.norvegicus]	539664 IMAGE Consortium	AA123048
2605 C	2 021QAMML	ESTs	540285 IMAGE Consortium	AA154947
2606 C	4 021QAMML	ESTs, Moderately similar to PEROXISOME BIOSYNTHESIS PROTEIN PAY4 [Yarrowia lipolytica]	540792 IMAGE Consortium	AI892539
2607 C	6 021QAMML	CCCTC-binding factor	539047 IMAGE Consortium	AA122668
2608 C	8 021QAMML	ESTs, Highly similar to CHROMODOMAIN-HELICASE-DNA-BINDING PROTEIN CHD-1 [Mus musculus]	539269 IMAGE Consortium	AA116690
2609 C	10 021QAMML	ESTs	539427 IMAGE Consortium	AA123139
2610 C	12 021QAMML	ESTs	538941 IMAGE Consortium	AI550209
2611 E	2 021QAMML	proteasome (prosome, macropain) subunit, alpha type 6	540395 IMAGE Consortium	AA162010
2612 E	4 021QAMML	ESTs, Highly similar to ATP-CITRATE [Rattus norvegicus]	539417 IMAGE Consortium	AI325815
2613 E	6 021QAMML	ESTs	539081 IMAGE Consortium	AA117353
2614 E	8 021QAMML	ESTs	539288 IMAGE Consortium	AA116829
2615 E	10 021QAMML	stromal cell derived factor 2	538767 IMAGE Consortium	AA123282
2616 E	12 021QAMML	ESTs, Weakly similar to PARATHYMOSIN [Rattus norvegicus]	538958 IMAGE Consortium	AA122581
2617 G	2 021QAMML	ESTs, Weakly similar to ORF YKL201c [S.cerevisiae]	540429 IMAGE Consortium	AA155249
2618 G	4 021QAMML	diacylglycerol acyltransferase	539487 IMAGE Consortium	AA123152
2619 G	6 021QAMML	ESTs, Highly similar to ENV POLYPROTEIN PRECURSOR [Gibbon leukemia virus]	539102 IMAGE Consortium	AA116610
2620 G	8 021QAMML	teratocarcinoma expressed, serine rich	539294 IMAGE Consortium	AA116831
2621 G	10 021QAMML	DNA segment, Chr 3, ERATO Doi 194, expressed	538797 IMAGE Consortium	AI550196
2622 G	12 021QAMML	ESTs, Weakly similar to CG14041 gene product [D.melanogaster]	538966 IMAGE Consortium	AA117292
2623 A	2 021XAMMM	ESTs, Moderately similar to AF116721_71 PRO1853 [H.sapiens]	541231 IMAGE Consortium	AI550613

2624 A	4 021XAMMM			
2625 A	6 021XAMMM			
2626 A	8 021XAMMM	DNA segment, Chr 19, Wayne State University 54, expressed	805911 IMAGE Consortium	AA399766
2627 A	10 021XAMMM	ubiquitin C	560220 IMAGE Consortium	AA210396
2628 A	12 021XAMMM	ribosomal protein L3	560372 IMAGE Consortium	AA210554
2629 C	2 021XAMMM	ESTs, Highly similar to PHOSPHORIBOSYLAMINOIMIDAZOLECARBOXAMIDE FORMYLTRANSFERASE [Gallus gallus]	541256 IMAGE Consortium	AI892547
2630 C	4 021XAMMM	ESTs, Highly similar to POSSIBLE GLOBAL TRANSCRIPTION ACTIVATOR SNF2L1 [Homo sapiens]	540786 IMAGE Consortium	AA154634
2631 C	6 021XAMMM	ESTs, Highly similar to hypothetical SBBI03 protein [H.sapiens]	559860 IMAGE Consortium	AA211954
2632 C	8 021XAMMM	ESTs, Weakly similar to JAK3_MOUSE TYROSINE-PROTEIN KINASE JAK3 [M.musculus]	805918 IMAGE Consortium	AA399878
2633 C	10 021XAMMM	protein kinase inhibitor, gamma	560241 IMAGE Consortium	AA210230
2634 C	12 021XAMMM	FK506 binding protein 12-rapamycin associated protein 1	560376 IMAGE Consortium	AA210345
2635 E	2 021XAMMM	ESTs, Highly similar to unnamed protein product [H.sapiens]	541299 IMAGE Consortium	AI550619
2636 E	4 021XAMMM	ESTs	540847 IMAGE Consortium	AA162348
2637 E	6 021XAMMM	ESTs, Highly similar to ERGIC-53 PROTEIN PRECURSOR [Homo sapiens]	559956 IMAGE Consortium	AA265127
2638 E	8 021XAMMM	ESTs	805968 IMAGE Consortium	AA399725
2639 E	10 021XAMMM	ESTs	560276 IMAGE Consortium	AA210249
2640 E	12 021XAMMM	heterogeneous nuclear ribonucleoprotein A1	560388 IMAGE Consortium	AA210421
2641 G	2 021XAMMM	stimulated by retinoic acid gene 11	541316 IMAGE Consortium	AA139161
2642 G	4 021XAMMM	ESTs, Moderately similar to unnamed protein product [H.sapiens]	540875 IMAGE Consortium	AI550597
2643 G	6 021XAMMM	ESTs	559965 IMAGE Consortium	AA265164
2644 G	8 021XAMMM	secreted frizzled-related sequence protein 3	805986 IMAGE Consortium	AA399912
2645 G	10 021XAMMM	ESTs, Highly similar to MALATE OXIDOREDUCTASE [Homo sapiens]	560292 IMAGE Consortium	AA210237
2646 G	12 021XAMMM	ESTs	560394 IMAGE Consortium	AA210559
2647 A	2 0214AMMN	ESTs	560577 IMAGE Consortium	AA209796
2648 A	4 0214AMMN	gene trap locus 3	806098 IMAGE Consortium	AA432483
2649 A	6 0214AMMN	mannosidase 2, alpha B1	822750 IMAGE Consortium	AA474812
2650 A	8 0214AMMN	casein gamma	865060 IMAGE Consortium	AA469646
2651 A	10 0214AMMN	ESTs, Highly similar to AF223576_1 cysteine-rich protein NFX-1 [M.musculus]	822461 IMAGE Consortium	AI549643
2652 A	12 0214AMMN	Mus musculus cellular apoptosis susceptibility protein mRNA, complete cds	821867 IMAGE Consortium	AI549625
2653 C	2 0214AMMN	DNA segment, Chr 11, KL Mohlke 35	560749 IMAGE Consortium	AA212010
2654 C	4 0214AMMN	ESTs	806118 IMAGE Consortium	AA432495
2655 C	6 0214AMMN	ESTs	822760 IMAGE Consortium	AA474882

2656 C	8 0214AMMN	ADP-ribosylation factor 1	865104 IMAGE Consortium	AI614743
2657 C	10 0214AMMN	ethanol induced 6	822470 IMAGE Consortium	AI549642
2658 C	12 0214AMMN	ESTs, Weakly similar to AF217484_1 DnaQL [M.musculus]	821891 IMAGE Consortium	AA413170
2659 E	2 0214AMMN	RAB9, member RAS oncogene family	560871 IMAGE Consortium	AA212042
2660 E	4 0214AMMN	ESTs, Highly similar to U4/U6-associated RNA splicing factor [H.sapiens]	822581 IMAGE Consortium	AI549656
2661 E	6 0214AMMN	Mus musculus TSC22-related inducible leucine zipper 2 (Tilz2) mRNA, complete cds	822819 IMAGE Consortium	AI549681
2662 E	8 0214AMMN	Mus musculus cleavage and polyadenylation specificity factor 1 (Cpsf1) mRNA, complete cds	865201 IMAGE Consortium	AA450531
2663 E	10 0214AMMN	ESTs, Highly similar to AP50_HUMAN CLATHRIN COAT ASSEMBLY PROTEIN AP50 [M.musculus]	822480 IMAGE Consortium	AA474264
2664 E	12 0214AMMN	calcyclin binding protein	821944 IMAGE Consortium	AA474042
2665 G	2 0214AMMN	ESTs	560891 IMAGE Consortium	AA212046
2666 G	4 0214AMMN	centrin 2	822605 IMAGE Consortium	AA474223
2667 G	6 0214AMMN	ESTs, Moderately similar to SPC1_HUMAN MICROSOMAL SIGNAL PEPTIDASE 25 KDA SUBUNIT [H.sapiens]	822863 IMAGE Consortium	AI549687
2668 G	8 0214AMMN	BCL2/adenovirus E1B 19 kDa-interacting protein 1, NIP2	865276 IMAGE Consortium	AA450831
2669 G	10 0214AMMN	gene rich cluster, B gene	865394 IMAGE Consortium	AI643197
2670 G	12 0214AMMN	ESTs, Weakly similar to weak similarity to SP:YAD5_CLOAB [C.elegans]	821992 IMAGE Consortium	AI549632
2671 A	2 021BAMMO	DNA segment, Chr 15, Wayne State University 59, expressed	821405 IMAGE Consortium	AI552356
2672 A	4 021BAMMO	ESTs	821737 IMAGE Consortium	AA413037
2673 A	6 021BAMMO	cd164 antigen	865052 IMAGE Consortium	AA469634
2674 A	8 021BAMMO	Trk-fused gene	865151 IMAGE Consortium	AA450450
2675 A	10 021BAMMO	heat shock 10 kDa protein 1 (chaperonin 10)	865208 IMAGE Consortium	AA450526
2676 A	12 021BAMMO	synaptosomal-associated protein, 23kD	865280 IMAGE Consortium	AA450833
2677 C	2 021BAMMO	glucosamine-6-phosphate deaminase	821421 IMAGE Consortium	AA438071
2678 C	4 021BAMMO	serine/threonine kinase 5	806440 IMAGE Consortium	AA470284
2679 C	6 021BAMMO	ESTs, Moderately similar to T08694 hypothetical protein DKFZp564O092.1 - human [H.sapiens]	865069 IMAGE Consortium	AA469657
2680 C	8 021BAMMO	Mus musculus reticulon 3 (Rtn3) mRNA, complete cds	865159 IMAGE Consortium	AA450462
2681 C	10 021BAMMO	glutamine synthetase pseudogene 1	865213 IMAGE Consortium	AA450534
2682 C	12 021BAMMO	embigin	865286 IMAGE Consortium	AA450836
2683 E	2 021BAMMO	ESTs	806349 IMAGE Consortium	AA432749
2684 E	4 021BAMMO	DNA segment, Chr 2, ERATO Doi 435, expressed	806469 IMAGE Consortium	AA470294
2685 E	6 021BAMMO	ESTs, Highly similar to elongation factor SIII p15 subunit [R.norvegicus]	865076 IMAGE Consortium	AI614738
2686 E	8 021BAMMO	von Hippel-Lindau binding protein 1	865165 IMAGE Consortium	AA450464
2687 E	10 021BAMMO	protein kinase Chk2	865220 IMAGE Consortium	AA450529
2688 E	12 021BAMMO	cell division cycle 42 homolog (S. cerevisiae)	865302 IMAGE Consortium	AA450848

2689 G	2 021BAMMO	ESTs, Highly similar to PROTEIN TRANSPORT PROTEIN SEC13 [Saccharomyces cerevisiae]	821674 IMAGE Consortium	AA412780
2690 G	4 021BAMMO	carbonic anhydrase 14	821794 IMAGE Consortium	AI549611
2691 G	6 021BAMMO	calumenin	865097 IMAGE Consortium	AA469669
2692 G	8 021BAMMO	PPAR interacting protein PRIP	865169 IMAGE Consortium	AI614752
2693 G	10 021BAMMO	ESTs	865222 IMAGE Consortium	AA450530
2694 G	12 021BAMMO	chaperonin subunit 6a (zeta)	865310 IMAGE Consortium	AA450850
2695 A	2 021IAMMP	ESTs, Weakly similar to R32611_2 [H.sapiens]	865336 IMAGE Consortium	AA450905
2696 A	4 021IAMMP	ESTs, Highly similar to THRR_MOUSE THROMBIN RECEPTOR PRECURSOR [M.musculus]	865468 IMAGE Consortium	AI643210
2697 A	6 021IAMMP	ESTs, Moderately similar to T00075 hypothetical protein KIAA0461 - human [H.sapiens]	906063 IMAGE Consortium	AI553356
2698 A	8 021IAMMP	DEAD (aspartate-glutamate-alanine-aspartate) box polypeptide 9	905848 IMAGE Consortium	AA529409
2699 A	10 021IAMMP	ubiquitin-activating enzyme E1, Chr X	907025 IMAGE Consortium	AI551960
2700 A	12 021IAMMP	stress-induced phosphoprotein 1	907424 IMAGE Consortium	AA591433
2701 C	2 021IAMMP	brain expressed X-linked 3	865355 IMAGE Consortium	AA450920
2702 C	4 021IAMMP	syndecan binding protein 2	865547 IMAGE Consortium	AI552931
2703 C	6 021IAMMP	Mus musculus brain cDNA, clone MNCb-3848, similar to Homo sapiens CGI-28 protein mRNA	906070 IMAGE Consortium	AA529663
2704 C	8 021IAMMP	DnaJ homolog 10	905866 IMAGE Consortium	AA529424
2705 C	10 021IAMMP	DNA segment, Chr 2, Wayne State University 23, expressed	907097 IMAGE Consortium	AA591806
2706 C	12 021IAMMP	ESTs	906734 IMAGE Consortium	AA522320
2707 E	2 021IAMMP	ESTs	865362 IMAGE Consortium	AA450917
2708 E	4 021IAMMP	ESTs, Highly similar to The KIAA0191 gene is expressed ubiquitously. [H.sapiens]	865628 IMAGE Consortium	AA445064
2709 E	6 021IAMMP	ESTs, Moderately similar to dJ202I21.1 [H.sapiens]	906119 IMAGE Consortium	AA529734
2710 E	8 021IAMMP	ESTs	905916 IMAGE Consortium	AA529486
2711 E	10 021IAMMP	DNA polymerase delta 1, catalytic domain	907233 IMAGE Consortium	AA591989
2712 E	12 021IAMMP	laminin B1 subunit 1	906740 IMAGE Consortium	AI551931
2713 G	2 021IAMMP	ESTs, Moderately similar to AF099186_1 EH domain-containing protein EHD1 [M.musculus]	865373 IMAGE Consortium	AA444581
2714 G	4 021IAMMP	fragile X mental retardation syndrome 1 homolog	865652 IMAGE Consortium	AA445079
2715 G	6 021IAMMP	heat shock protein, 86 kDa 1	906182 IMAGE Consortium	AA529377
2716 G	8 021IAMMP	ESTs, Highly similar to unnamed protein product [H.sapiens]	866036 IMAGE Consortium	AI552958
2717 G	10 021IAMMP	ESTs	907245 IMAGE Consortium	AA591992
2718 G	12 021IAMMP	Mus musculus brain cDNA, clone MNCb-1820	907486 IMAGE Consortium	AA588981
2719 A	2 021PAMMQ	ESTs	906880 IMAGE Consortium	AA522414
2720 A	4 021PAMMQ	twist gene homolog, (Drosophila)	908717 IMAGE Consortium	AI613926
2721 A	6 021PAMMQ			
2722 A	8 021PAMMQ	mesoderm posterior 1	514023 IMAGE Consortium	AA062101



2723 A	10 021PAMMQ	meiosis expressed gene 1	514389 IMAGE Consortium	AI326529
2724 A	12 021PAMMQ	mammalian relative of DnaJ	515284 IMAGE Consortium	AI595605
2725 C	2 021PAMMQ	ribosomal protein L28	907608 IMAGE Consortium	AA555905
2726 C	4 021PAMMQ	ESTs, Moderately similar to Trip230 [H.sapiens]	908788 IMAGE Consortium	AI597478
2727 C	6 021PAMMQ	Notch gene homolog 1, (Drosophila)	514609 IMAGE Consortium	AA063885
2728 C	8 021PAMMQ	ESTs, Highly similar to RAS-RELATED PROTEIN RAB-11A [Homo sapiens; Rattus norvegicus; Canis familiaris; Oryctolagus cuniculus]	514051 IMAGE Consortium	AA062342
2729 C	10 021PAMMQ	ESTs, Highly similar to PHOSPHATIDYLSERINE DECARBOXYLASE PROENZYME [Cricetulus griseus]	515101 IMAGE Consortium	AI385983
2730 C	12 021PAMMQ	ESTs	515294 IMAGE Consortium	AI595606
2731 E	2 021PAMMQ	hepatoma-derived growth factor	906907 IMAGE Consortium	AI551948
2732 E	4 021PAMMQ	immunosuperfamily protein BI2	514412 IMAGE Consortium	AA061145
2733 E	6 021PAMMQ	carnitine palmitoyltransferase 1, muscle	513915 IMAGE Consortium	AA061685
2734 E	8 021PAMMQ	ESTs, Moderately similar to dJ206D15.3 [H.sapiens]	514150 IMAGE Consortium	AI595120
2735 E	10 021PAMMQ	ESTs	515111 IMAGE Consortium	AA061303
2736 E	12 021PAMMQ	ESTs, Weakly similar to unnamed protein product [H.sapiens]	514573 IMAGE Consortium	AA066091
2737 G	2 021PAMMQ	Mus musculus EST from clone 555961, full insert	907664 IMAGE Consortium	AA555659
2738 G	4 021PAMMQ	protein kinase inhibitor beta, cAMP dependent, testis specific	514418 IMAGE Consortium	AI326530
2739 G	6 021PAMMQ	ESTs	513935 IMAGE Consortium	AI595100
2740 G	8 021PAMMQ	DNA segment, Chr 15, Wayne State University 75, expressed	514200 IMAGE Consortium	AI595124
2741 G	10 021PAMMQ	actin-like 7b	515132 IMAGE Consortium	AA064431
2742 G	12 021PAMMQ	aquaporin 7	515317 IMAGE Consortium	AI595611
2743 A	2 021WAMMR	ESTs	515405 IMAGE Consortium	AI429111
2744 A	4 021WAMMR	protamine 2	514849 IMAGE Consortium	AI892208
2745 A	6 021WAMMR	chondroitin 4-sulfotransferase	515516 IMAGE Consortium	AI595005
2746 A	8 021WAMMR	DJ-1 protein	515611 IMAGE Consortium	AA065934
2747 A	10 021WAMMR	ESTs, Highly similar to CENTROMERIC PROTEIN E [Homo sapiens]	515854 IMAGE Consortium	AI385998
2748 A	12 021WAMMR	U2 small nuclear ribonucleoprotein polypeptide A'	516471 IMAGE Consortium	AI892224
2749 C	2 021WAMMR	acid phosphatase 1, soluble	514716 IMAGE Consortium	AA063861
2750 C	4 021WAMMR	choline/ethanolamine kinase	514888 IMAGE Consortium	AI385978
2751 C	6 021WAMMR	ESTs, Moderately similar to ACYL-COA DEHYDROGENASE, SHORT-CHAIN SPECIFIC PRECURSOR [Homo sapiens]	515590 IMAGE Consortium	AA065473
2752 C	8 021WAMMR	homeodomain interacting protein kinase 3	516346 IMAGE Consortium	AI386016
2753 C	10 021WAMMR	RAB11B, member RAS oncogene family	515860 IMAGE Consortium	AA066041
2754 C	12 021WAMMR	ESTs	517227 IMAGE Consortium	AA062427
2755 E	2 021WAMMR	ESTs, Weakly similar to SERINE/THREONINE-PROTEIN KINASE KKIALRE [Homo sapiens]	515483 IMAGE Consortium	AI385990

2756 E	4 021WAMMR	ESTs, Moderately similar to POLLEN SPECIFIC PROTEIN SF3 [ <i>Helianthus annuus</i> ]	514931 IMAGE Consortium	AI595272
2757 E	6 021WAMMR	t-complex-associated testis expressed 3	515679 IMAGE Consortium	AI326533
2758 E	8 021WAMMR	ESTs, Weakly similar to CG7872 gene product [ <i>D.melanogaster</i> ]	515661 IMAGE Consortium	AI595024
2759 E	10 021WAMMR	ESTs	515862 IMAGE Consortium	AI595534
2760 E	12 021WAMMR	stromal cell derived factor 1	517242 IMAGE Consortium	AA066069
2761 G	2 021WAMMR	Mus musculus mRNA for hypothetical protein (ORF2 ortholog)	515503 IMAGE Consortium	AI595007
2762 G	4 021WAMMR	diacylglycerol kinase, alpha (80 kDa)	514993 IMAGE Consortium	AA061283
2763 G	6 021WAMMR	t-complex protein 10b	516034 IMAGE Consortium	AI326540
2764 G	8 021WAMMR	ESTs	515677 IMAGE Consortium	AI595508
2765 G	10 021WAMMR	ESTs, Weakly similar to similarity to a number of different protein with tandem repeats of E-P [ <i>C.elegans</i> ]	515915 IMAGE Consortium	AI595539
2766 G	12 021WAMMR	CRIP1 protein	517258 IMAGE Consortium	AA060667
2767 A	2 0213AMMS	ESTs	517335 IMAGE Consortium	AA062298
2768 A	4 0213AMMS	ESTs, Highly similar to SUCCINATE DEHYDROGENASE [ <i>Homo sapiens</i> ]	517467 IMAGE Consortium	AA059601
2769 A	6 0213AMMS	ESTs, Weakly similar to AF161363_1 HSPC100 [ <i>H.sapiens</i> ]	516901 IMAGE Consortium	AI596627
2770 A	8 0213AMMS	cadherin 16	517590 IMAGE Consortium	AA066458
2771 A	10 0213AMMS	lactate dehydrogenase 2, B chain	517772 IMAGE Consortium	AA066634
2772 A	12 0213AMMS	ESTs, Moderately similar to ZINC FINGER PROTEIN GLO3 [ <i>Saccharomyces cerevisiae</i> ]	519243 IMAGE Consortium	AA106531
2773 C	2 0213AMMS	ESTs, Highly similar to HYPOTHETICAL 47.9 KD PROTEIN B0303.3 IN CHROMOSOME III [ <i>Caenorhabditis elegans</i> ]	516646 IMAGE Consortium	AI596600
2774 C	4 0213AMMS	a disintegrin and metalloprotease domain 2	516742 IMAGE Consortium	AI326547
2775 C	6 0213AMMS	endonuclease G	516928 IMAGE Consortium	AI892230
2776 C	8 0213AMMS	ESTs	517601 IMAGE Consortium	AI597111
2777 C	10 0213AMMS	paraoxonase 3	517793 IMAGE Consortium	AA066640
2778 C	12 0213AMMS	ESTs	519303 IMAGE Consortium	AI597344
2779 E	2 0213AMMS	heat shock protein, DNAJ-like 3	516668 IMAGE Consortium	AA108189
2780 E	4 0213AMMS	MAP kinase-activated protein kinase 5	516749 IMAGE Consortium	AI892226
2781 E	6 0213AMMS	protein tyrosine phosphatase, receptor type, G	516984 IMAGE Consortium	AI596632
2782 E	8 0213AMMS	solute carrier family 2 (facilitated glucose transporter), member 4	517625 IMAGE Consortium	AA066475
2783 E	10 0213AMMS	villin 2	517799 IMAGE Consortium	AA066685
2784 E	12 0213AMMS	ESTs, Weakly similar to ACDM_MOUSE ACYL-COA DEHYDROGENASE, MEDIUM-CHAIN SPECIFIC PRECURSOR [ <i>M.musculus</i> ]	519375 IMAGE Consortium	AA107083
2785 G	2 0213AMMS	ESTs, Moderately similar to C211_HUMAN PUTATIVE SURFACE GLYCOPROTEIN C21ORF1 PRECURSOR [ <i>H.sapiens</i> ]	517398 IMAGE Consortium	AA062452

2786 G	4 0213AMMS	ESTs, Highly similar to mRNA cleavage factor I 25 kDa subunit [H.sapiens]	516758 IMAGE Consortium	AA062282
2787 G	6 0213AMMS	ESTs, Weakly similar to RRAS_MOUSE RAS-RELATED PROTEIN R-RAS [M.musculus]	517728 IMAGE Consortium	AA066594
2788 G	8 0213AMMS	coatomer protein complex subunit alpha	517630 IMAGE Consortium	AA066474
2789 G	10 0213AMMS	ESTs, Moderately similar to unnamed protein product [H.sapiens]	517989 IMAGE Consortium	AA068028
2790 G	12 0213AMMS	ESTs, Weakly similar to GDIS_MOUSE RHO GDP-DISSOCIATION INHIBITOR 2 [M.musculus]	518682 IMAGE Consortium	AA105950
2791 A	2 021AAMMT	syntaxin binding protein 2	518923 IMAGE Consortium	AA106765
2792 A	4 021AAMMT	voltage-dependent anion channel 1	518812 IMAGE Consortium	AA089179
2793 A	6 021AAMMT	ESTs	519143 IMAGE Consortium	AA108057
2794 A	8 021AAMMT	cholecystokinin A receptor	519986 IMAGE Consortium	AA109799
2795 A	10 021AAMMT			
2796 A	12 021AAMMT	adducin 1 (alpha)	521078 IMAGE Consortium	AI386042
2797 C	2 021AAMMT	ESTs, Weakly similar to hypothetical protein [H.sapiens]	518951 IMAGE Consortium	AA106774
2798 C	4 021AAMMT	uromodulin	518837 IMAGE Consortium	AA089188
2799 C	6 021AAMMT			
2800 C	8 021AAMMT	ESTs	520009 IMAGE Consortium	AI552658
2801 C	10 021AAMMT	ESTs, Highly similar to SNC3_HUMAN SNRNA ACTIVATING PROTEIN COMPLEX 50 KDA SUBUNIT [H.sapiens]	520320 IMAGE Consortium	AA110912
2802 C	12 021AAMMT	inhibitor of DNA binding 3	521100 IMAGE Consortium	AA066816
2803 E	2 021AAMMT	COMPLEMENT FACTOR H PRECURSOR	518970 IMAGE Consortium	AA106098
2804 E	4 021AAMMT	solute carrier family 25 (mitochondrial carrier; adenine nucleotide translocator), member 5	518860 IMAGE Consortium	AA106338
2805 E	6 021AAMMT	ESTs, Weakly similar to HYPOTHETICAL 20.3 KD PROTEIN IN GCD14-POS18 INTERGENIC REGION [Saccharomyces cerevisiae]	519465 IMAGE Consortium	AI325578
2806 E	8 021AAMMT	ESTs, Highly similar to MITOCHONDRIAL RESPIRATORY CHAIN COMPLEXES ASSEMBLY PROTEIN RCA1 [Saccharomyces cerevisiae]	520056 IMAGE Consortium	AA109830
2807 E	10 021AAMMT			
2808 E	12 021AAMMT	ESTs, Weakly similar to HYPOTHETICAL 54.9 KD PROTEIN C02F5.7 IN CHROMOSOME III [Caenorhabditis elegans]	520410 IMAGE Consortium	AA105320
2809 G	2 021AAMMT	Mus musculus major histocompatibility locus class III regions Hsc70t gene, partial cds; smRNP, G7A, NG23, MutS homolog, CLCP, NG24, NG25, and NG26 genes, complete cds; and unknown genes	518988 IMAGE Consortium	AA106153
2810 G	4 021AAMMT	hydroxysteroid dehydrogenase-1, delta<5>-3-beta	518881 IMAGE Consortium	AI597312
2811 G	6 021AAMMT	ESTs	519529 IMAGE Consortium	AI597359
2812 G	8 021AAMMT	amine N-sulfotransferase	520093 IMAGE Consortium	AI892274
2813 G	10 021AAMMT	thiopurine methyltransferase	520354 IMAGE Consortium	AI552680

2814 G	12 021AAMMT	ESTs	520436 IMAGE Consortium	AI552688
2815 A	2 021HAMMU	growth hormone receptor	521281 IMAGE Consortium	AI386055
2816 A	4 021HAMMU	BCL2/adenovirus E1B 19 kDa-interacting protein 3-like	520879 IMAGE Consortium	AI552708
2817 A	6 021HAMMU	ESTs	443571 IMAGE Consortium	AA014081
2818 A	8 021HAMMU	ESTs	439776 IMAGE Consortium	AA008331
2819 A	10 021HAMMU	ESTs, Weakly similar to CALD_HUMAN CALDESMON [H.sapiens]	522462 IMAGE Consortium	AA067914
2820 A	12 021HAMMU	ESTs, Weakly similar to TIG1_HUMAN RETINOIC ACID RECEPTOR RESPONDER PROTEIN 1 [H.sapiens]	522805 IMAGE Consortium	AA087569
2821 C	2 021HAMMU	ESTs, Highly similar to FIBRINOGEN GAMMA-A AND B CHAIN PRECURSORS [Rattus norvegicus]	521341 IMAGE Consortium	AA066867
2822 C	4 021HAMMU	creatine kinase, muscle	520908 IMAGE Consortium	AI552711
2823 C	6 021HAMMU	trefoil factor 3, intestinal	445033 IMAGE Consortium	AI894059
2824 C	8 021HAMMU	cornichon (Drosophila)-like	440043 IMAGE Consortium	AA016502
2825 C	10 021HAMMU	cyclin-dependent kinase inhibitor 2B (p15, inhibits CDK4)	523205 IMAGE Consortium	AA067836
2826 C	12 021HAMMU	ESTs, Highly similar to UBIQUITIN-CONJUGATING ENZYME E2-19 KD [Arabidopsis thaliana]	522819 IMAGE Consortium	AI892297
2827 E	2 021HAMMU	ESTs	520635 IMAGE Consortium	AI552700
2828 E	4 021HAMMU	ESTs, Moderately similar to CPJ6_MOUSE CYTOCHROME P450 2J6 [M.musculus]	520992 IMAGE Consortium	AA066877
2829 E	6 021HAMMU	wingless-related MMTV integration site 4	445179 IMAGE Consortium	AI894060
2830 E	8 021HAMMU	procollagen, type XIII, alpha 1	440572 IMAGE Consortium	AA014637
2831 E	10 021HAMMU	ESTs	523249 IMAGE Consortium	AI614585
2832 E	12 021HAMMU	ribosomal protein L44	522865 IMAGE Consortium	AI552827
2833 G	2 021HAMMU	solute carrier family 22 (organic cation transporter), member 1-like	520685 IMAGE Consortium	AA066386
2834 G	4 021HAMMU	wingless related MMTV integration site 10b	439485 IMAGE Consortium	AI894037
2835 G	6 021HAMMU	ESTs, Moderately similar to HYPOTHETICAL PROTEIN HI1455 [Haemophilus influenzae]	445279 IMAGE Consortium	AA016799
2836 G	8 021HAMMU	ESTs, Weakly similar to KIAA0946 protein [H.sapiens]	441321 IMAGE Consortium	AA009264
2837 G	10 021HAMMU	ESTs, Weakly similar to NPT1_MOUSE RENAL SODIUM-DEPENDENT PHOSPHATE TRANSPORT PROTEIN 1 [M.musculus]	522551 IMAGE Consortium	AI552795
2838 G	12 021HAMMU	ESTs, Highly similar to AF132965_1 CGI-31 protein [H.sapiens]	522894 IMAGE Consortium	AA065863
2839 A	2 021OAMMV	keratin complex 2, gene 6g	522345 IMAGE Consortium	AI552779
2840 A	4 021OAMMV	ESTs, Highly similar to CREATINE KINASE, SARCOMERIC MITOCHONDRIAL PRECURSOR [Rattus norvegicus]	521882 IMAGE Consortium	AI325608
2841 A	6 021OAMMV	ESTs	522083 IMAGE Consortium	AI552771
2842 A	8 021OAMMV	nuclear factor, erythroid derived 2,-like 1	523198 IMAGE Consortium	AA067104
2843 A	10 021OAMMV			
2844 A	12 021OAMMV	ESTs	523648 IMAGE Consortium	AI614679
2845 C	2 021OAMMV	glucose-6-phosphate dehydrogenase X-linked	522368 IMAGE Consortium	AI326573

2846 C	4 021OAMMV	mitsugumin 29	521911 IMAGE Consortium	AA106336
2847 C	6 021OAMMV	coagulation factor VII	522105 IMAGE Consortium	AI326572
2848 C	8 021OAMMV	G protein gamma 3 linked gene	523266 IMAGE Consortium	AA067122
2849 C	10 021OAMMV	ESTs	523406 IMAGE Consortium	AA080175
2850 C	12 021OAMMV	ESTs	522927 IMAGE Consortium	AI552841
2851 E	2 021OAMMV	ESTs	522398 IMAGE Consortium	AA067625
2852 E	4 021OAMMV	carbonic anhydrase 4	521956 IMAGE Consortium	AA106894
2853 E	6 021OAMMV	serum albumin variant	521512 IMAGE Consortium	AI386058
2854 E	8 021OAMMV	ESTs, Highly similar to KIAA1004 protein [H.sapiens]	523278 IMAGE Consortium	AA080122
2855 E	10 021OAMMV	ESTs, Moderately similar to bA108L7.4 [H.sapiens]	523410 IMAGE Consortium	AA080257
2856 E	12 021OAMMV	malic enzyme, supernatant	523682 IMAGE Consortium	AA067147
2857 G	2 021OAMMV	histocompatibility 2, class II antigen E alpha	521720 IMAGE Consortium	AI326566
2858 G	4 021OAMMV	complement component factor i	521132 IMAGE Consortium	AI326560
2859 G	6 021OAMMV	ESTs, Moderately similar to UDP- GLUCURONOSYLTRANSFERASE 2B12 PRECURSOR, MICROSOMAL [Rattus norvegicus]	521599 IMAGE Consortium	AI386060
2860 G	8 021OAMMV	ESTs, Highly similar to N121_RAT NUCLEAR ENVELOPE PORE MEMBRANE PROTEIN POM 121 [R.norvegicus]	523284 IMAGE Consortium	AA079940
2861 G	10 021OAMMV	integrin-associated protein	523435 IMAGE Consortium	AA080266
2862 G	12 021OAMMV	ESTs, Highly similar to HYPOTHETICAL 272.0 KD PROTEIN C50C3.6 IN CHROMOSOME III [Caenorhabditis elegans]	523922 IMAGE Consortium	AI386244
2863 A	2 021VAMMW			
2864 A	4 021VAMMW	ESTs, Highly similar to pot. glutaminase [R.norvegicus]	524280 IMAGE Consortium	AA065691
2865 A	6 021VAMMW	ESTs, Highly similar to CNTF_MOUSE CILIARY NEUROTROPHIC FACTOR [M.musculus]	523861 IMAGE Consortium	AA067364
2866 A	8 021VAMMW	ESTs	524511 IMAGE Consortium	AA068540
2867 A	10 021VAMMW	ESTs	524748 IMAGE Consortium	AA067775
2868 A	12 021VAMMW	DNA segment, Chr 15, ERATO Doi 621, expressed	524077 IMAGE Consortium	AA066151
2869 C	2 021VAMMW	Hbs1-like (S. cerevisiae)	524091 IMAGE Consortium	AI386251
2870 C	4 021VAMMW	ESTs	524287 IMAGE Consortium	AA066204
2871 C	6 021VAMMW			
2872 C	8 021VAMMW	ESTs, Weakly similar to AF119384_1 putative CAMP protein [M.musculus]	524580 IMAGE Consortium	AI614949
2873 C	10 021VAMMW	ATPase, H+ transporting, lysosomal (vacuolar proton pump), alpha 70 kDa, isoform 2	524759 IMAGE Consortium	AA068612
2874 C	12 021VAMMW	t-complex protein 1, related sequence 1	524085 IMAGE Consortium	AA066253
2875 E	2 021VAMMW			
2876 E	4 021VAMMW			
2877 E	6 021VAMMW	heat shock protein cognate 70	523937 IMAGE Consortium	AA067797
2878 E	8 021VAMMW			
2879 E	10 021VAMMW	tetraspanin TM4-A homolog	524798 IMAGE Consortium	AA067830

2880 E	12 021VAMMW	Max interacting protein 1	524106 IMAGE Consortium	AI326586
2881 G	2 021VAMMW	ribosomal protein L8	524168 IMAGE Consortium	AA065602
2882 G	4 021VAMMW			
2883 G	6 021VAMMW	semaphorin cytoplasmic domain-associated protein 3A	523973 IMAGE Consortium	AA071995
2884 G	8 021VAMMW	ESTs, Weakly similar to A57369 anillin - fruit fly [D.melanogaster]	524610 IMAGE Consortium	AA068559
2885 G	10 021VAMMW	DNA segment, Chr 13, Wayne State University 64, expressed	524815 IMAGE Consortium	AA097042
2886 G	12 021VAMMW	ESTs	524111 IMAGE Consortium	AI614012
2887 A	2 0212AMMX	ESTs	524270 IMAGE Consortium	AI614796
2888 A	4 0212AMMX			
2889 A	6 0212AMMX	ESTs	524763 IMAGE Consortium	AA068212
2890 A	8 0212AMMX	ESTs, Highly similar to UBIQUINOL-CYTOCHROME C REDUCTASE COMPLEX 6.4 KD PROTEIN [Bos taurus]	524842 IMAGE Consortium	AA097097
2891 A	10 0212AMMX	DNA segment, Chr 15, ERATO Doi 405, expressed	524966 IMAGE Consortium	AA096982
2892 A	12 0212AMMX	ESTs, Highly similar to SERINE HYDROXYMETHYLTRANSFERASE, MITOCHONDRIAL [Oryctolagus cuniculus]	534113 IMAGE Consortium	AI386288
2893 C	2 0212AMMX	uridine phosphorylase	524272 IMAGE Consortium	AI614797
2894 C	4 0212AMMX	ESTs, Highly similar to T17343 hypothetical protein DKFZp586K0524.1 - human [H.sapiens]	524475 IMAGE Consortium	AA067818
2895 C	6 0212AMMX	ESTs	525071 IMAGE Consortium	AA089032
2896 C	8 0212AMMX	enoyl coenzyme A hydratase 1, peroxisomal	524846 IMAGE Consortium	AA097098
2897 C	10 0212AMMX			
2898 C	12 0212AMMX	ESTs, Weakly similar to CG7332 gene product [D.melanogaster]	534150 IMAGE Consortium	AI552046
2899 E	2 0212AMMX	Public domain EST	524292 IMAGE Consortium	
2900 E	4 0212AMMX			
2901 E	6 0212AMMX	ESTs	525114 IMAGE Consortium	AI643103
2902 E	8 0212AMMX	TPA regulated locus	524853 IMAGE Consortium	AA088959
2903 E	10 0212AMMX	high mobility group AT-hook 2	524992 IMAGE Consortium	AA097062
2904 E	12 0212AMMX	ESTs, Weakly similar to PERIPLASMIC DIVALENT CATION TOLERANCE PROTEIN CUTA [Escherichia coli]	534168 IMAGE Consortium	AI386290
2905 G	2 0212AMMX	junction-mediating and regulatory protein	524294 IMAGE Consortium	AI614803
2906 G	4 0212AMMX	synaptojanin 2 binding protein	524616 IMAGE Consortium	AI614958
2907 G	6 0212AMMX	ESTs	525182 IMAGE Consortium	AA096764
2908 G	8 0212AMMX	dynein, cytoplasmic, intermediate chain 2	524873 IMAGE Consortium	AA088964
2909 G	10 0212AMMX	ESTs, Moderately similar to AF231110_1 muscleblind [M.musculus]	533126 IMAGE Consortium	AA068812
2910 G	12 0212AMMX	ESTs	533454 IMAGE Consortium	AI551994
2911 A	2 0219AMMY			
2912 A	4 0219AMMY	ESTs	533993 IMAGE Consortium	AA072901
2913 A	6 0219AMMY	actin, alpha, cardiac	536543 IMAGE Consortium	AI325745

2914 A	8 0219AMMY	ribosomal protein L13	535209 IMAGE Consortium	AI325708
2915 A	10 0219AMMY	aryl-hydrocarbon receptor-interacting protein	535648 IMAGE Consortium	AA110870
2916 A	12 0219AMMY	ESTs	477230 IMAGE Consortium	AA048073
2917 C	2 0219AMMY	ESTs	533584 IMAGE Consortium	AA105069
2918 C	4 0219AMMY	calpain 6	536007 IMAGE Consortium	AA073358
2919 C	6 0219AMMY	ESTs	536591 IMAGE Consortium	AA079931
2920 C	8 0219AMMY	ADP-ribosylation factor-like protein 3	535258 IMAGE Consortium	AI892399
2921 C	10 0219AMMY	aristaless homeobox gene homolog (Drosophila)	534970 IMAGE Consortium	AA073367
2922 C	12 0219AMMY	ESTs	478026 IMAGE Consortium	AA049357
2923 E	2 0219AMMY	retinoblastoma 1	534320 IMAGE Consortium	AA072780
2924 E	4 0219AMMY	ESTs, Weakly similar to cDNA EST CEMSF67FB comes from this gene [C.elegans]	536154 IMAGE Consortium	AA073446
2925 E	6 0219AMMY	junction plakoglobin	536606 IMAGE Consortium	AA073952
2926 E	8 0219AMMY	ESTs, Weakly similar to ORF YGR066c [S.cerevisiae]	535310 IMAGE Consortium	AA086768
2927 E	10 0219AMMY	solute carrier family 17 (sodium/hydrogen exchanger), member 1	535707 IMAGE Consortium	AA106229
2928 E	12 0219AMMY	neurogranin (protein kinase C substrate, RC3)	476693 IMAGE Consortium	AA050507
2929 G	2 0219AMMY	caspase 8	533745 IMAGE Consortium	AI386280
2930 G	4 0219AMMY	Jun proto-oncogene related gene d1	550760 IMAGE Consortium	AA098137
2931 G	6 0219AMMY	guanylate cyclase 1, soluble, beta 3	536624 IMAGE Consortium	AI325750
2932 G	8 0219AMMY	RNA polymerase II 4 (14 kDa subunit)	535336 IMAGE Consortium	AA087794
2933 G	10 0219AMMY	hairy/enhancer-of-split related with YRPW motif 1	535777 IMAGE Consortium	AI552130
2934 G	12 0219AMMY	cathelin-like protein	478131 IMAGE Consortium	AI323038
2935 A	2 021GAMMZ	ESTs, Weakly similar to KIAA0699 protein [H.sapiens]	479147 IMAGE Consortium	AI510109
2936 A	4 021GAMMZ	paralemmin	479479 IMAGE Consortium	AA049097
2937 A	6 021GAMMZ	ESTs	572921 IMAGE Consortium	AA111613
2938 A	8 021GAMMZ	ESTs	572197 IMAGE Consortium	AI587825
2939 A	10 021GAMMZ	ESTs, Moderately similar to OOCYTE ZINC FINGER PROTEIN XLCOF7.1 [Xenopus laevis]	480211 IMAGE Consortium	AI385818
2940 A	12 021GAMMZ	trefoil factor 1	481581 IMAGE Consortium	AA060105
2941 C	2 021GAMMZ			
2942 C	4 021GAMMZ	transcriptional regulator protein	478924 IMAGE Consortium	AI385789
2943 C	6 021GAMMZ	ESTs	572938 IMAGE Consortium	AI587865
2944 C	8 021GAMMZ	ESTs	571586 IMAGE Consortium	AA108788
2945 C	10 021GAMMZ	ESTs	480314 IMAGE Consortium	AA058211
2946 C	12 021GAMMZ	zinc finger protein 95	482321 IMAGE Consortium	AI385486
2947 E	2 021GAMMZ	ESTs	478505 IMAGE Consortium	AA050034
2948 E	4 021GAMMZ	ESTs, Weakly similar to Similarity to H.influenza ribonuclease PH [C.elegans]	475434 IMAGE Consortium	AI509942
2949 E	6 021GAMMZ	ESTs	573864 IMAGE Consortium	AI530466
2950 E	8 021GAMMZ	Mus musculus stromal protein associated with thymii and lymph nodes short isoform (Spatial) mRNA, complete cds; alternatively spliced	573071 IMAGE Consortium	AI587887

2951 E	10 021GAMMZ	ESTs, Highly similar to METHIONYL-TRNA SYNTHETASE [Bacillus stearothermophilus]	481174 IMAGE Consortium	AI385460
2952 E	12 021GAMMZ	ESTs, Moderately similar to HYPOTHETICAL 9.8 KD PROTEIN ZK652.3 IN CHROMOSOME III [Caenorhabditis elegans]	481671 IMAGE Consortium	AI385481
2953 G	2 021GAMMZ	ESTs, Weakly similar to S59856 collagen alpha 1(III) chain precursor - mouse [M.musculus]	477865 IMAGE Consortium	AA048649
2954 G	4 021GAMMZ	glial cell line derived neurotrophic factor family receptor alpha 3	475497 IMAGE Consortium	AA050083
2955 G	6 021GAMMZ	mannosidase 1, beta	570537 IMAGE Consortium	AI587725
2956 G	8 021GAMMZ	zinc finger protein 235	570945 IMAGE Consortium	AI509384
2957 G	10 021GAMMZ	crystallin, alpha 1	481224 IMAGE Consortium	AI323082
2958 G	12 021GAMMZ	insulin-like growth factor binding protein 1	483145 IMAGE Consortium	AI892189
2959 A	2 021GAMN0	nuclear factor, interleukin 3, regulated	481915 IMAGE Consortium	AA060731
2960 A	4 021GAMN0	ESTs, Moderately similar to ZINC FINGER PROTEIN MFG1 [Mus musculus]	574368 IMAGE Consortium	AA119564
2961 A	6 021GAMN0	ESTs	574639 IMAGE Consortium	AI530474
2962 A	8 021GAMN0	NK6 transcription factor related, locus 2 (Drosophila)	576409 IMAGE Consortium	AA120355
2963 A	10 021GAMN0	ESTs	987456 IMAGE Consortium	AA572640
2964 A	12 021GAMN0			
2965 C	2 021GAMN0	ESTs, Weakly similar to LAR PROTEIN PRECURSOR [Homo sapiens]	481955 IMAGE Consortium	AA059639
2966 C	4 021GAMN0	nuclear factor of kappa light chain gene enhancer in B-cells 1, p105	575105 IMAGE Consortium	AI323229
2967 C	6 021GAMN0	ESTs, Weakly similar to MDC-3.13 isoform 1 [H.sapiens]	573953 IMAGE Consortium	AI594979
2968 C	8 021GAMN0	ESTs	575780 IMAGE Consortium	AA120431
2969 C	10 021GAMN0	ESTs, Weakly similar to t-BOP [M.musculus]	987549 IMAGE Consortium	AA575346
2970 C	12 021GAMN0	phosphatidylserine synthase 1	1001384 IMAGE Consortium	AA771633
2971 E	2 021GAMN0	immunoglobulin superfamily containing leucine-rich repeat	481983 IMAGE Consortium	AI892179
2972 E	4 021GAMN0	ESTs	573796 IMAGE Consortium	AI530452
2973 E	6 021GAMN0	ESTs	575429 IMAGE Consortium	AI530673
2974 E	8 021GAMN0	ESTs	576634 IMAGE Consortium	AA123447
2975 E	10 021GAMN0	mannose binding lectin, serum (C)	987696 IMAGE Consortium	AA793252
2976 E	12 021GAMN0	ectoplacental cone, invasive trophoblast giant cells, extraembryonic ectoderm and chorion sequence 21	1001397 IMAGE Consortium	AA771647
2977 G	2 021GAMN0	Mus musculus PACSIN3 (Pacsin3) mRNA, complete cds	482719 IMAGE Consortium	AI594932
2978 G	4 021GAMN0	ESTs	573802 IMAGE Consortium	AI530453
2979 G	6 021GAMN0	ESTs	574712 IMAGE Consortium	AA125231
2980 G	8 021GAMN0			
2981 G	10 021GAMN0	ficolin A	1001011 IMAGE Consortium	AA771355
2982 G	12 021GAMN0	ESTs, Weakly similar to rjs [M.musculus]	1001433 IMAGE Consortium	AI587946
2983 A	2 021NAMN1	phospholipid transfer protein	1001588 IMAGE Consortium	AA666595
2984 A	4 021NAMN1	ESTs	1002526 IMAGE Consortium	AA596737



2985 A	6 021NAMN1	ESTs, Highly similar to unnamed protein product [H.sapiens]	1002803 IMAGE Consortium	AA600485
2986 A	8 021NAMN1	CD53 antigen	1020649 IMAGE Consortium	AA646061
2987 A	10 021NAMN1	ESTs, Weakly similar to CTM [M.musculus]	1020943 IMAGE Consortium	AA646332
2988 A	12 021NAMN1			
2989 C	2 021NAMN1	ESTs	1001661 IMAGE Consortium	AI587972
2990 C	4 021NAMN1	ESTs, Highly similar to HYPOTHETICAL 94.2 KD PROTEIN C38D4.5 IN CHROMOSOME III [Caenorhabditis elegans]	1002545 IMAGE Consortium	AA596745
2991 C	6 021NAMN1	ESTs	1002829 IMAGE Consortium	AA600502
2992 C	8 021NAMN1	TATA box binding protein (Tbp)-associated factor, RNA polymerase I, B	1020674 IMAGE Consortium	AA646073
2993 C	10 021NAMN1	26S proteasome-associated pad1 homolog	1002086 IMAGE Consortium	AA684366
2994 C	12 021NAMN1	ESTs	1002328 IMAGE Consortium	AA596543
2995 E	2 021NAMN1	EST	1001674 IMAGE Consortium	AI504065
2996 E	4 021NAMN1	ESTs	1002569 IMAGE Consortium	AA596752
2997 E	6 021NAMN1	ESTs	1002836 IMAGE Consortium	AA600497
2998 E	8 021NAMN1	ESTs	1020772 IMAGE Consortium	AI594170
2999 E	10 021NAMN1	ESTs, Weakly similar to G10D_MOUSE PROBABLE G PROTEIN-COUPLED RECEPTOR G10D [M.musculus]	1002127 IMAGE Consortium	AA608087
3000 E	12 021NAMN1	palmitoyl-protein thioesterase	1002364 IMAGE Consortium	AA596482
3001 G	2 021NAMN1	hydroxysteroid 11-beta dehydrogenase 2	1001696 IMAGE Consortium	AI587979
3002 G	4 021NAMN1	ESTs	1002573 IMAGE Consortium	AA596754
3003 G	6 021NAMN1	ESTs	1002877 IMAGE Consortium	AI506831
3004 G	8 021NAMN1	ESTs	1020807 IMAGE Consortium	AA645765
3005 G	10 021NAMN1			
3006 G	12 021NAMN1	ESTs	1002376 IMAGE Consortium	AA596557
3007 A	2 021UAMN2	ESTs, Highly similar to IDN3-B [H.sapiens]	1001780 IMAGE Consortium	AA666703
3008 A	4 021UAMN2	nidogen 2	1021101 IMAGE Consortium	AA645955
3009 A	6 021UAMN2	ESTs, Weakly similar to AF156605_1 DXHXS6673E protein [M.musculus]	1038843 IMAGE Consortium	AA624556
3010 A	8 021UAMN2	ectonucleoside triphosphate diphosphohydrolase 5	1023593 IMAGE Consortium	AA647099
3011 A	10 021UAMN2	WD repeat domain 1	1039210 IMAGE Consortium	AA624634
3012 A	12 021UAMN2	ADP-ribosylation factor 4	1038975 IMAGE Consortium	AA624867
3013 C	2 021UAMN2	ESTs	1002528 IMAGE Consortium	AA596651
3014 C	4 021UAMN2	annexin A6	1038601 IMAGE Consortium	AA671168
3015 C	6 021UAMN2	Mus musculus mRNA for Clast1, complete cds	1038875 IMAGE Consortium	AA624578
3016 C	8 021UAMN2	beta-D-galactosidase fusion protein	1023634 IMAGE Consortium	AA799010
3017 C	10 021UAMN2	cyclin-dependent kinase inhibitor 1C (P57)	1038618 IMAGE Consortium	AA671166
3018 C	12 021UAMN2	myosin heavy chain, cardiac muscle, fetal	1038996 IMAGE Consortium	AA671135
3019 E	2 021UAMN2	T-cell receptor gamma, variable 4	1001830 IMAGE Consortium	AA666779
3020 E	4 021UAMN2	protein inhibitor of activated STAT protein-gamma)	1038636 IMAGE Consortium	AA671180
3021 E	6 021UAMN2	basigin	1038915 IMAGE Consortium	AA624721

3022 E	8 021UAMN2	ESTs, Weakly similar to AF113615_1 FH1/FH2 domain-containing protein FHOS [H.sapiens]	1023725 IMAGE Consortium	AA799097
3023 E	10 021UAMN2	a disintegrin and metalloproteinase domain 15 (metargidin)	1038670 IMAGE Consortium	AA671271
3024 E	12 021UAMN2	ESTs, Moderately similar to triadin [H.sapiens]	1039990 IMAGE Consortium	AA611321
3025 G	2 021UAMN2	ESTs	1001925 IMAGE Consortium	AA684073
3026 G	4 021UAMN2	pre B-cell leukemia transcription factor 2	1038654 IMAGE Consortium	AA671265
3027 G	6 021UAMN2	ribosomal protein S7	1038936 IMAGE Consortium	AA624837
3028 G	8 021UAMN2	microtubule-associated protein 4	1023794 IMAGE Consortium	AA799148
3029 G	10 021UAMN2	peroxiredoxin 4	1038694 IMAGE Consortium	AA671285
3030 G	12 021UAMN2	sema domain, immunoglobulin domain (Ig), transmembrane domain (TM) and short cytoplasmic domain, (semaphorin) 4A	1040009 IMAGE Consortium	AA611350
3031 A	2 0211AMN3	ESTs, Highly similar to hypothetical protein, similar to [H.sapiens]	1040190 IMAGE Consortium	AA615286
3032 A	4 0211AMN3	serine protease inhibitor 2	1077424 IMAGE Consortium	AA733490
3033 A	6 0211AMN3	aspartyl aminopeptidase	1077610 IMAGE Consortium	AA821963
3034 A	8 0211AMN3	major urinary protein 2	1077766 IMAGE Consortium	AA822105
3035 A	10 0211AMN3	myosin light chain 2	1077603 IMAGE Consortium	AA821970
3036 A	12 0211AMN3	ESTs	1040536 IMAGE Consortium	AA612331
3037 C	2 0211AMN3	ESTs, Moderately similar to I38909 damage-specific DNA binding protein 2 - human [H.sapiens]	1040237 IMAGE Consortium	AA611680
3038 C	4 0211AMN3	ESTs, Highly similar to NADH-UBIQUINONE OXIDOREDUCTASE 9 KD SUBUNIT PRECURSOR [Bos taurus]	1077477 IMAGE Consortium	AA727982
3039 C	6 0211AMN3	myoglobin	1077645 IMAGE Consortium	AA821997
3040 C	8 0211AMN3	ESTs	1040853 IMAGE Consortium	AA607702
3041 C	10 0211AMN3	catenin beta	1077648 IMAGE Consortium	AA822077
3042 C	12 0211AMN3	DNA segment, Chr 4, ERATO Doi 196, expressed	1040588 IMAGE Consortium	AA612369
3043 E	2 0211AMN3	ESTs	1040265 IMAGE Consortium	AI607260
3044 E	4 0211AMN3	DNA segment, Chr 7, Wayne State University 86, expressed	1077501 IMAGE Consortium	AA821903
3045 E	6 0211AMN3	cytochrome P450, 2c40	1077682 IMAGE Consortium	AA822002
3046 E	8 0211AMN3	squamous cell carcinoma antigen recognized by T-cells 1	1040886 IMAGE Consortium	AA607769
3047 E	10 0211AMN3	alcohol dehydrogenase 1, complex	1077724 IMAGE Consortium	AI507919
3048 E	12 0211AMN3	ESTs	1040600 IMAGE Consortium	AA612420
3049 G	2 0211AMN3	deleted in polyposis 1	1040404 IMAGE Consortium	AA612377
3050 G	4 0211AMN3	ESTs, Weakly similar to HSPC035 protein [H.sapiens]	1077503 IMAGE Consortium	AA821904
3051 G	6 0211AMN3	SEC61, alpha subunit (S. cerevisiae)	1077712 IMAGE Consortium	AA822026
3052 G	8 0211AMN3	pigment epithelium-derived factor	1077458 IMAGE Consortium	AA727967
3053 G	10 0211AMN3	ESTs	1077745 IMAGE Consortium	AA822106
3054 G	12 0211AMN3	ESTs, Moderately similar to interferon-g induced GTPase [M.musculus]	1040630 IMAGE Consortium	AA608395

3055 A	2 0218AMN4			
3056 A	4 0218AMN4	Treacher Collins Franceschetti syndrome 1, homolog	1092810 IMAGE Consortium	AA674285
3057 A	6 0218AMN4	cytochrome P450, 4a10	1092244 IMAGE Consortium	AA755385
3058 A	8 0218AMN4	EST	1140097 IMAGE Consortium	AA646049
3059 A	10 0218AMN4	ESTs	1139699 IMAGE Consortium	AA645595
3060 A	12 0218AMN4	ESTs	1139819 IMAGE Consortium	AI591753
3061 C	2 0218AMN4	ESTs, Weakly similar to CAC-1 [H.sapiens]	1040778 IMAGE Consortium	AA607572
3062 C	4 0218AMN4	ESTs	1092877 IMAGE Consortium	AA674392
3063 C	6 0218AMN4	ESTs, Weakly similar to Similarity to Yeast D-lactate dehydrogenase [C.elegans]	1092267 IMAGE Consortium	AA822473
3064 C	8 0218AMN4	ESTs, Weakly similar to HC1 ORF [M.musculus]	1148760 IMAGE Consortium	AA673705
3065 C	10 0218AMN4	5'(3')-deoxyribonucleotidase	1139708 IMAGE Consortium	AI591736
3066 C	12 0218AMN4			
3067 E	2 0218AMN4			
3068 E	4 0218AMN4	pancreatic lipase related protein 1	1092891 IMAGE Consortium	AA674409
3069 E	6 0218AMN4	proteasome (prosome, macropain) subunit, beta type 9 (large multifunctional protease 2)	1092290 IMAGE Consortium	AA822485
3070 E	8 0218AMN4	ESTs	1148764 IMAGE Consortium	AA673707
3071 E	10 0218AMN4	ESTs	1148464 IMAGE Consortium	AA646304
3072 E	12 0218AMN4			
3073 G	2 0218AMN4	ESTs, Weakly similar to pancreatic lipase related protein 1 [M.musculus]	1093179 IMAGE Consortium	AA675111
3074 G	4 0218AMN4	ESTs, Weakly similar to APA4_MOUSE APOLIPOPROTEIN A-IV PRECURSOR [M.musculus]	1092978 IMAGE Consortium	AA674450
3075 G	6 0218AMN4	ESTs	1139766 IMAGE Consortium	AI591743
3076 G	8 0218AMN4	colony stimulating factor 2 receptor, beta 2, low-affinity (granulocyte-macrophage)	1148775 IMAGE Consortium	AA673722
3077 G	10 0218AMN4	histocompatibility 2, T region locus 17	1148497 IMAGE Consortium	AA646461
3078 G	12 0218AMN4			
3079 A	2 021FAMN5			
3080 A	4 021FAMN5			
3081 A	6 021FAMN5	ribosomal protein L27a	1149890 IMAGE Consortium	AA822609
3082 A	8 021FAMN5	protease (prosome, macropain) 28 subunit, beta	1148961 IMAGE Consortium	AA655570
3083 A	10 021FAMN5	methyl-CpG binding domain protein 4	1149091 IMAGE Consortium	AA655907
3084 A	12 021FAMN5			
3085 C	2 021FAMN5			
3086 C	4 021FAMN5			
3087 C	6 021FAMN5	ESTs, Moderately similar to N-copine [M.musculus]	1148819 IMAGE Consortium	AA717161
3088 C	8 021FAMN5	actin-crosslinking protein 7	1149004 IMAGE Consortium	AA655639
3089 C	10 021FAMN5	putative lysophosphatidic acid acyltransferase	1149101 IMAGE Consortium	AA655761
3090 C	12 021FAMN5			
3091 E	2 021FAMN5			
3092 E	4 021FAMN5			

3093 E	6 021FAMN5	cytotoxic T lymphocyte-associated protein 2 alpha	1148836 IMAGE Consortium	AA717167
3094 E	8 021FAMN5	ESTs, Moderately similar to KIAA0301 [H.sapiens]	1149017 IMAGE Consortium	AA655654
3095 E	10 021FAMN5	male-specific lethal-3 homolog 1 (Drosophila)	1149107 IMAGE Consortium	AI644923
3096 E	12 021FAMN5	ESTs, Highly similar to RAS-RELATED PROTEIN RAB-14 [Rattus norvegicus]	1149669 IMAGE Consortium	AA656067
3097 G	2 021FAMN5			
3098 G	4 021FAMN5	ESTs, Highly similar to CLEAVAGE SIGNAL-1 PROTEIN [Homo sapiens]	1149605 IMAGE Consortium	AA655938
3099 G	6 021FAMN5	esterase 10	1148861 IMAGE Consortium	AA717191
3100 G	8 021FAMN5	ESTs, Highly similar to SET PROTEIN [Homo sapiens]	1149031 IMAGE Consortium	AA655672
3101 G	10 021FAMN5	DNA segment, Chr 1, Lubeck 1	1149123 IMAGE Consortium	AA655782
3102 G	12 021FAMN5	protein tyrosine phosphatase 4a2	1149712 IMAGE Consortium	AA656143
3103 A	2 021MAMN6	ESTs, Weakly similar to DEPP [H.sapiens]	1150208 IMAGE Consortium	AA929863
3104 A	4 021MAMN6	centrin 3	1150638 IMAGE Consortium	AA930531
3105 A	6 021MAMN6	ESTs	1193195 IMAGE Consortium	AA718459
3106 A	8 021MAMN6	peptide N-glycanase homolog (S.cerevisiae)	1193529 IMAGE Consortium	AA727030
3107 A	10 021MAMN6	ESTs, Weakly similar to T31099 myosin-RhoGAP protein, Myr 7 - rat [R.norvegicus]	1193659 IMAGE Consortium	AA726975
3108 A	12 021MAMN6	Kruppel-like factor 4 (gut)	1197054 IMAGE Consortium	AA733600
3109 C	2 021MAMN6	ESTs, Moderately similar to antigen NY-CO-3 [H.sapiens]	1149980 IMAGE Consortium	AA791621
3110 C	4 021MAMN6	ESTs	1150811 IMAGE Consortium	AA791924
3111 C	6 021MAMN6	mutS homolog 3 (E. coli)	1193217 IMAGE Consortium	AA718476
3112 C	8 021MAMN6	interleukin 7 receptor	1193536 IMAGE Consortium	AA727024
3113 C	10 021MAMN6	ESTs, Weakly similar to Zinc finger protein s11-6 [M.musculus]	1193690 IMAGE Consortium	AA726998
3114 C	12 021MAMN6	crystallin, mu	1197080 IMAGE Consortium	AA733693
3115 E	2 021MAMN6	gasdermin	1150269 IMAGE Consortium	AA930013
3116 E	4 021MAMN6	ESTs	1150918 IMAGE Consortium	AA795713
3117 E	6 021MAMN6	ESTs	1193220 IMAGE Consortium	AA718468
3118 E	8 021MAMN6	actin related protein 2/3 complex, subunit 3 (21 kDa)	1193545 IMAGE Consortium	AA726908
3119 E	10 021MAMN6	ESTs	1193698 IMAGE Consortium	AA727002
3120 E	12 021MAMN6	ESTs, Highly similar to nucleoporin Nup84 [R.norvegicus]	1197119 IMAGE Consortium	AA733633
3121 G	2 021MAMN6	ESTs	1151010 IMAGE Consortium	AA791555
3122 G	4 021MAMN6	angiotensin II, type I receptor-associated protein	1150956 IMAGE Consortium	AA791447
3123 G	6 021MAMN6	ESTs, Highly similar to seven transmembrane domain protein [H.sapiens]	1193244 IMAGE Consortium	AA718488
3124 G	8 021MAMN6	ESTs, Highly similar to PUFF SPECIFIC PROTEIN BX42 [Drosophila melanogaster]	1193580 IMAGE Consortium	AA726922
3125 G	10 021MAMN6	ESTs	1193702 IMAGE Consortium	AA727004
3126 G	12 021MAMN6	ESTs	1196445 IMAGE Consortium	AA794176
3127 A	2 021TAMN7	sphingosine kinase 1	1198042 IMAGE Consortium	AA725966
3128 A	4 021TAMN7	ADP-ribosylation factor 5	1197558 IMAGE Consortium	AA794574
3129 A	6 021TAMN7	ESTs	1209311 IMAGE Consortium	AA726264

3130 A	8 021TAMN7	ESTs	1210076 IMAGE Consortium	AA727580
3131 A	10 021TAMN7	protein tyrosine kinase 9	1223029 IMAGE Consortium	AA733857
3132 A	12 021TAMN7	M.musculus mRNA for ventricular alkali myosin light chain	1224217 IMAGE Consortium	AA795264
3133 C	2 021TAMN7	metaxin	1197343 IMAGE Consortium	AA681687
3134 C	4 021TAMN7	keratin complex 1, acidic, gene 16	1198320 IMAGE Consortium	AA726162
3135 C	6 021TAMN7	ESTs	1209406 IMAGE Consortium	AA726343
3136 C	8 021TAMN7	nuclear receptor subfamily 1, group H, member 2	1210152 IMAGE Consortium	AA727208
3137 C	10 021TAMN7	ESTs	1223076 IMAGE Consortium	AA733937
3138 C	12 021TAMN7	transcription factor 12	1224266 IMAGE Consortium	AA709476
3139 E	2 021TAMN7	ESTs, Highly similar to T42092 s-afadin - rat [R.norvegicus]	1196631 IMAGE Consortium	AA716921
3140 E	4 021TAMN7	ESTs	1196942 IMAGE Consortium	AI429054
3141 E	6 021TAMN7	ESTs	1209879 IMAGE Consortium	AA727298
3142 E	8 021TAMN7	DNA segment, Chr 2, ERATO Doi 473, expressed	1210220 IMAGE Consortium	AA727255
3143 E	10 021TAMN7	hypothetical protein, clone 1-82	1223125 IMAGE Consortium	AA734040
3144 E	12 021TAMN7	calcium channel, voltage-dependent, alpha2/delta subunit 1	1223549 IMAGE Consortium	AA734015
3145 G	2 021TAMN7	ESTs	1196643 IMAGE Consortium	AA716935
3146 G	4 021TAMN7	ESTs	1196983 IMAGE Consortium	AI429064
3147 G	6 021TAMN7	placental protein 11 related	1209901 IMAGE Consortium	AA727317
3148 G	8 021TAMN7	ESTs	1210238 IMAGE Consortium	AA727270
3149 G	10 021TAMN7	ESTs, Moderately similar to KIAA0830 protein [H.sapiens]	1223132 IMAGE Consortium	AA734032
3150 G	12 021TAMN7			
3151 A	2 0210AMN8	ESTs, Weakly similar to Ig heavy chain precursor [M.musculus]	1223891 IMAGE Consortium	AA667565
3152 A	4 0210AMN8			
3153 A	6 0210AMN8	phosphodiesterase 4B, cAMP specific	1211260 IMAGE Consortium	AA656268
3154 A	8 0210AMN8	hemopoietic cell kinase	597097 IMAGE Consortium	AA145674
3155 A	10 0210AMN8	Mus musculus germline J and C regions of immuoglobulin lambda-2 and germline J and C regions of the immuoglobulin lambda-4 pseudo-gene	596633 IMAGE Consortium	AA138265
3156 A	12 0210AMN8	myxovirus (influenza virus) resistance 1	596973 IMAGE Consortium	AA138836
3157 C	2 0210AMN8	potassium voltage-gated channel, Isk-related subfamily, member 1	1223990 IMAGE Consortium	AA667785
3158 C	4 0210AMN8	troponin C, cardiac/slow skeletal	1222578 IMAGE Consortium	AA815689
3159 C	6 0210AMN8	ferritin heavy chain	1211290 IMAGE Consortium	AA656293
3160 C	8 0210AMN8	procollagen-lysine, 2-oxoglutarate 5-dioxygenase 3	597159 IMAGE Consortium	AI893023
3161 C	10 0210AMN8	interferon consensus sequence binding protein	596643 IMAGE Consortium	AI386405
3162 C	12 0210AMN8	ATPase, H <sup>+</sup> transporting lysosomal (vacuolar proton pump), 9.2 kDa	1224378 IMAGE Consortium	AA755768
3163 E	2 0210AMN8	ESTs, Weakly similar to NTC1_MOUSE NEUROGENIC LOCUS NOTCH HOMOLOG PROTEIN 1 PRECURSOR [M.musculus]	1211398 IMAGE Consortium	AI593096

3164 E	4 0210AMN8	ESTs, Weakly similar to CA11_MOUSE COLLAGEN ALPHA 1(I) CHAIN PRECURSOR [M.musculus]	1222753 IMAGE Consortium	AA667546
3165 E	6 0210AMN8	synaptogyrin 1	1211308 IMAGE Consortium	AA656350
3166 E	8 0210AMN8	ESTs, Highly similar to hypothetical protein [H.sapiens]	597233 IMAGE Consortium	AA138401
3167 E	10 0210AMN8	DEAD/H (Asp-Glu-Ala-Asp/His) box polypeptide 20	596668 IMAGE Consortium	AI594219
3168 E	12 0210AMN8	human EST 573010	1224468 IMAGE Consortium	AA755846
3169 G	2 0210AMN8	ESTs, Weakly similar to CARG-BINDING FACTOR-A [Mus musculus]	1211452 IMAGE Consortium	AA655041
3170 G	4 0210AMN8	vesicle-associated membrane protein 5	1222787 IMAGE Consortium	AA667646
3171 G	6 0210AMN8	growth arrest specific 7	597375 IMAGE Consortium	AI324359
3172 G	8 0210AMN8	ESTs	596531 IMAGE Consortium	AA146439
3173 G	10 0210AMN8	ESTs	596681 IMAGE Consortium	AI594224
3174 G	12 0210AMN8	natriuretic peptide precursor type A	1224496 IMAGE Consortium	AA755870
3175 A	2 0217AMN9	actin, gamma 2, smooth muscle, enteric	1224697 IMAGE Consortium	AA756136
3176 A	4 0217AMN9	keratin-associated protein 5-4	1227337 IMAGE Consortium	AA739024
3177 A	6 0217AMN9	P40-8, functional	1227394 IMAGE Consortium	AA739104
3178 A	8 0217AMN9	ESTs, Weakly similar to S15660 [M.musculus]	1227570 IMAGE Consortium	AA738938
3179 A	10 0217AMN9	prprotein phosphatase 1G (formerly 2C), magnesium-dependent, gamma isoform	1227857 IMAGE Consortium	AA760317
3180 A	12 0217AMN9	uncoupling protein 2, mitochondrial	1228288 IMAGE Consortium	AA755022
3181 C	2 0217AMN9	ESTs	1224710 IMAGE Consortium	AI428582
3182 C	4 0217AMN9	ESTs	1227412 IMAGE Consortium	AA739124
3183 C	6 0217AMN9	complement component 4 (within H-2S)	1227418 IMAGE Consortium	AA739125
3184 C	8 0217AMN9	esterase 1	1227633 IMAGE Consortium	AA760070
3185 C	10 0217AMN9	ESTs, Highly similar to PSD-95/SAP90-associated protein-4 [R.norvegicus]	1228050 IMAGE Consortium	AA760121
3186 C	12 0217AMN9	tumor-associated calcium signal transducer 1	1228293 IMAGE Consortium	AA755032
3187 E	2 0217AMN9	DNA segment, Chr 17, human D6S51E	1224719 IMAGE Consortium	AI428588
3188 E	4 0217AMN9	ESTs, Highly similar to 2-OXOGLUTARATE DEHYDROGENASE E1 COMPONENT PRECURSOR [Homo sapiens]	1227434 IMAGE Consortium	AA738742
3189 E	6 0217AMN9	ESTs, Moderately similar to unnamed protein product [H.sapiens]	1227444 IMAGE Consortium	AA738745
3190 E	8 0217AMN9	keratin-associated protein 9-1	1227720 IMAGE Consortium	AA760130
3191 E	10 0217AMN9	annexin A2	1228117 IMAGE Consortium	AA760184
3192 E	12 0217AMN9	proteasome (prosome, macropain) subunit, beta type 7	1227976 IMAGE Consortium	AA760022
3193 G	2 0217AMN9	ESTs, Highly similar to p53 tumor suppressor-binding protein 1 [H.sapiens]	1224753 IMAGE Consortium	AA756177
3194 G	4 0217AMN9	ephrin A1	1227505 IMAGE Consortium	AA738854
3195 G	6 0217AMN9	UDP-N-acetyl-alpha-D-galactosamine:(N-acetylneuraminy)-galactosylglucosylceramide-beta-1, 4-N-acetylgalactosaminyltransferase	1227446 IMAGE Consortium	AA738746
3196 G	8 0217AMN9	procollagen, type IV, alpha 1	1227730 IMAGE Consortium	AA760135

3197 G	10 0217AMN9	ESTs, Highly similar to TRANSLOCON-ASSOCIATED PROTEIN, ALPHA SUBUNIT PRECURSOR [Canis familiaris]	1228153 IMAGE Consortium	AA760252
3198 G	12 0217AMN9	carboxypeptidase E	1227989 IMAGE Consortium	AA760047
3199 A	2 021RAMNA	thrombospondin 3	1229766 IMAGE Consortium	AA791128
3200 A	4 021RAMNA	mesoderm specific transcript	1259788 IMAGE Consortium	AI465074
3201 A	6 021RAMNA	ESTs	1261530 IMAGE Consortium	AA856109
3202 A	8 021RAMNA	ESTs, Weakly similar to AF198489_1 LBP-32 [H.sapiens]	1262013 IMAGE Consortium	AI466361
3203 A	10 021RAMNA	transmembrane domain protein regulated in adipocytes 40 kDa	1259999 IMAGE Consortium	AA839984
3204 A	12 021RAMNA	ATP-binding cassette, sub-family D (ALD), member 1	1260389 IMAGE Consortium	AA855347
3205 C	2 021RAMNA	ESTs	1229910 IMAGE Consortium	AA791250
3206 C	4 021RAMNA	ESTs	1243193 IMAGE Consortium	AA822240
3207 C	6 021RAMNA	ESTs	1261600 IMAGE Consortium	AA856164
3208 C	8 021RAMNA	ESTs, Highly similar to HP1-BP74 protein [M.musculus]	1262043 IMAGE Consortium	AA874100
3209 C	10 021RAMNA	ESTs, Weakly similar to DnaJ-like protein [M.musculus]	1260162 IMAGE Consortium	AA855618
3210 C	12 021RAMNA	casein beta	1260475 IMAGE Consortium	AA855396
3211 E	2 021RAMNA	ESTs, Highly similar to GASTRULA ZINC FINGER PROTEIN XLCGF52.1 [Xenopus laevis]	1229989 IMAGE Consortium	AA789386
3212 E	4 021RAMNA	ESTs, Highly similar to MYOSIN LIGHT CHAIN 1, SKELETAL MUSCLE ISOFORM [Mus musculus]	1243203 IMAGE Consortium	AA822257
3213 E	6 021RAMNA	ESTs	1261819 IMAGE Consortium	AA873958
3214 E	8 021RAMNA	ESTs, Weakly similar to AF209778_1 312 protein [D.melanogaster]	1261316 IMAGE Consortium	AA855944
3215 E	10 021RAMNA	Mus musculus monocarboxylate transporter 4 mRNA, complete cds	1260230 IMAGE Consortium	AA855235
3216 E	12 021RAMNA	ESTs, Moderately similar to polycystic kidney disease 1 protein [M.musculus]	1260507 IMAGE Consortium	AI464039
3217 G	2 021RAMNA	ESTs, Highly similar to DNA-DIRECTED RNA POLYMERASE III LARGEST SUBUNIT [Saccharomyces cerevisiae]	1230071 IMAGE Consortium	AA798891
3218 G	4 021RAMNA	phosphate cytidyltransferase 1, choline, alpha isoform	1259908 IMAGE Consortium	AA839928
3219 G	6 021RAMNA	mammalian relative of DnaJ	1261970 IMAGE Consortium	AA874130
3220 G	8 021RAMNA	ESTs	1261329 IMAGE Consortium	AA855968
3221 G	10 021RAMNA	dynein, cytoplasmic, light chain 1	1260273 IMAGE Consortium	AI463929
3222 G	12 021RAMNA	ras association (RalGDS/AF-6) domain family 1	1260544 IMAGE Consortium	AA855703
3223 A	2 021YAMNB	heterogeneous nuclear ribonucleoprotein D	1262348 IMAGE Consortium	AA880507
3224 A	4 021YAMNB	ESTs	1276832 IMAGE Consortium	AA880780
3225 A	6 021YAMNB	cyclin D2	1277386 IMAGE Consortium	AA879568
3226 A	8 021YAMNB	ESTs, Weakly similar to nuclear transcriptional repressor Mph1 [M.musculus]	1277906 IMAGE Consortium	AA882264
3227 A	10 021YAMNB	growth hormone	330360 IMAGE Consortium	W14555
3228 A	12 021YAMNB	ESTs	330887 IMAGE Consortium	AI390851

3229 C	2 021YAMNB	excision repair cross-complementing rodent repair deficiency, complementation group 4	1262374 IMAGE Consortium	AA880530
3230 C	4 021YAMNB			
3231 C	6 021YAMNB	ESTs, Weakly similar to 3-OXOACYL-[ACYL-CARRIER PROTEIN] REDUCTASE [Escherichia coli]	1276747 IMAGE Consortium	AA880732
3232 C	8 021YAMNB	Eph receptor A4	329775 IMAGE Consortium	AI325333
3233 C	10 021YAMNB	protein tyrosine phosphatase, receptor type, T	330377 IMAGE Consortium	AI390812
3234 C	12 021YAMNB	ESTs	318192 IMAGE Consortium	AI536342
3235 E	2 021YAMNB	ESTs	1262447 IMAGE Consortium	AA880601
3236 E	4 021YAMNB	ESTs	1262139 IMAGE Consortium	AA880430
3237 E	6 021YAMNB	Mus musculus thymus chemokine 1 mRNA, complete cds	1277525 IMAGE Consortium	AA882184
3238 E	8 021YAMNB	ESTs	329406 IMAGE Consortium	AI536378
3239 E	10 021YAMNB	cholinergic receptor, muscarinic 5	330410 IMAGE Consortium	AI327507
3240 E	12 021YAMNB	ESTs	318219 IMAGE Consortium	AI536344
3241 G	2 021YAMNB	bromodomain-containing 7	1262548 IMAGE Consortium	AA839190
3242 G	4 021YAMNB	hypothetical protein MNCb-5680	1262156 IMAGE Consortium	AI643009
3243 G	6 021YAMNB	ESTs	1276836 IMAGE Consortium	AA880782
3244 G	8 021YAMNB	ESTs	329551 IMAGE Consortium	AI451747
3245 G	10 021YAMNB	heparan sulfate (glucosamine) 3-O-sulfotransferase 3A	330423 IMAGE Consortium	W14854
3246 G	12 021YAMNB	peroxisome proliferator activated receptor gamma	317536 IMAGE Consortium	W34083
3247 A	2 0215AMNC	G protein-regulated inducer of neurite outgrowth 1	317921 IMAGE Consortium	W12945
3248 A	4 0215AMNC			
3249 A	6 0215AMNC			
3250 A	8 0215AMNC			
3251 A	10 0215AMNC	ESTs, Weakly similar to KIAA1140 protein [H.sapiens]	1279752 IMAGE Consortium	AA896165
3252 A	12 0215AMNC	sodium channel, nonvoltage-gated 1 beta	1280716 IMAGE Consortium	AA790595
3253 C	2 0215AMNC	annexin A10	317982 IMAGE Consortium	W12985
3254 C	4 0215AMNC			
3255 C	6 0215AMNC			
3256 C	8 0215AMNC			
3257 C	10 0215AMNC	ESTs	1279847 IMAGE Consortium	AA896294
3258 C	12 0215AMNC	EST AA673177	1280802 IMAGE Consortium	AA790651
3259 E	2 0215AMNC	ESTs, Highly similar to Y161_HUMAN HYPOTHETICAL PROTEIN KIAA0161 [H.sapiens]	318752 IMAGE Consortium	AI536363
3260 E	4 0215AMNC			
3261 E	6 0215AMNC			
3262 E	8 0215AMNC			
3263 E	10 0215AMNC	ESTs	1279943 IMAGE Consortium	AA896566
3264 E	12 0215AMNC	ESTs	1279408 IMAGE Consortium	AA983053
3265 G	2 0215AMNC	ESTs	318789 IMAGE Consortium	AI536365
3266 G	4 0215AMNC			
3267 G	6 0215AMNC			
3268 G	8 0215AMNC			



3269 G	10 0215AMNC	islet cell autoantigen 1, 69 kDa	1280107 IMAGE Consortium	AA895034
3270 G	12 0215AMNC	DNA segment, Chr 3, ERATO Doi 250, expressed	1279579 IMAGE Consortium	AA981316
3271 A	2 021CAMND	ESTs, Highly similar to RTS1 PROTEIN [Saccharomyces cerevisiae]	1279019 IMAGE Consortium	AA896671
3272 A	4 021CAMND	membrane bound C2 domain containing protein	1278569 IMAGE Consortium	AA895306
3273 A	6 021CAMND	advanced glycosylation end product-specific receptor	1350054 IMAGE Consortium	AA982240
3274 A	8 021CAMND	vitronectin	1350350 IMAGE Consortium	AA980366
3275 A	10 021CAMND	ESTs	598138 IMAGE Consortium	AI447461
3276 A	12 021CAMND	ESTs, Weakly similar to dJ977B1.1 [H.sapiens]	598399 IMAGE Consortium	AI592124
3277 C	2 021CAMND	ESTs, Highly similar to KIAA1017 protein [H.sapiens]	1278316 IMAGE Consortium	AA895132
3278 C	4 021CAMND	core promoter element binding protein	1278584 IMAGE Consortium	AA895718
3279 C	6 021CAMND	apolipoprotein H	1350386 IMAGE Consortium	AA980395
3280 C	8 021CAMND	ESTs	598738 IMAGE Consortium	AA168431
3281 C	10 021CAMND	ESTs	597523 IMAGE Consortium	AI592016
3282 C	12 021CAMND	ESTs, Weakly similar to TSGA_RAT TESTIS SPECIFIC PROTEIN A [R.norvegicus]	597681 IMAGE Consortium	AA154034
3283 E	2 021CAMND			
3284 E	4 021CAMND	ESTs, Highly similar to UDP-GalNAc:polypeptide N-acetylgalactosaminyltransferase T6 [R.norvegicus]	1279315 IMAGE Consortium	AA981116
3285 E	6 021CAMND	ESTs, Weakly similar to cytohesin-1 [M.musculus]	1367105 IMAGE Consortium	AI020517
3286 E	8 021CAMND	ESTs	598748 IMAGE Consortium	AA163023
3287 E	10 021CAMND	histocompatibility 2, O region beta locus	597562 IMAGE Consortium	AA154371
3288 E	12 021CAMND	ESTs, Weakly similar to zinc finger protein 64 [M.musculus]	599113 IMAGE Consortium	AA154358
3289 G	2 021CAMND	ESTs, Moderately similar to HSPC007 [H.sapiens]	1279123 IMAGE Consortium	AA981024
3290 G	4 021CAMND	ESTs, Moderately similar to RAS-LIKE PROTEIN TC10 [Homo sapiens]	1279344 IMAGE Consortium	AA981141
3291 G	6 021CAMND	ESTs, Moderately similar to ZINC FINGER PROTEIN 46 [Homo sapiens]	1367157 IMAGE Consortium	AI020307
3292 G	8 021CAMND	ESTs, Weakly similar to cell fate specification homolog MAB21L1 [M.musculus]	598801 IMAGE Consortium	AI592184
3293 G	10 021CAMND			
3294 G	12 021CAMND	ESTs, Weakly similar to similar to GTPase-activating proteins [H.sapiens]	598402 IMAGE Consortium	AI592121
3295 A	2 021JAMNE	lymphoid enhancer binding factor 1	597815 IMAGE Consortium	AA163420
3296 A	4 021JAMNE	dystrobrevin	775144 IMAGE Consortium	AA386892
3297 A	6 021JAMNE	fibroblast growth factor binding protein 1	775275 IMAGE Consortium	AA387137
3298 A	8 021JAMNE	ESTs, Highly similar to unnamed protein product [H.sapiens]	775411 IMAGE Consortium	AA386920
3299 A	10 021JAMNE	branched chain aminotransferase 2, mitochondrial	774935 IMAGE Consortium	AA387185
3300 A	12 021JAMNE	ubiquitin B	805186 IMAGE Consortium	AA473251
3301 C	2 021JAMNE	interferon-induced protein with tetratricopeptide repeats 2	597840 IMAGE Consortium	AA164018
3302 C	4 021JAMNE	Mus musculus mRNA for Ran binding protein 16	775165 IMAGE Consortium	AA386878

3303 C	6 021JAMNE	ESTs	775280 IMAGE Consortium	AA387132
3304 C	8 021JAMNE	histone 4 protein	775450 IMAGE Consortium	AA387386
3305 C	10 021JAMNE	ESTs, Highly similar to ZINC FINGER PROTEIN 46 [Homo sapiens]	789983 IMAGE Consortium	AA387945
3306 C	12 021JAMNE	ubiquitin-conjugating enzyme HR6A	804492 IMAGE Consortium	AI551532
3307 E	2 021JAMNE	6-pyruvoyl-tetrahydropterin synthase	597880 IMAGE Consortium	AA163644
3308 E	4 021JAMNE	ESTs	775180 IMAGE Consortium	AA386929
3309 E	6 021JAMNE	hepatoma-derived growth factor, related protein 2	775299 IMAGE Consortium	AA387018
3310 E	8 021JAMNE	ESTs, Weakly similar to Cox19p [S.cerevisiae]	775470 IMAGE Consortium	AA388124
3311 E	10 021JAMNE	ESTs	790029 IMAGE Consortium	AA387753
3312 E	12 021JAMNE			
3313 G	2 021JAMNE	EST	597892 IMAGE Consortium	AI447439
3314 G	4 021JAMNE	p21 (CDKN1A)-activated kinase 3	775190 IMAGE Consortium	AA387195
3315 G	6 021JAMNE	ferredoxin reductase	775316 IMAGE Consortium	AA386839
3316 G	8 021JAMNE	ESTs, Weakly similar to CG9591 gene product [D.melanogaster]	775475 IMAGE Consortium	AA388213
3317 G	10 021JAMNE	cartilage homeo protein 1	790077 IMAGE Consortium	AA388194
3318 G	12 021JAMNE	ESTs	805239 IMAGE Consortium	AA473304
3319 A	2 021QAMNF	ubiquitin-activating enzyme E1C	805401 IMAGE Consortium	AA473404
3320 A	4 021QAMNF	ESTs, Moderately similar to S65785 mel-13a protein - mouse [M.musculus]	789987 IMAGE Consortium	AA387961
3321 A	6 021QAMNF	myosin IXb	819726 IMAGE Consortium	AA437514
3322 A	8 021QAMNF	casein kinase I-alpha	819962 IMAGE Consortium	AA437784
3323 A	10 021QAMNF	ESTs, Moderately similar to AF161525_1 HSPC177 [H.sapiens]	805518 IMAGE Consortium	AI551664
3324 A	12 021QAMNF	v-crk-associated tyrosine kinase substrate	834133 IMAGE Consortium	AA413084
3325 C	2 021QAMNF	dolichyl-di-phosphooligosaccharide-protein glycotransferase	805451 IMAGE Consortium	AA474585
3326 C	4 021QAMNF	replication factor C, 140 kDa	790008 IMAGE Consortium	AA387984
3327 C	6 021QAMNF	transmembrane 4 superfamily member 6	819796 IMAGE Consortium	AA437557
3328 C	8 021QAMNF	ESTs	819968 IMAGE Consortium	AA437787
3329 C	10 021QAMNF	fructose bisphosphatase 2	805536 IMAGE Consortium	AA397241
3330 C	12 021QAMNF	H2A histone family, member Y	834197 IMAGE Consortium	AI596515
3331 E	2 021QAMNF	ESTs, Moderately similar to KIAA0092 gene product is distantly related to smooth muscle myosin. [H.sapiens]	804731 IMAGE Consortium	AA474398
3332 E	4 021QAMNF	ESTs, Weakly similar to SH2 domain-containing protein [M.musculus]	790042 IMAGE Consortium	AA387935
3333 E	6 021QAMNF	prediabetic NOD sera-reactive autoantigen	819820 IMAGE Consortium	AA437918
3334 E	8 021QAMNF	SWAP complex protein, 70 kDa	819985 IMAGE Consortium	AA437805
3335 E	10 021QAMNF	coding region determinant-binding protein	805632 IMAGE Consortium	AI604682
3336 E	12 021QAMNF	ESTs	834219 IMAGE Consortium	AA413192
3337 G	2 021QAMNF	ESTs	804738 IMAGE Consortium	AI551677
3338 G	4 021QAMNF	fibulin 2	790062 IMAGE Consortium	AA388038

3339 G	6 021QAMNF	distal-less homeobox 1	819842 IMAGE Consortium	AA437933
3340 G	8 021QAMNF	ESTs, Highly similar to PCAF associated factor 65 alpha [H.sapiens]	819989 IMAGE Consortium	AA437807
3341 G	10 021QAMNF	DNA segment, Chr 7, ERATO Doi 177, expressed	805647 IMAGE Consortium	AI604685
3342 G	12 021QAMNF	ESTs, Weakly similar to y+L amino acid transporter 1 [R.norvegicus]	834224 IMAGE Consortium	AA413187
3343 A	2 021XAMNG	ribosomal protein L26	834514 IMAGE Consortium	AI596527
3344 A	4 021XAMNG	telomeric repeat binding factor 1	819781 IMAGE Consortium	AA437554
3345 A	6 021XAMNG	tumor protein D52	819928 IMAGE Consortium	AA403797
3346 A	8 021XAMNG	ESTs, Highly similar to KIAA0594 protein [H.sapiens]	894023 IMAGE Consortium	AA516681
3347 A	10 021XAMNG	DNA segment, Chr 17, Wayne State University 166, expressed	894344 IMAGE Consortium	AA517413
3348 A	12 021XAMNG	protein phosphatase 2a, catalytic subunit, alpha isoform	894484 IMAGE Consortium	AA517691
3349 C	2 021XAMNG	m6a methyltransferase	833808 IMAGE Consortium	AA412911
3350 C	4 021XAMNG	ESTs, Highly similar to S68418 protein phosphatase 1M chain M110 isoform - rat [R.norvegicus]	819791 IMAGE Consortium	AA437678
3351 C	6 021XAMNG	COP9 (constitutive photomorphogenic), subunit 3 (Arabidopsis)	819930 IMAGE Consortium	AA403798
3352 C	8 021XAMNG	ESTs, Highly similar to KIAA0155 gene product [H.sapiens]	894189 IMAGE Consortium	AA517049
3353 C	10 021XAMNG	repeat family 3 gene	894389 IMAGE Consortium	AA517682
3354 C	12 021XAMNG	alkaline phosphatase 2, liver	894513 IMAGE Consortium	AA517588
3355 E	2 021XAMNG	ESTs	834552 IMAGE Consortium	AI596532
3356 E	4 021XAMNG	ESTs, Highly similar to JC4899 proline rich protein - rat [R.norvegicus]	819811 IMAGE Consortium	AA437683
3357 E	6 021XAMNG	ESTs, Highly similar to AF161530_1 HSPC182 [H.sapiens]	819935 IMAGE Consortium	AA403804
3358 E	8 021XAMNG	RNA polymerase II 1	894235 IMAGE Consortium	AA517228
3359 E	10 021XAMNG	ESTs, Highly similar to MITOCHONDRIAL LON PROTEASE HOMOLOG PRECURSOR [Homo sapiens]	933168 IMAGE Consortium	AA538210
3360 E	12 021XAMNG	ESTs, Highly similar to non-muscle alpha-actinin 1 [R.norvegicus]	894582 IMAGE Consortium	AA517712
3361 G	2 021XAMNG	ESTs, Moderately similar to dJ37E16.6 [H.sapiens]	833844 IMAGE Consortium	AA413005
3362 G	4 021XAMNG	heat shock protein, DNAJ-like 2	819831 IMAGE Consortium	AA437931
3363 G	6 021XAMNG	ESTs	819957 IMAGE Consortium	AA403674
3364 G	8 021XAMNG	Mus musculus WAVE-1 mRNA, complete cds	894283 IMAGE Consortium	AA517337
3365 G	10 021XAMNG	ESTs	894403 IMAGE Consortium	AA517689
3366 G	12 021XAMNG	synovial sarcoma, translocated to X chromosome	894584 IMAGE Consortium	AA517713
3367 A	2 0214AMNH	ESTs	893027 IMAGE Consortium	AA501135
3368 A	4 0214AMNH	ESTs	835372 IMAGE Consortium	AA445786
3369 A	6 0214AMNH	ESTs, Highly similar to NEDD-4 PROTEIN [Homo sapiens]	933012 IMAGE Consortium	AA538062
3370 A	8 0214AMNH	ESTs, Weakly similar to hypothetical protein [H.sapiens]	934277 IMAGE Consortium	AI551028
3371 A	10 0214AMNH	selective hybridizing clone	933934 IMAGE Consortium	AI550931

3372 A	12 0214AMNH	serine/threonine kinase receptor associated protein	934450 IMAGE Consortium	AA537006
3373 C	2 0214AMNH	ESTs, Moderately similar to unnamed protein product [H.sapiens]	892320 IMAGE Consortium	AA501245
3374 C	4 0214AMNH	ESTs, Moderately similar to putative transmembrane protein [H.sapiens]	834766 IMAGE Consortium	AA467087
3375 C	6 0214AMNH	nucleobindin	933228 IMAGE Consortium	AA541936
3376 C	8 0214AMNH	defender against cell death 1	933689 IMAGE Consortium	AA544207
3377 C	10 0214AMNH	ESTs, Highly similar to T14792 hypothetical protein DKFZp586G0322.1 - human [H.sapiens]	933943 IMAGE Consortium	AA545634
3378 C	12 0214AMNH	ESTs, Moderately similar to p80 [R.norvegicus]	934477 IMAGE Consortium	AA537028
3379 E	2 0214AMNH	ESTs	892418 IMAGE Consortium	AA501317
3380 E	4 0214AMNH	ESTs	834814 IMAGE Consortium	AI596546
3381 E	6 0214AMNH	ESTs	934073 IMAGE Consortium	AA544570
3382 E	8 0214AMNH	ESTs	933751 IMAGE Consortium	AI550913
3383 E	10 0214AMNH	ESTs, Weakly similar to nuclear protein np95 [M.musculus]	934172 IMAGE Consortium	AA544754
3384 E	12 0214AMNH	DNA segment, Chr 4, Wayne State University 27, expressed	934543 IMAGE Consortium	AA536695
3385 G	2 0214AMNH	tubulin, beta 5	835161 IMAGE Consortium	AA467239
3386 G	4 0214AMNH	ESTs	834953 IMAGE Consortium	AA467289
3387 G	6 0214AMNH	ESTs	934079 IMAGE Consortium	AA544572
3388 G	8 0214AMNH	ESTs	934489 IMAGE Consortium	AA536628
3389 G	10 0214AMNH	ESTs, Weakly similar to AF167320_1 zinc finger protein ZFP113 [M.musculus]	934219 IMAGE Consortium	AA544678
3390 G	12 0214AMNH	ESTs	934574 IMAGE Consortium	AA536706
3391 A	2 021BAMNI	Mus musculus transcription factor GATA-6 (Gata6) mRNA, complete cds	934696 IMAGE Consortium	AA536899
3392 A	4 021BAMNI	Mus musculus low density lipoprotein receptor-related protein 2 LRP2 (Lrp2) mRNA, partial cds	934860 IMAGE Consortium	AA617362
3393 A	6 021BAMNI	ESTs, Weakly similar to NED4_MOUSE NEDD-4 PROTEIN [M.musculus]	617281 IMAGE Consortium	AA174621
3394 A	8 021BAMNI	ESTs	618351 IMAGE Consortium	AI639809
3395 A	10 021BAMNI	ESTs, Weakly similar to cDNA EST EMBL:C08125 comes from this gene [C.elegans]	962624 IMAGE Consortium	AA562129
3396 A	12 021BAMNI	ESTs	946026 IMAGE Consortium	AI585440
3397 C	2 021BAMNI			
3398 C	4 021BAMNI	ESTs, Weakly similar to Edp1 protein [M.musculus]	934891 IMAGE Consortium	AA617448
3399 C	6 021BAMNI	nuclear receptor interacting protein 1	617333 IMAGE Consortium	AA174819
3400 C	8 021BAMNI	ESTs	617634 IMAGE Consortium	AA170569
3401 C	10 021BAMNI			
3402 C	12 021BAMNI			
3403 E	2 021BAMNI	DNA segment, Chr 18, Wayne State University 181, expressed	934773 IMAGE Consortium	AA617418
3404 E	4 021BAMNI	ESTs, Highly similar to UBIQUITIN-CONJUGATING	934911 IMAGE Consortium	AA617391

3405 E	6 021BAMNI	ENZYME E2-17 KD [ <i>Drosophila melanogaster</i> ]	616642 IMAGE Consortium	AI640047
3406 E	8 021BAMNI	ESTs, Weakly similar to Rfxank [ <i>M.musculus</i> ]	617980 IMAGE Consortium	AA175847
3407 E	10 021BAMNI	schlafen 3	947441 IMAGE Consortium	AI558178
3408 E	12 021BAMNI	ESTs, Moderately similar to dJ622L5.2 [ <i>H.sapiens</i> ]	946107 IMAGE Consortium	AI550945
3409 G	2 021BAMNI	ESTs, Weakly similar to DNA helicase [ <i>H.sapiens</i> ]	934796 IMAGE Consortium	AA617422
3410 G	4 021BAMNI	ESTs, Weakly similar to transcription factor BFCOL1 [ <i>M.musculus</i> ]	934916 IMAGE Consortium	AA617452
3411 G	6 021BAMNI	ESTs	616688 IMAGE Consortium	AI640053
3412 G	8 021BAMNI	DNA segment, Chr 7, ERATO Doi 684, expressed	599080 IMAGE Consortium	AI592230
3413 G	10 021BAMNI	ESTs	962761 IMAGE Consortium	AI597196
3414 G	12 021BAMNI	zinc finger protein 259	946146 IMAGE Consortium	AI550950
3415 A	2 021IAMNJ	ESTs, Highly similar to GASTRULATION SPECIFIC PROTEIN G12 [ <i>Danio rerio</i> ]	945667 IMAGE Consortium	AA546346
3416 A	4 021IAMNJ	DNA segment, Chr 2, ERATO Doi 434, expressed	935207 IMAGE Consortium	AI558253
3417 A	6 021IAMNJ	ESTs	945236 IMAGE Consortium	AA545611
3418 A	8 021IAMNJ	RaBP1 associated Eps domain containing protein	962770 IMAGE Consortium	AA561920
3419 A	10 021IAMNJ	ESTs	977267 IMAGE Consortium	AI645420
3420 A	12 021IAMNJ	ESTs, Weakly similar to collagen alpha 1(XVIII) chain [ <i>M.musculus</i> ]	976880 IMAGE Consortium	AA562544
3421 C	2 021IAMNJ	DNA segment, Chr 11, ERATO Doi 676, expressed	935015 IMAGE Consortium	AA617560
3422 C	4 021IAMNJ	t-complex protein 1, related sequence 1	935266 IMAGE Consortium	AA617224
3423 C	6 021IAMNJ	beta-transducin repeat containing protein	945292 IMAGE Consortium	AA546559
3424 C	8 021IAMNJ	ESTs, Weakly similar to AF119384_1 putative CAMP protein [ <i>M.musculus</i> ]	962795 IMAGE Consortium	AA561934
3425 C	10 021IAMNJ	ESTs	977283 IMAGE Consortium	AA560314
3426 C	12 021IAMNJ	ESTs	976885 IMAGE Consortium	AA562555
3427 E	2 021IAMNJ	ESTs	935084 IMAGE Consortium	AI558241
3428 E	4 021IAMNJ	apolipoprotein CII	935269 IMAGE Consortium	AA617232
3429 E	6 021IAMNJ	ESTs	945384 IMAGE Consortium	AA546759
3430 E	8 021IAMNJ	junction cell adhesion molecule	962825 IMAGE Consortium	AI597200
3431 E	10 021IAMNJ	ESTs	977341 IMAGE Consortium	AI645426
3432 E	12 021IAMNJ	isocitrate dehydrogenase 2 (NADP+), mitochondrial	976890 IMAGE Consortium	AA562548
3433 G	2 021IAMNJ	coronin, actin binding protein 1B	935121 IMAGE Consortium	AA617588
3434 G	4 021IAMNJ	ESTs, Highly similar to LYAR_MOUSE CELL GROWTH REGULATING NUCLEOLAR PROTEIN□ [ <i>M.musculus</i> ]	935275 IMAGE Consortium	AI558261
3435 G	6 021IAMNJ	ESTs, Weakly similar to CABA_MOUSE CARG-BINDING FACTOR-A [ <i>M.musculus</i> ]	945464 IMAGE Consortium	AI585413
3436 G	8 021IAMNJ	guanine nucleotide binding protein, alpha 12	962853 IMAGE Consortium	AA561797
3437 G	10 021IAMNJ	ESTs, Highly similar to N107_RAT NUCLEAR PORE COMPLEX PROTEIN NUP107 [ <i>R.norvegicus</i> ]	977481 IMAGE Consortium	AA560885
3438 G	12 021IAMNJ	solute carrier family 25 (mitochondrial carnitine/acylcarnitine translocase), member 20	976902 IMAGE Consortium	AA562563

3439 A	2 021PAMNK	telomerase binding protein p23	976974 IMAGE Consortium	AA562974
3440 A	4 021PAMNK	gap junction membrane channel protein beta 3	988523 IMAGE Consortium	AA638198
3441 A	6 021PAMNK	GTP cyclohydrolase 1	988764 IMAGE Consortium	AA638354
3442 A	8 021PAMNK	EST	977687 IMAGE Consortium	AI504848
3443 A	10 021PAMNK	ESTs	991435 IMAGE Consortium	AA574784
3444 A	12 021PAMNK	ubiquitin specific protease 25	991698 IMAGE Consortium	AI604608
3445 C	2 021PAMNK	ESTs, Highly similar to D-BETA-HYDROXYBUTYRATE DEHYDROGENASE PRECURSOR [Rattus norvegicus]	976981 IMAGE Consortium	AI645404
3446 C	4 021PAMNK	ESTs, Moderately similar to PTD017 [H.sapiens]	988531 IMAGE Consortium	AA638200
3447 C	6 021PAMNK	Mus musculus 2P1 protein mRNA, complete cds	988794 IMAGE Consortium	AI586351
3448 C	8 021PAMNK	ESTs	977944 IMAGE Consortium	AA563125
3449 C	10 021PAMNK	small acidic protein sid2057p	991481 IMAGE Consortium	AA612050
3450 C	12 021PAMNK	DNA segment, Chr 5, ERATO Doi 287, expressed	991743 IMAGE Consortium	AI604615
3451 E	2 021PAMNK	ESTs, Highly similar to R32184_3 [H.sapiens]	976989 IMAGE Consortium	AA562986
3452 E	4 021PAMNK	ESTs, Weakly similar to hypothetical protein [H.sapiens]	989282 IMAGE Consortium	AA571416
3453 E	6 021PAMNK	DNA segment, Chr 6, ERATO Doi 263, expressed	988875 IMAGE Consortium	AI586357
3454 E	8 021PAMNK	ESTs	989049 IMAGE Consortium	AA571252
3455 E	10 021PAMNK	ESTs	991520 IMAGE Consortium	AA612076
3456 E	12 021PAMNK	solute carrier family 35 (UDP-galactose transporter), member 2	991866 IMAGE Consortium	AA574838
3457 G	2 021PAMNK	ESTs, Highly similar to ABC3_HUMAN ATP-BINDING CASSETTE, SUB-FAMILY A, MEMBER 3 [H.sapiens]	977736 IMAGE Consortium	AA562577
3458 G	4 021PAMNK			
3459 G	6 021PAMNK	ESTs, Moderately similar to I49636 DNA-binding protein - mouse [M.musculus]	988893 IMAGE Consortium	AA572422
3460 G	8 021PAMNK	eukaryotic translation initiation factor 4A1	989093 IMAGE Consortium	AA575544
3461 G	10 021PAMNK	glia maturation factor, beta	991537 IMAGE Consortium	AA575789
3462 G	12 021PAMNK	tyrosine 3-monooxygenase/tryptophan 5-monooxygenase activation protein, sigma polypeptide	989155 IMAGE Consortium	AA571363
3463 A	2 021WAMNL	ESTs	989786 IMAGE Consortium	AA793446
3464 A	4 021WAMNL	ESTs	990236 IMAGE Consortium	AA590095
3465 A	6 021WAMNL	ESTs, Weakly similar to NAH1_MOUSE SODIUM/HYDROGEN EXCHANGER 1 [M.musculus]	1004123 IMAGE Consortium	AI606379
3466 A	8 021WAMNL	ESTs, Highly similar to PUTATIVE 30.7 KD METHYLTRANSFERASE IN TSM1-ARE1 INTERGENIC REGION [Saccharomyces cerevisiae]	1004339 IMAGE Consortium	AA600437
3467 A	10 021WAMNL	myelocytomatosis oncogene	1004629 IMAGE Consortium	AA608146
3468 A	12 021WAMNL	ESTs, Moderately similar to restricted expressed proliferation associated protein 100 [H.sapiens]	992617 IMAGE Consortium	AA607867
3469 C	2 021WAMNL	ESTs, Weakly similar to KIAA0841 protein [H.sapiens]	989854 IMAGE Consortium	AA793527
3470 C	4 021WAMNL	sphingomyelin phosphodiesterase 1, acid lysosomal	990556 IMAGE Consortium	AA589977
3471 C	6 021WAMNL	ESTs, Highly similar to DUS6_RAT DUAL SPECIFICITY PROTEIN PHOSPHATASE 6 [R.norvegicus]	1004139 IMAGE Consortium	AA600596

3472 C	8 021WAMNL	muscle and microspikes RAS	1004375 IMAGE Consortium	AA600451
3473 C	10 021WAMNL	testis expressed gene 10	1004645 IMAGE Consortium	AA608159
3474 C	12 021WAMNL	open reading frame 5	992822 IMAGE Consortium	AA608002
3475 E	2 021WAMNL			
3476 E	4 021WAMNL	ESTs, Weakly similar to trg [R.norvegicus]	990643 IMAGE Consortium	AA590160
3477 E	6 021WAMNL	ESTs	1004157 IMAGE Consortium	AA600600
3478 E	8 021WAMNL	thioredoxin reductase 1	1004386 IMAGE Consortium	AA600454
3479 E	10 021WAMNL	ESTs	1004668 IMAGE Consortium	AA608282
3480 E	12 021WAMNL	DNA segment, Chr 5, Bucan 26 expressed	992861 IMAGE Consortium	AI604638
3481 G	2 021WAMNL	ESTs	989948 IMAGE Consortium	AI607043
3482 G	4 021WAMNL	ESTs	990780 IMAGE Consortium	AA590624
3483 G	6 021WAMNL	signal sequence receptor, delta	1004217 IMAGE Consortium	AA600634
3484 G	8 021WAMNL	DNA j protein b9	1004406 IMAGE Consortium	AA600462
3485 G	10 021WAMNL	Jun-B oncogene	1004694 IMAGE Consortium	AA608298
3486 G	12 021WAMNL	ESTs, Highly similar to hypothetical protein [H.sapiens]	1005007 IMAGE Consortium	AA608175
3487 A	2 0213AMNM	ESTs, Weakly similar to I49625 testosterone 16a-hydroxylase type b - mouse [M.musculus]	1005063 IMAGE Consortium	AA608330
3488 A	4 0213AMNM	nidogen 1	1005138 IMAGE Consortium	AA606605
3489 A	6 0213AMNM	N-methylpurine-DNA glycosylase	1004828 IMAGE Consortium	AA606852
3490 A	8 0213AMNM	polymerase, gamma	1004924 IMAGE Consortium	AA607085
3491 A	10 0213AMNM			
3492 A	12 0213AMNM	protective protein for beta-galactosidase	1005637 IMAGE Consortium	AA607008
3493 C	2 0213AMNM	ESTs, Highly similar to HYPOTHETICAL 13.6 KD PROTEIN IN NUP170-ILS1 INTERGENIC REGION [Saccharomyces cerevisiae]	1005078 IMAGE Consortium	AA608333
3494 C	4 0213AMNM	TBX1 protein	1005145 IMAGE Consortium	AA606770
3495 C	6 0213AMNM	recombining binding protein suppressor of hairless (Drosophila)	1004845 IMAGE Consortium	AA606873
3496 C	8 0213AMNM	ESTs, Highly similar to EUKARYOTIC INITIATION FACTOR 4 GAMMA [Oryctolagus cuniculus]	1004940 IMAGE Consortium	AA607098
3497 C	10 0213AMNM	ESTs	1006021 IMAGE Consortium	AA607065
3498 C	12 0213AMNM	prohibitin	1005668 IMAGE Consortium	AA607019
3499 E	2 0213AMNM	guanylate kinase 1	1005084 IMAGE Consortium	AA608335
3500 E	4 0213AMNM	ESTs, Weakly similar to HYPOTHETICAL 92.1 KD PROTEIN C24H6.03 IN CHROMOSOME I [Schizosaccharomyces pombe]	1004638 IMAGE Consortium	AA608141
3501 E	6 0213AMNM	surfeit gene 4	1004858 IMAGE Consortium	AA606969
3502 E	8 0213AMNM	tyrosine 3-monooxygenase/tryptophan 5-monooxygenase activation protein, eta polypeptide	1004943 IMAGE Consortium	AA607106
3503 E	10 0213AMNM	protein tyrosine phosphatase, non-receptor type 21	1006111 IMAGE Consortium	AA607174
3504 E	12 0213AMNM	ESTs	1005692 IMAGE Consortium	AA607115
3505 G	2 0213AMNM	adenylosuccinate lyase	1005097 IMAGE Consortium	AA606587
3506 G	4 0213AMNM	adenylate kinase 4	1004729 IMAGE Consortium	AI594232

3507 G	6 0213AMNM	transcription elongation factor B (SIII), polypeptide 1 (15 kDa),-like	1004862 IMAGE Consortium	AA606971
3508 G	8 0213AMNM	ESTs	1004950 IMAGE Consortium	AA607103
3509 G	10 0213AMNM	ESTs, Highly similar to similar to Schizosaccharomyces pombe cut1+ protein which regulates spindle pole body duplication. [H.sapiens]	1006179 IMAGE Consortium	AA624147
3510 G	12 0213AMNM	ESTs, Highly similar to 2208369A signal peptidase:SUBUNIT	1005694 IMAGE Consortium	AA607116
3511 A	2 021AAMNN	Yamaguchi sarcoma viral (v-yes) oncogene homolog	1005808 IMAGE Consortium	AA606937
3512 A	4 021AAMNN	ribosomal protein S29	1005884 IMAGE Consortium	AA606952
3513 A	6 021AAMNN			
3514 A	8 021AAMNN	ESTs, Highly similar to SRPR_HUMAN SIGNAL RECOGNITION PARTICLE RECEPTOR ALPHA SUBUNIT [H.sapiens]	1005439 IMAGE Consortium	AA608348
3515 A	10 021AAMNN	heterogeneous nuclear ribonucleoprotein U	1005751 IMAGE Consortium	AA607267
3516 A	12 021AAMNN	fusion 2 (human)	620208 IMAGE Consortium	AA177916
3517 C	2 021AAMNN	DNA segment, Chr 17, human D6S52E	1005811 IMAGE Consortium	AA606941
3518 C	4 021AAMNN	actin, gamma, cytoplasmic	1005899 IMAGE Consortium	AA606826
3519 C	6 021AAMNN	ESTs, Highly similar to HYPOTHETICAL 66.5 KD PROTEIN F02A9.5 IN CHROMOSOME III [Caenorhabditis elegans]	1005990 IMAGE Consortium	AA607041
3520 C	8 021AAMNN	telomerase associated protein 1	1005464 IMAGE Consortium	AA608355
3521 C	10 021AAMNN	eomesodermin homolog (Xenopus laevis)	1005792 IMAGE Consortium	AA606660
3522 C	12 021AAMNN	ESTs, Moderately similar to KINESIN LIGHT CHAIN [Rattus norvegicus]	620274 IMAGE Consortium	AA172733
3523 E	2 021AAMNN	C184L-22	1005829 IMAGE Consortium	AA606681
3524 E	4 021AAMNN	DNA segment, Chr 11, ERATO Doi 333, expressed	1005904 IMAGE Consortium	AA606955
3525 E	6 021AAMNN	ESTs, Weakly similar to HYPOTHETICAL 39.2 KD PROTEIN IN EGT2-KRE1 INTERGENIC REGION [Saccharomyces cerevisiae]	1005996 IMAGE Consortium	AA607043
3526 E	8 021AAMNN			
3527 E	10 021AAMNN	DNA segment, Chr 9, ERATO Doi 660, expressed	1005817 IMAGE Consortium	AA606678
3528 E	12 021AAMNN	ESTs	621007 IMAGE Consortium	AA178450
3529 G	2 021AAMNN	glutathione peroxidase 4	1005832 IMAGE Consortium	AA606943
3530 G	4 021AAMNN	ESTs, Weakly similar to GENERAL NEGATIVE REGULATOR OF TRANSCRIPTION SUBUNIT 1 [Saccharomyces cerevisiae]	1005926 IMAGE Consortium	AA606835
3531 G	6 021AAMNN	procollagen, type IV, alpha 2	1005368 IMAGE Consortium	AA607232
3532 G	8 021AAMNN			
3533 G	10 021AAMNN	H2A histone family, member Z	1005846 IMAGE Consortium	AA606684
3534 G	12 021AAMNN	ESTs	620303 IMAGE Consortium	AI604209
3535 A	2 021HAMNO	ESTs	620464 IMAGE Consortium	AI604226
3536 A	4 021HAMNO	ESTs	620936 IMAGE Consortium	AA177238
3537 A	6 021HAMNO	ESTs	1006166 IMAGE Consortium	AI594303



3538 A	8 021HAMNO	ESTs	1021789 IMAGE Consortium	AI048063
3539 A	10 021HAMNO	ESTs, Weakly similar to acetylglucosaminyltransferase-like protein [M.musculus]	1006886 IMAGE Consortium	AI594129
3540 A	12 021HAMNO	DNA segment, Chr 1, ERATO Doi 251, expressed	1038083 IMAGE Consortium	AA624525
3541 C	2 021HAMNO	ESTs	621235 IMAGE Consortium	AI592298
3542 C	4 021HAMNO	retinoblastoma binding protein 4	1037544 IMAGE Consortium	AA623561
3543 C	6 021HAMNO	DNA segment, Chr 2, ERATO Doi 612, expressed	1022269 IMAGE Consortium	AI647127
3544 C	8 021HAMNO	ESTs	1006503 IMAGE Consortium	AA624291
3545 C	10 021HAMNO	X-linked lymphocyte-regulated complex	1022928 IMAGE Consortium	AA623498
3546 C	12 021HAMNO	DNA segment, Chr 8, ERATO Doi 325, expressed	1038097 IMAGE Consortium	AI642314
3547 E	2 021HAMNO	ESTs	621270 IMAGE Consortium	AI592302
3548 E	4 021HAMNO	DNA segment, Chr 12, ERATO Doi 771, expressed	1037575 IMAGE Consortium	AA623586
3549 E	6 021HAMNO	ESTs, Weakly similar to T25G3.1 [C.elegans]	1021582 IMAGE Consortium	AI047949
3550 E	8 021HAMNO	thimet oligopeptidase 1	1006531 IMAGE Consortium	AA624339
3551 E	10 021HAMNO	ESTs	1022988 IMAGE Consortium	AA623448
3552 E	12 021HAMNO	Ser/Arg-related nuclear matrix protein	1038099 IMAGE Consortium	AA624541
3553 G	2 021HAMNO	protocadherin 7	620676 IMAGE Consortium	AI604245
3554 G	4 021HAMNO			
3555 G	6 021HAMNO	ESTs	1023007 IMAGE Consortium	AA623473
3556 G	8 021HAMNO	S-adenosylmethionine decarboxylase 2	1006558 IMAGE Consortium	AA615119
3557 G	10 021HAMNO	TATA box binding protein-like protein	1006968 IMAGE Consortium	AA637018
3558 G	12 021HAMNO	ESTs, Moderately similar to AF077037_1 HSPC013 [H.sapiens]	1038120 IMAGE Consortium	AA624675
3559 A	2 021OAMNP	ESTs, Weakly similar to protein B [M.musculus]	1037589 IMAGE Consortium	AA623591
3560 A	4 021OAMNP	ribosomal protein L22	1037768 IMAGE Consortium	AA638878
3561 A	6 021OAMNP	cell division cycle 37 homolog (S. cerevisiae)	1037882 IMAGE Consortium	AA638943
3562 A	8 021OAMNP	ESTs	1038267 IMAGE Consortium	AA671203
3563 A	10 021OAMNP	ESTs	1038353 IMAGE Consortium	AA671314
3564 A	12 021OAMNP	metaxin 2	1038449 IMAGE Consortium	AA624514
3565 C	2 021OAMNP	testis expressed gene 261	1037650 IMAGE Consortium	AA638764
3566 C	4 021OAMNP	asparagine synthetase	1037772 IMAGE Consortium	AA638879
3567 C	6 021OAMNP	D-dopachrome tautomerase	1037884 IMAGE Consortium	AA638944
3568 C	8 021OAMNP	glyceronephosphate O-acyltransferase	1038284 IMAGE Consortium	AA671200
3569 C	10 021OAMNP	Mus musculus core1 UDP-galactose:N-acetylgalactosamine-alpha-R beta 1,3-galactosyltransferase (C1galt1) mRNA, complete cds	1038376 IMAGE Consortium	AA671409
3570 C	12 021OAMNP			
3571 E	2 021OAMNP	ESTs, Moderately similar to YCE7_HUMAN HYPOTHETICAL PROTEIN CGI-147 [H.sapiens]	1037658 IMAGE Consortium	AA638768
3572 E	4 021OAMNP	ESTs	1037777 IMAGE Consortium	AI642265
3573 E	6 021OAMNP	ESTs, Highly similar to activity-dependent neuroprotective protein [M.musculus]	1037917 IMAGE Consortium	AA636938
3574 E	8 021OAMNP	glycine C-acetyltransferase (2-amino-3-ketobutyrate-	1038289 IMAGE Consortium	AA671226

		coenzyme A ligase)		
3575 E	10 021OAMNP	ESTs	1038386 IMAGE Consortium	AA671422
3576 E	12 021OAMNP	DNA segment, Chr 14, ERATO Doi 310, expressed	1038479 IMAGE Consortium	AA624595
3577 G	2 021OAMNP	ESTs	1037661 IMAGE Consortium	AA638776
3578 G	4 021OAMNP	calcium binding protein P22	1037783 IMAGE Consortium	AI642266
3579 G	6 021OAMNP	ESTs, Weakly similar to AF220193_1 uncharacterized hypothalamus protein HT007 [H.sapiens]	1037920 IMAGE Consortium	AA636931
3580 G	8 021OAMNP			
3581 G	10 021OAMNP	cyclin B2	1038400 IMAGE Consortium	AA671428
3582 G	12 021OAMNP	ATP-binding cassette, sub-family F (GCN20), member 2	1038482 IMAGE Consortium	AA624596
3583 A	2 021VAMNQ	ESTs, Moderately similar to GEPHYRIN [Rattus norvegicus]	1038515 IMAGE Consortium	AA624733
3584 A	4 021VAMNQ	acetyl-Coenzyme A transporter	1109316 IMAGE Consortium	AA666972
3585 A	6 021VAMNQ	chromobox homolog 1 (Drosophila HP1 beta)	1109837 IMAGE Consortium	AA823935
3586 A	8 021VAMNQ	solute carrier family 25 (mitochondrial carrier; adenine nucleotide translocator), member 10	1111011 IMAGE Consortium	AA683883
3587 A	10 021VAMNQ	ESTs	1110716 IMAGE Consortium	AA683639
3588 A	12 021VAMNQ	ESTs	2101991 IMAGE Consortium	AW210270
3589 C	2 021VAMNQ	ESTs, Weakly similar to cDNA EST yk325c7.5 comes from this gene [C.elegans]	1038558 IMAGE Consortium	AA624872
3590 C	4 021VAMNQ	GATA-binding protein 3	1109348 IMAGE Consortium	AA666992
3591 C	6 021VAMNQ	RNA binding motif protein 6	1109869 IMAGE Consortium	AA823956
3592 C	8 021VAMNQ	ESTs, Moderately similar to 36 kDa FK506 binding protein [H.sapiens]	1110310 IMAGE Consortium	AA795608
3593 C	10 021VAMNQ	recombining binding protein suppressor of hairless (Drosophila)	1110845 IMAGE Consortium	AA683765
3594 C	12 021VAMNQ	ESTs	2102017 IMAGE Consortium	AW210301
3595 E	2 021VAMNQ	capping protein (actin filament), gelsolin-like	1038566 IMAGE Consortium	AA624875
3596 E	4 021VAMNQ	ESTs, Highly similar to AF125100_1 HSPC039 protein [H.sapiens]	1109381 IMAGE Consortium	AA667061
3597 E	6 021VAMNQ	capping protein beta 1	1109913 IMAGE Consortium	AA824021
3598 E	8 021VAMNQ	ESTs, Highly similar to cyclin D-interacting myb-like protein [M.musculus]	1111052 IMAGE Consortium	AA683898
3599 E	10 021VAMNQ	Eph receptor B3	1110951 IMAGE Consortium	AA683854
3600 E	12 021VAMNQ	fibrillin 1	2102050 IMAGE Consortium	AW210317
3601 G	2 021VAMNQ	ribosomal protein S16	1038589 IMAGE Consortium	AA624897
3602 G	4 021VAMNQ	cytochrome c oxidase, subunit Vb	1109616 IMAGE Consortium	AA822723
3603 G	6 021VAMNQ	serine/threonine kinase 2	1109990 IMAGE Consortium	AA795165
3604 G	8 021VAMNQ	ESTs	1111098 IMAGE Consortium	AA688749
3605 G	10 021VAMNQ	maternal embryonic message 3	2102229 IMAGE Consortium	AW209270
3606 G	12 021VAMNQ	tight junction protein 1	2102053 IMAGE Consortium	AW210329
3607 A	2 021AMNR	ESTs, Highly similar to N-ACETYLGLUCOSAMINE-6-SULFATASE PRECURSOR [Homo sapiens]	2102153 IMAGE Consortium	AW210416

3608 A	4 0212AMNR	cadherin 11	2101710 IMAGE Consortium	AW210019
3609 A	6 0212AMNR	ESTs, Weakly similar to CG6482 gene product [D.melanogaster]	2101209 IMAGE Consortium	AW209599
3610 A	8 0212AMNR	ESTs, Moderately similar to EPIDERMAL GROWTH FACTOR RECEPTOR SUBSTRATE SUBSTRATE 15 [Homo sapiens] {IMAGE:804453}	804453 IMAGE Consortium	AA474757
3611 A	10 0212AMNR	ESTs, Weakly similar to predicted using Genefinder [C.elegans] {IMAGE:945059}	945059 IMAGE Consortium	AA537323
3612 A	12 0212AMNR	IMAGE EST {IMAGE:775033}	775033 IMAGE Consortium	AA387356
3613 C	2 0212AMNR	peroxisomal delta3, delta2-enoyl-Coenzyme A isomerase	2102165 IMAGE Consortium	AW210430
3614 C	4 0212AMNR	vesicle-associated membrane protein 3	2101811 IMAGE Consortium	AW210120
3615 C	6 0212AMNR	heat shock factor 1	2101278 IMAGE Consortium	AW209653
3616 C	8 0212AMNR	Kinesin heavy chain member 5B, ubiquitous {IMAGE:760837}	760837 IMAGE Consortium	AA386680
3617 C	10 0212AMNR	ESTs {IMAGE:944942}	944942 IMAGE Consortium	AA537095
3618 C	12 0212AMNR	ESTs {IMAGE:804709}	804709 IMAGE Consortium	AA474386
3619 E	2 0212AMNR	protein kinase, DNA activated, catalytic polypeptide interacting protein	2102178 IMAGE Consortium	AW210425
3620 E	4 0212AMNR			
3621 E	6 0212AMNR	carnitine palmitoyltransferase 2	2101335 IMAGE Consortium	AW209706
3622 E	8 0212AMNR	ESTs, Highly similar to UDP-GLUCOSE 4-EPIMERASE [Rattus norvegicus] {IMAGE:760889}	760889 IMAGE Consortium	AA386807
3623 E	10 0212AMNR	ESTs {IMAGE:833601}	833601 IMAGE Consortium	AA432967
3624 E	12 0212AMNR	ESTs {IMAGE:760857}	760857 IMAGE Consortium	AA386550
3625 G	2 0212AMNR			
3626 G	4 0212AMNR			
3627 G	6 0212AMNR	ESTs, Weakly similar to GTP-binding protein [H.sapiens]	2101351 IMAGE Consortium	AW209723
3628 G	8 0212AMNR	ESTs {IMAGE:760847}	760847 IMAGE Consortium	AA386794
3629 G	10 0212AMNR	ESTs {IMAGE:833919}	833919 IMAGE Consortium	AA413174
3630 G	12 0212AMNR	ESTs, Highly similar to UTP--GLUCOSE-1-PHOSPHATE URIDYLTRANSFERASE [Homo sapiens] {IMAGE:805218}	805218 IMAGE Consortium	AA473123
3631 A	2 0219AMNS	ESTs {IMAGE:834291}	834291 IMAGE Consortium	AA413296
3632 A	4 0219AMNS			
3633 A	6 0219AMNS	ESTs	619105 IMAGE Consortium	AI592536
3634 A	8 0219AMNS	caspase 11	618560 IMAGE Consortium	AA175904
3635 A	10 0219AMNS	interleukin 6 receptor, alpha	619519 IMAGE Consortium	AA175680
3636 A	12 0219AMNS	ESTs	619787 IMAGE Consortium	AI604122
3637 C	2 0219AMNS	ESTs {IMAGE:834572}	834572 IMAGE Consortium	AA470303
3638 C	4 0219AMNS	IMAGE EST {IMAGE:775092}	775092 IMAGE Consortium	AA386832
3639 C	6 0219AMNS	ESTs	619125 IMAGE Consortium	AI592537
3640 C	8 0219AMNS	ESTs	619296 IMAGE Consortium	AA175733
3641 C	10 0219AMNS	ESTs	619615 IMAGE Consortium	AA175783

3642 C	12 0219AMNS	ESTs	619829 IMAGE Consortium	AI604133
3643 E	2 0219AMNS	ESTs, Moderately similar to JAK3 [M.musculus] {IMAGE:834549}	834549 IMAGE Consortium	AA470301
3644 E	4 0219AMNS	IMAGE EST {IMAGE:774808}	774808 IMAGE Consortium	AA386709
3645 E	6 0219AMNS	ESTs	619161 IMAGE Consortium	AA175641
3646 E	8 0219AMNS	ESTs	619301 IMAGE Consortium	AA175738
3647 E	10 0219AMNS	ESTs	618901 IMAGE Consortium	AA175461
3648 E	12 0219AMNS	ESTs	619906 IMAGE Consortium	AA177995
3649 G	2 0219AMNS	IMAGE EST {IMAGE:524133}	524133 IMAGE Consortium	AA066133
3650 G	4 0219AMNS	ESTs {IMAGE:775538}	775538 IMAGE Consortium	AA388304
3651 G	6 0219AMNS	fucosyltransferase 7	619197 IMAGE Consortium	AA170772
3652 G	8 0219AMNS	ESTs, Highly similar to AA2A_MOUSE ADENOSINE A2A RECEPTOR [M.musculus]	618613 IMAGE Consortium	AA174942
3653 G	10 0219AMNS	ESTs	619639 IMAGE Consortium	AI604096
3654 G	12 0219AMNS	gene trap ROSA 26, Philippe Soriano	619948 IMAGE Consortium	AA178020
3655 A	2 021GAMNT	ESTs {IMAGE:524641}	524641 IMAGE Consortium	AA068605
3656 A	4 021GAMNT			
3657 A	6 021GAMNT	ESTs, Moderately similar to GLYPICAN-3 PRECURSOR [Rattus norvegicus] {IMAGE:657264}	657264 IMAGE Consortium	AA240923
3658 A	8 021GAMNT	ESTs, Moderately similar to HYPOTHETICAL 49.7 KD PROTEIN IN GIN2-STE3 INTERGENIC REGION [Saccharomyces cerevisiae] {IMAGE:779257}	779257 IMAGE Consortium	AA445775
3659 A	10 021GAMNT	IMAGE EST {IMAGE:315913}	315913 IMAGE Consortium	W08983
3660 A	12 021GAMNT	ESTs {IMAGE:318938}	318938 IMAGE Consortium	W34387
3661 C	2 021GAMNT	IMAGE EST {IMAGE:533085}	533085 IMAGE Consortium	AA068665
3662 C	4 021GAMNT	Mus musculus PAF acetylhydrolase mRNA, complete cds {IMAGE:536464}	536464 IMAGE Consortium	AA073963
3663 C	6 021GAMNT	ESTs, Highly similar to ETn insert [M.musculus] {IMAGE:633484}	633484 IMAGE Consortium	AA184421
3664 C	8 021GAMNT	ESTs, Highly similar to N33 protein form 2 [H.sapiens] {IMAGE:779020}	779020 IMAGE Consortium	AA413956
3665 C	10 021GAMNT	IMAGE EST {IMAGE:316143}	316143 IMAGE Consortium	W12536
3666 C	12 021GAMNT	ESTs {IMAGE:333486}	333486 IMAGE Consortium	AA052356
3667 E	2 021GAMNT	ESTs {IMAGE:523046}	523046 IMAGE Consortium	AA066439
3668 E	4 021GAMNT	ESTs, Highly similar to HYPOTHETICAL 36.7 KD PROTEIN AH6.2 IN CHROMOSOME II [Caenorhabditis elegans] {IMAGE:532139}	532139 IMAGE Consortium	AA068366
3669 E	6 021GAMNT	Cytochrome P450, 2e1, ethanol inducible {IMAGE:656925}	656925 IMAGE Consortium	AA240739
3670 E	8 021GAMNT	Mus musculus calmodulin-binding protein SHA1 (Sha1) mRNA, complete cds {IMAGE:791460}	791460 IMAGE Consortium	AA416426
3671 E	10 021GAMNT	ESTs {IMAGE:313629}	313629 IMAGE Consortium	W10658
3672 E	12 021GAMNT	Mus musculus Bet1p homolog (mbet1) mRNA, complete cds {IMAGE:333232}	333232 IMAGE Consortium	W18376

3673 G	2 021GAMNT	IMAGE EST {IMAGE:533489}	533489 IMAGE Consortium	AA105093
3674 G	4 021GAMNT	IMAGE EST {IMAGE:523726}	523726 IMAGE Consortium	AA067346
3675 G	6 021GAMNT	IMAGE EST {IMAGE:651304}	651304 IMAGE Consortium	AA214855
3676 G	8 021GAMNT	ESTs, Weakly similar to predicted using Genefinder [C.elegans] {IMAGE:793605}	793605 IMAGE Consortium	AA415799
3677 G	10 021GAMNT	ESTs {IMAGE:317515}	317515 IMAGE Consortium	W34080
3678 G	12 021GAMNT	ESTs, Weakly similar to Similarity to Sheep transthyretin precursor [C.elegans] {IMAGE:331792}	331792 IMAGE Consortium	W08097
3679 A	2 021NAMNU	Mus musculus Eph-and Elk-related kinase (eek) mRNA, complete cds {IMAGE:352644}	352644 IMAGE Consortium	W34187
3680 A	4 021NAMNU	IMAGE EST {IMAGE:420020}	420020 IMAGE Consortium	W90879
3681 A	6 021NAMNU			
3682 A	8 021NAMNU	ACETYLCHOLINE RECEPTOR PROTEIN, GAMMA CHAIN PRECURSOR {IMAGE:425717}	425717 IMAGE Consortium	AA000299
3683 A	10 021NAMNU	IMAGE EST {IMAGE:439436}	439436 IMAGE Consortium	AA004142
3684 A	12 021NAMNU	ESTs {IMAGE:457055}	457055 IMAGE Consortium	AA023720
3685 C	2 021NAMNU	M.musculus mRNA for N-glycan alpha 2,8-sialyltransferase {IMAGE:355197}	355197 IMAGE Consortium	W47914
3686 C	4 021NAMNU	ESTs {IMAGE:418690}	418690 IMAGE Consortium	W96925
3687 C	6 021NAMNU	ESTs {IMAGE:423473}	423473 IMAGE Consortium	W98815
3688 C	8 021NAMNU	ESTs {IMAGE:426172}	426172 IMAGE Consortium	AA002994
3689 C	10 021NAMNU	Homeo box B6 {IMAGE:439758}	439758 IMAGE Consortium	AA008326
3690 C	12 021NAMNU	ESTs, Weakly similar to brain carcinoembryonic antigen [M.musculus] {IMAGE:456237}	456237 IMAGE Consortium	AA023268
3691 E	2 021NAMNU	ESTs, Moderately similar to MANNOSE-1-PHOSPHATE GUANYLTRANSFERASE [Saccharomyces cerevisiae] {IMAGE:373922}	373922 IMAGE Consortium	W59400
3692 E	4 021NAMNU	IMAGE EST {IMAGE:420468}	420468 IMAGE Consortium	W90847
3693 E	6 021NAMNU	Homeo box B9 {IMAGE:423995}	423995 IMAGE Consortium	W91526
3694 E	8 021NAMNU	IMAGE EST {IMAGE:425482}	425482 IMAGE Consortium	AA000044
3695 E	10 021NAMNU	ESTs {IMAGE:437601}	437601 IMAGE Consortium	AA003728
3696 E	12 021NAMNU	ESTs {IMAGE:455371}	455371 IMAGE Consortium	AA020415
3697 G	2 021NAMNU			
3698 G	4 021NAMNU	IMAGE EST {IMAGE:421965}	421965 IMAGE Consortium	W97408
3699 G	6 021NAMNU	ESTs {IMAGE:424563}	424563 IMAGE Consortium	W98594
3700 G	8 021NAMNU	ESTs {IMAGE:427218}	427218 IMAGE Consortium	AA003081
3701 G	10 021NAMNU	ESTs {IMAGE:439902}	439902 IMAGE Consortium	AA008727
3702 G	12 021NAMNU	ESTs {IMAGE:458513}	458513 IMAGE Consortium	AA023949
3703 A	2 021UAMNV	ESTs, Highly similar to TRYPSINOGEN [Bos taurus] {IMAGE:464085}	464085 IMAGE Consortium	AA028356
3704 A	4 021UAMNV	Kruppel-like factor 3 (basic)	622046 IMAGE Consortium	AI591871
3705 A	6 021UAMNV	ESTs	621685 IMAGE Consortium	AI591831
3706 A	8 021UAMNV	ESTs	622927 IMAGE Consortium	AI591964

3707 A	10 021UAMNV	ESTs {IMAGE:476369}	476369 IMAGE Consortium	AA050212
3708 A	12 021UAMNV	Tumor necrosis factor induced protein 6 {IMAGE:478579}	478579 IMAGE Consortium	AA051341
3709 C	2 021UAMNV	IMAGE EST {IMAGE:464498}	464498 IMAGE Consortium	AA033029
3710 C	4 021UAMNV	ESTs	622242 IMAGE Consortium	AI591887
3711 C	6 021UAMNV	ESTs	621720 IMAGE Consortium	AI591833
3712 C	8 021UAMNV	ESTs, Moderately similar to KIAA0853 protein [H.sapiens]	634126 IMAGE Consortium	AA184651
3713 C	10 021UAMNV	ESTs {IMAGE:475631}	475631 IMAGE Consortium	AA049807
3714 C	12 021UAMNV	IMAGE EST {IMAGE:480219}	480219 IMAGE Consortium	AA058084
3715 E	2 021UAMNV	IMAGE EST {IMAGE:466905}	466905 IMAGE Consortium	AA031230
3716 E	4 021UAMNV	ESTs	622256 IMAGE Consortium	AI591889
3717 E	6 021UAMNV	ESTs	622507 IMAGE Consortium	AI591911
3718 E	8 021UAMNV	adrenergic receptor, beta 2	634188 IMAGE Consortium	AA182071
3719 E	10 021UAMNV	ESTs {IMAGE:476972}	476972 IMAGE Consortium	AA048101
3720 E	12 021UAMNV	IMAGE EST {IMAGE:480181}	480181 IMAGE Consortium	AA058187
3721 G	2 021UAMNV	IMAGE EST {IMAGE:472801}	472801 IMAGE Consortium	AA037995
3722 G	4 021UAMNV	ESTs	621548 IMAGE Consortium	AI592339
3723 G	6 021UAMNV	EST	621827 IMAGE Consortium	AI451147
3724 G	8 021UAMNV	ESTs	634362 IMAGE Consortium	AA185219
3725 G	10 021UAMNV	Mus musculus mRNA for p67phox, complete cds {IMAGE:475840}	475840 IMAGE Consortium	AA050149
3726 G	12 021UAMNV	ESTs {IMAGE:481481}	481481 IMAGE Consortium	AA060306
3727 A	2 0211AMNW	ESTs {IMAGE:483578}	483578 IMAGE Consortium	AA061744
3728 A	4 0211AMNW	Lipocalin 2 {IMAGE:493658}	493658 IMAGE Consortium	AA087193
3729 A	6 0211AMNW	IMAGE EST {IMAGE:573581}	573581 IMAGE Consortium	AA119375
3730 A	8 0211AMNW	ESTs, Weakly similar to cDNA EST EMBL:D70402 comes from this gene [C.elegans] {IMAGE:576761}	576761 IMAGE Consortium	AA120013
3731 A	10 0211AMNW	ESTs {IMAGE:583033}	583033 IMAGE Consortium	AA153663
3732 A	12 0211AMNW	ESTs {IMAGE:597280}	597280 IMAGE Consortium	AA138290
3733 C	2 0211AMNW	IMAGE EST {IMAGE:482886}	482886 IMAGE Consortium	AA062144
3734 C	4 0211AMNW	ESTs {IMAGE:493434}	493434 IMAGE Consortium	AA086952
3735 C	6 0211AMNW	Hepatic lipase {IMAGE:576579}	576579 IMAGE Consortium	AA124071
3736 C	8 0211AMNW	ESTs {IMAGE:576131}	576131 IMAGE Consortium	AA119797
3737 C	10 0211AMNW	Mus musculus Ets-family transcription factor Elf-1 mRNA, complete cds {IMAGE:581665}	581665 IMAGE Consortium	AA152626
3738 C	12 0211AMNW	ESTs, Weakly similar to cDNA EST EMBL:C13590 comes from this gene [C.elegans] {IMAGE:596404}	596404 IMAGE Consortium	AA138546
3739 E	2 0211AMNW	ESTs {IMAGE:483572}	483572 IMAGE Consortium	AA061650
3740 E	4 0211AMNW	ESTs {IMAGE:493652}	493652 IMAGE Consortium	AA098345
3741 E	6 0211AMNW	ESTs {IMAGE:574527}	574527 IMAGE Consortium	AA125185
3742 E	8 0211AMNW	ESTs {IMAGE:576767}	576767 IMAGE Consortium	AA120015
3743 E	10 0211AMNW	ESTs {IMAGE:577555}	577555 IMAGE Consortium	AA138025
3744 E	12 0211AMNW	IMAGE EST {IMAGE:598824}	598824 IMAGE Consortium	AA168416
3745 G	2 0211AMNW	IMAGE EST {IMAGE:481307}	481307 IMAGE Consortium	AA060448

3746 G	4 0211AMNW	ESTs {IMAGE:573243}	573243 IMAGE Consortium	AA119313
3747 G	6 0211AMNW	Tax-transcriptionally activated glycoprotein 1 {IMAGE:573690}	573690 IMAGE Consortium	AA110765
3748 G	8 0211AMNW	IMAGE EST {IMAGE:577135}	577135 IMAGE Consortium	AA124522
3749 G	10 0211AMNW	IMAGE EST {IMAGE:578097}	578097 IMAGE Consortium	AA124744
3750 G	12 0211AMNW	IG ALPHA CHAIN C REGION {IMAGE:596604}	596604 IMAGE Consortium	AA146478
3751 A	2 0218AMNX	Mus musculus PS-2 mRNA, complete cds {IMAGE:597527}	597527 IMAGE Consortium	AA144382
3752 A	4 0218AMNX	ESTs {IMAGE:619632}	619632 IMAGE Consortium	AA175784
3753 A	6 0218AMNX	Small inducible cytokine A4 {IMAGE:621095}	621095 IMAGE Consortium	AA178155
3754 A	8 0218AMNX	ESTs {IMAGE:619696}	619696 IMAGE Consortium	AA172813
3755 A	10 0218AMNX	ESTs {IMAGE:622722}	622722 IMAGE Consortium	AA177554
3756 A	12 0218AMNX	ESTs, Weakly similar to EGF repeat transmembrane protein [M.musculus] {IMAGE:636129}	636129 IMAGE Consortium	AA189462
3757 C	2 0218AMNX	ESTs {IMAGE:598060}	598060 IMAGE Consortium	AA265239
3758 C	4 0218AMNX	ESTs {IMAGE:617294}	617294 IMAGE Consortium	AA174315
3759 C	6 0218AMNX	ESTs {IMAGE:617197}	617197 IMAGE Consortium	AA174776
3760 C	8 0218AMNX	IMAGE EST {IMAGE:617992}	617992 IMAGE Consortium	AA170335
3761 C	10 0218AMNX	Interleukin 4 receptor, alpha {IMAGE:635530}	635530 IMAGE Consortium	AA166418
3762 C	12 0218AMNX	ESTs, Weakly similar to ZINC FINGER PROTEIN 91 [H.sapiens] {IMAGE:622878}	622878 IMAGE Consortium	AA184296
3763 E	2 0218AMNX	ESTs {IMAGE:597336}	597336 IMAGE Consortium	AA146020
3764 E	4 0218AMNX	ESTs {IMAGE:620766}	620766 IMAGE Consortium	AA178217
3765 E	6 0218AMNX	ESTs {IMAGE:617010}	617010 IMAGE Consortium	AA163405
3766 E	8 0218AMNX	ESTs {IMAGE:621412}	621412 IMAGE Consortium	AA178467
3767 E	10 0218AMNX	IMAGE EST {IMAGE:622653}	622653 IMAGE Consortium	AA177538
3768 E	12 0218AMNX	IMAGE EST {IMAGE:622128}	622128 IMAGE Consortium	AA184039
3769 G	2 0218AMNX	ESTs {IMAGE:597201}	597201 IMAGE Consortium	AA145326
3770 G	4 0218AMNX	ESTs {IMAGE:618584}	618584 IMAGE Consortium	AA175911
3771 G	6 0218AMNX	IMAGE EST {IMAGE:620086}	620086 IMAGE Consortium	AA172493
3772 G	8 0218AMNX	Janus kinase 2 {IMAGE:621226}	621226 IMAGE Consortium	AA178495
3773 G	10 0218AMNX	ESTs {IMAGE:622920}	622920 IMAGE Consortium	AA184365
3774 G	12 0218AMNX	ESTs {IMAGE:634992}	634992 IMAGE Consortium	AA154838
3775 A	2 021FAMNY	IMAGE EST {IMAGE:622914}	622914 IMAGE Consortium	AA177703
3776 A	4 021FAMNY	ESTs, Highly similar to HYPOTHETICAL PROTEIN KIAA0036 [H.sapiens] {IMAGE:640287}	640287 IMAGE Consortium	AA198784
3777 A	6 021FAMNY	IMAGE EST {IMAGE:638739}	638739 IMAGE Consortium	AA200942
3778 A	8 021FAMNY	IMAGE EST {IMAGE:653484}	653484 IMAGE Consortium	AA215268
3779 A	10 021FAMNY	ESTs {IMAGE:653402}	653402 IMAGE Consortium	AA240299
3780 A	12 021FAMNY	ESTs	635256 IMAGE Consortium	AA189494
3781 C	2 021FAMNY	Transcription factor 21 {IMAGE:634816}	634816 IMAGE Consortium	AA154683
3782 C	4 021FAMNY			
3783 C	6 021FAMNY	ESTs {IMAGE:640919}	640919 IMAGE Consortium	AA204360

3784 C	8 021FAMNY	ESTs {IMAGE:641687}	641687 IMAGE Consortium	AA209990
3785 C	10 021FAMNY	Mus musculus CTP synthetase mRNA, complete cds {IMAGE:642329}	642329 IMAGE Consortium	AA185252
3786 C	12 021FAMNY	ESTs	635274 IMAGE Consortium	AA189770
3787 E	2 021FAMNY	ESTs {IMAGE:636375}	636375 IMAGE Consortium	AA183022
3788 E	4 021FAMNY	ESTs {IMAGE:638401}	638401 IMAGE Consortium	AA217009
3789 E	6 021FAMNY	IMAGE EST {IMAGE:641462}	641462 IMAGE Consortium	AA200457
3790 E	8 021FAMNY	ESTs {IMAGE:652544}	652544 IMAGE Consortium	AA217333
3791 E	10 021FAMNY	DNA segment, Chr 7, ERATO Doi 758, expressed	635009 IMAGE Consortium	AA155495
3792 E	12 021FAMNY	Rad51 homolog ( <i>S. cerevisiae</i> )	635294 IMAGE Consortium	AI592821
3793 G	2 021FAMNY	ESTs {IMAGE:636308}	636308 IMAGE Consortium	AA182050
3794 G	4 021FAMNY	ESTs {IMAGE:640611}	640611 IMAGE Consortium	AA203985
3795 G	6 021FAMNY	IMAGE EST {IMAGE:639062}	639062 IMAGE Consortium	AA199299
3796 G	8 021FAMNY	ESTs {IMAGE:643257}	643257 IMAGE Consortium	AA186075
3797 G	10 021FAMNY	Mus musculus inducible 6-phosphofructo-2-kinase mRNA, complete cds	635021 IMAGE Consortium	AA163244
3798 G	12 021FAMNY	ESTs	636036 IMAGE Consortium	AI645130
3799 A	2 021MAMNZ	transducer of ErbB-2.1	635510 IMAGE Consortium	AA166386
3800 A	4 021MAMNZ	ESTs	634993 IMAGE Consortium	AI592788
3801 A	6 021MAMNZ	peroxisome biogenesis factor 7	316746 IMAGE Consortium	W11104
3802 A	8 021MAMNZ	matrix metalloproteinase 9	331802 IMAGE Consortium	W14436
3803 A	10 021MAMNZ	ESTs, Weakly similar to RHO1 PROTEIN [Schizosaccharomyces pombe]	335280 IMAGE Consortium	W17640
3804 A	12 021MAMNZ	Epstein-Barr virus induced gene 3	385689 IMAGE Consortium	W62918
3805 C	2 021MAMNZ	ESTs	635580 IMAGE Consortium	AI509252
3806 C	4 021MAMNZ	ESTs	635735 IMAGE Consortium	AA163495
3807 C	6 021MAMNZ	natriuretic peptide receptor 1	317611 IMAGE Consortium	W13098
3808 C	8 021MAMNZ	endothelial differentiation sphingolipid G-protein-coupled receptor 1	332169 IMAGE Consortium	W08198
3809 C	10 021MAMNZ	neural cell adhesion molecule	347892 IMAGE Consortium	W29590
3810 C	12 021MAMNZ	forkhead box G1	388688 IMAGE Consortium	AI893944
3811 E	2 021MAMNZ	ESTs	634893 IMAGE Consortium	AA154781
3812 E	4 021MAMNZ	ESTs	635775 IMAGE Consortium	AI509287
3813 E	6 021MAMNZ	ephrin A4	317658 IMAGE Consortium	W12777
3814 E	8 021MAMNZ	protein tyrosine phosphatase, non-receptor type 11	332650 IMAGE Consortium	W08730
3815 E	10 021MAMNZ	kallikrein 26	350112 IMAGE Consortium	W34620
3816 E	12 021MAMNZ	glutamate receptor, ionotropic, NMDA1 (zeta 1)	400025 IMAGE Consortium	AI893952
3817 G	2 021MAMNZ	ESTs	635611 IMAGE Consortium	AA163085
3818 G	4 021MAMNZ	ESTs, Weakly similar to APC-binding protein EB1 homolog [M.musculus]	635856 IMAGE Consortium	AA185325
3819 G	6 021MAMNZ	transglutaminase 3, E polypeptide	317794 IMAGE Consortium	AI325332
3820 G	8 021MAMNZ	insulin-like growth factor binding protein 6	332940 IMAGE Consortium	W15804
3821 G	10 021MAMNZ	phosphatidylinositol membrane-associated	352852 IMAGE Consortium	W34690



3822 G	12 021MAMNZ	nuclear factor of activated T-cells, cytoplasmic 1	401579 IMAGE Consortium	W78454
3823 A	2 021MAMOO	myelin-associated glycoprotein	391791 IMAGE Consortium	AI893951
3824 A	4 021MAMOO	interleukin 3 receptor, alpha chain	445664 IMAGE Consortium	AI894066
3825 A	6 021MAMOO	transcription elongation factor A (SII), 3	423026 IMAGE Consortium	W97993
3826 A	8 021MAMOO	CD59a antigen	578532 IMAGE Consortium	AA162378
3827 A	10 021MAMOO	nuclear factor of activated T-cells, cytoplasmic 2	620068 IMAGE Consortium	AA176009
3828 A	12 021MAMOO	discs, large homolog 4 (Drosophila)	639235 IMAGE Consortium	AA200993
3829 C	2 021MAMOO	hairy and enhancer of split 5, (Drosophila)	400761 IMAGE Consortium	AI893957
3830 C	4 021MAMOO	aortic preferentially expressed gene 1	457583 IMAGE Consortium	AI894082
3831 C	6 021MAMOO	epsin 1	441278 IMAGE Consortium	AA011791
3832 C	8 021MAMOO	linker for activation of T cells	582840 IMAGE Consortium	AA153591
3833 C	10 021MAMOO	wingless-related MMTV integration site 11	621780 IMAGE Consortium	AA183414
3834 C	12 021MAMOO	dihydrofolate reductase	642271 IMAGE Consortium	AA212405
3835 E	2 021MAMOO	urokinase plasminogen activator receptor	404131 IMAGE Consortium	W82324
3836 E	4 021MAMOO	forkhead box A3	476343 IMAGE Consortium	AA050454
3837 E	6 021MAMOO	adenosine deaminase	456324 IMAGE Consortium	AA023331
3838 E	8 021MAMOO	B-cell leukemia/lymphoma 3	595882 IMAGE Consortium	AA138064
3839 E	10 021MAMOO	programmed cell death 4	622890 IMAGE Consortium	AA177698
3840 E	12 021MAMOO	corticotropin releasing hormone receptor	643358 IMAGE Consortium	AA189886
3841 G	2 021MAMOO	mitogen activated protein kinase 10	419942 IMAGE Consortium	AI893985
3842 G	4 021MAMOO	tumor necrosis factor receptor superfamily, member 5	477641 IMAGE Consortium	AA048574
3843 G	6 021MAMOO	decidual/trophoblast prolactin-related protein	458442 IMAGE Consortium	AA023998
3844 G	8 021MAMOO	CD19 antigen	596887 IMAGE Consortium	AA144880
3845 G	10 021MAMOO	cytotoxic T lymphocyte-associated protein 2 beta	635845 IMAGE Consortium	AA182308
3846 G	12 021MAMOO	Eph receptor A3	658197 IMAGE Consortium	AA198303
3847 A	2 021TAMO1	complement component 9	680492 IMAGE Consortium	AA237324
3848 A	4 021TAMO1	complement receptor 2	722309 IMAGE Consortium	AA254235
3849 A	6 021TAMO1	ESTs, Highly similar to HGFL_MOUSE HEPATOCYTE GROWTH FACTOR-LIKE PROTEIN PRECURSOR [M.musculus]	692292 IMAGE Consortium	AA238356
3850 A	8 021TAMO1	cytokine receptor family, class 1 (WSXWS), member 1	752212 IMAGE Consortium	AI508564
3851 A	10 021TAMO1	anterior gradient 2 (Xenopus laevis)	351305 IMAGE Consortium	W40608
3852 A	12 021TAMO1	sema domain, immunoglobulin domain (Ig), short basic domain, secreted, (semaphorin) 3E	670962 IMAGE Consortium	AA241390
3853 C	2 021TAMO1	surfeit gene 6	680911 IMAGE Consortium	AA241888
3854 C	4 021TAMO1	synaptotagmin 2	334490 IMAGE Consortium	AA052402
3855 C	6 021TAMO1	sulfotransferase, hydroxysteroid preferring 2	693650 IMAGE Consortium	AA266661
3856 C	8 021TAMO1	Mus musculus HMG-box transcription factor TCF4B (Tcf4) mRNA, complete cds	764951 IMAGE Consortium	AA273416
3857 C	10 021TAMO1	N-acetyl galactosaminidase, alpha	438635 IMAGE Consortium	AA050869
3858 C	12 021TAMO1	carbonic anhydrase-like sequence 1	720107 IMAGE Consortium	AA254942
3859 E	2 021TAMO1	interleukin 18	681648 IMAGE Consortium	AA237736
3860 E	4 021TAMO1	aldo-keto reductase	353040 IMAGE Consortium	W35987

3861 E	6 021TAMO1	mature T-cell proliferation 1	723189 IMAGE Consortium	AA254474
3862 E	8 021TAMO1	LIM homeobox protein 8	777339 IMAGE Consortium	AA276924
3863 E	10 021TAMO1	SRY-box containing gene 17	442058 IMAGE Consortium	AA013556
3864 E	12 021TAMO1	Rho interacting protein 2	776054 IMAGE Consortium	AA276026
3865 G	2 021TAMO1	retinoic acid early transcript delta	681805 IMAGE Consortium	AA244388
3866 G	4 021TAMO1	7-dehydrocholesterol reductase	367115 IMAGE Consortium	W54045
3867 G	6 021TAMO1	reelin	734262 IMAGE Consortium	AA268883
3868 G	8 021TAMO1	Friend leukemia integration 1	808955 IMAGE Consortium	AA467723
3869 G	10 021TAMO1	gamma-aminobutyric acid (GABA-A) transporter 2	445665 IMAGE Consortium	AA017866
3870 G	12 021TAMO1	vesicle-associated membrane protein 4	639685 IMAGE Consortium	AI785208
3871 A	2 021AMO2	ESTs, Weakly similar to S06318 endoplasmic reticulum protein ERp72 homolog - mouse [M.musculus]	639261 IMAGE Consortium	AI645950
3872 A	4 021AMO2	ESTs, Weakly similar to AF064806_1 vinexin alpha [M.musculus]	639683 IMAGE Consortium	AA197390
3873 A	6 021AMO2	interleukin 18 receptor accessory protein	640615 IMAGE Consortium	AA203986
3874 A	8 021AMO2	caspase 6	636228 IMAGE Consortium	AA183198
3875 A	10 021AMO2	ESTs, Weakly similar to endocrine regulator [H.sapiens]	636449 IMAGE Consortium	AA182058
3876 A	12 021AMO2	ESTs	636618 IMAGE Consortium	AI645199
3877 C	2 021AMO2	ESTs, Moderately similar to T08762 hypothetical protein DKFZp586P2219.1 - human [H.sapiens]	637810 IMAGE Consortium	AA189642
3878 C	4 021AMO2	ESTs	639793 IMAGE Consortium	AA197555
3879 C	6 021AMO2	ESTs	640718 IMAGE Consortium	AA203962
3880 C	8 021AMO2	ESTs	636258 IMAGE Consortium	AI645157
3881 C	10 021AMO2	erythrocyte protein band 4.1-like 3	636468 IMAGE Consortium	AA182223
3882 C	12 021AMO2	ESTs, Weakly similar to JAK3_MOUSE TYROSINE-PROTEIN KINASE JAK3 [M.musculus]	636648 IMAGE Consortium	AI646621
3883 E	2 021AMO2	mitogen activated protein kinase kinase 6	639277 IMAGE Consortium	AA199056
3884 E	4 021AMO2	immune associated nucleotide 3	639935 IMAGE Consortium	AA197670
3885 E	6 021AMO2	ESTs, Highly similar to DNA-REPAIR PROTEIN COMPLEMENTING XP-B CELLS HOMOLOG [Mus musculus]	638093 IMAGE Consortium	AI509373
3886 E	8 021AMO2	serine protease inhibitor 6	636322 IMAGE Consortium	AA189231
3887 E	10 021AMO2	ESTs, Weakly similar to ANX7_MOUSE ANNEXIN VII [M.musculus]	637211 IMAGE Consortium	AI645264
3888 E	12 021AMO2	ESTs	636662 IMAGE Consortium	AA189475
3889 G	2 021AMO2	ESTs	638580 IMAGE Consortium	AA200183
3890 G	4 021AMO2	ESTs	640074 IMAGE Consortium	AA197420
3891 G	6 021AMO2	ATPase, H <sup>+</sup> transporting, lysosomal (vacuolar proton pump), subunit 1	636755 IMAGE Consortium	AA183318
3892 G	8 021AMO2	ESTs	636331 IMAGE Consortium	AI645164
3893 G	10 021AMO2	ESTs, Highly similar to SNAG_HUMAN GAMMA-SOLUBLE NSF ATTACHMENT PROTEIN [H.sapiens]	637222 IMAGE Consortium	AI645263
3894 G	12 021AMO2	ESTs	636698 IMAGE Consortium	AA189266

3895 A	2 0217AMO3	dead ringer homolog 1 (Drosophila)	642414 IMAGE Consortium	AA184881
3896 A	4 0217AMO3	ethanol decreased 2	642611 IMAGE Consortium	AA184893
3897 A	6 0217AMO3	s17 protein	347837 IMAGE Consortium	AI391019
3898 A	8 0217AMO3	ESTs	643118 IMAGE Consortium	AA189921
3899 A	10 0217AMO3	complement receptor 2	644166 IMAGE Consortium	AA208784
3900 A	12 0217AMO3	ESTs	642886 IMAGE Consortium	AI627066
3901 C	2 0217AMO3	ESTs	642432 IMAGE Consortium	AI606016
3902 C	4 0217AMO3	potassium intermediate/small conductance calcium-activated channel, subfamily N, member 4	642677 IMAGE Consortium	AA185547
3903 C	6 0217AMO3	ESTs	336003 IMAGE Consortium	AI390160
3904 C	8 0217AMO3	ESTs	642427 IMAGE Consortium	AI606015
3905 C	10 0217AMO3	protein tyrosine phosphatase, receptor type, C polypeptide-associated protein	642720 IMAGE Consortium	AI627045
3906 C	12 0217AMO3	ESTs	643603 IMAGE Consortium	AA200306
3907 E	2 0217AMO3	ESTs	642459 IMAGE Consortium	AA184852
3908 E	4 0217AMO3	ESTs	338012 IMAGE Consortium	W29714
3909 E	6 0217AMO3	ESTs, Weakly similar to 2022314A granule cell marker protein [M.musculus]	336054 IMAGE Consortium	AI390162
3910 E	8 0217AMO3	ESTs	643245 IMAGE Consortium	AI627099
3911 E	10 0217AMO3	ESTs	643479 IMAGE Consortium	AI390453
3912 E	12 0217AMO3	ESTs	643612 IMAGE Consortium	AI390461
3913 G	2 0217AMO3	ESTs	642492 IMAGE Consortium	AA190127
3914 G	4 0217AMO3	cartilage oligomeric matrix protein	335946 IMAGE Consortium	AI390155
3915 G	6 0217AMO3	ESTs, Highly similar to 2-OXOGLUTARATE DEHYDROGENASE E1 COMPONENT PRECURSOR [Homo sapiens]	336183 IMAGE Consortium	AI322289
3916 G	8 0217AMO3	ESTs	643250 IMAGE Consortium	AI627101
3917 G	10 0217AMO3	ESTs	642769 IMAGE Consortium	AI627052
3918 G	12 0217AMO3	ESTs	644359 IMAGE Consortium	AA265229
3919 A	2 021EAMO4	T-cell acute lymphocytic leukemia 1	643709 IMAGE Consortium	AI390473
3920 A	4 021EAMO4	four jointed box 1 (Drosophila)	658549 IMAGE Consortium	AA221832
3921 A	6 021EAMO4	ESTs	659701 IMAGE Consortium	AA220032
3922 A	8 021EAMO4	Mus musculus transcription factor-like protein ODA-10 mRNA, complete cds	669809 IMAGE Consortium	AA230638
3923 A	10 021EAMO4	ESTs	652719 IMAGE Consortium	AI390606
3924 A	12 021EAMO4	ESTs	658214 IMAGE Consortium	AI390388
3925 C	2 021EAMO4	ESTs	643739 IMAGE Consortium	AI390478
3926 C	4 021EAMO4	ESTs	659359 IMAGE Consortium	AA217921
3927 C	6 021EAMO4	ESTs	659712 IMAGE Consortium	AI390197
3928 C	8 021EAMO4	myosin, heavy polypeptide 3, skeletal muscle, embryonic	669856 IMAGE Consortium	AA230591
3929 C	10 021EAMO4	ESTs	652760 IMAGE Consortium	AI390609
3930 C	12 021EAMO4	EST	653004 IMAGE Consortium	AI465532
3931 E	2 021EAMO4	Mus musculus Ets transcription factor Spi-B, partial cds	643831 IMAGE Consortium	AI450417

3932 E	4 021EAMO4	ESTs, Weakly similar to F53B1.2 [C.elegans]	659372 IMAGE Consortium	AA218404
3933 E	6 021EAMO4	ESTs	659865 IMAGE Consortium	AA219894
3934 E	8 021EAMO4	ESTs, Weakly similar to AF112151_1 developmental arteries and neural crest EGF-like protein [M.musculus]	660500 IMAGE Consortium	AA222756
3935 E	10 021EAMO4	ESTs	652774 IMAGE Consortium	AI390650
3936 E	12 021EAMO4	Mouse mRNA for neurotrophin receptor interacting factor (Zfp110 gene)	653050 IMAGE Consortium	AI390684
3937 G	2 021EAMO4	ESTs	643859 IMAGE Consortium	AI450423
3938 G	4 021EAMO4	ESTs	658768 IMAGE Consortium	AI390423
3939 G	6 021EAMO4	ESTs	659994 IMAGE Consortium	AI464317
3940 G	8 021EAMO4	ESTs	660571 IMAGE Consortium	AI390230
3941 G	10 021EAMO4	Mus musculus brain cDNA, clone MNCb-0467, similar to Homo sapiens cDNA FLJ10078, clone HEMBA1001869	652832 IMAGE Consortium	AI390656
3942 G	12 021EAMO4	leucine rich repeat protein 1, neuronal	653102 IMAGE Consortium	AA216890
3943 A	2 021LAMO5	ESTs	653341 IMAGE Consortium	AI390714
3944 A	4 021LAMO5	ESTs, Highly similar to calcium-independent phospholipase A2 [H.sapiens]	679779 IMAGE Consortium	AI451806
3945 A	6 021LAMO5	zinc finger protein, multitype 2	672455 IMAGE Consortium	AA231039
3946 A	8 021LAMO5	immunity-associated protein, 38 kDa	678649 IMAGE Consortium	AA238037
3947 A	10 021LAMO5	ESTs	681077 IMAGE Consortium	AA242653
3948 A	12 021LAMO5	ESTs, Weakly similar to CBPC_MOUSE MAST CELL CARBOXYPEPTIDASE A PRECURSOR [M.musculus]	680881 IMAGE Consortium	AI451862
3949 C	2 021LAMO5	ESTs	652651 IMAGE Consortium	AA217031
3950 C	4 021LAMO5	DNA segment, Chr 8, ERATO Doi 563, expressed	680058 IMAGE Consortium	AI451818
3951 C	6 021LAMO5	sulfotransferase-related protein SULT-X1	680668 IMAGE Consortium	AA237329
3952 C	8 021LAMO5	asialoglycoprotein receptor 1	680786 IMAGE Consortium	AA242413
3953 C	10 021LAMO5			
3954 C	12 021LAMO5	cytochrome P450 3A25	681642 IMAGE Consortium	AA237986
3955 E	2 021LAMO5	Msx-interacting-zinc finger	653396 IMAGE Consortium	AA240516
3956 E	4 021LAMO5	ESTs	672186 IMAGE Consortium	AI391368
3957 E	6 021LAMO5	plexin 2	677079 IMAGE Consortium	AA212820
3958 E	8 021LAMO5	ATP-binding cassette, sub-family B (MDR/TAP), member 1	692721 IMAGE Consortium	AA239727
3959 E	10 021LAMO5	ESTs, Moderately similar to PCP_HUMAN LYSOSOMAL PRO-X CARBOXYPEPTIDASE PRECURSOR [H.sapiens]	681298 IMAGE Consortium	AA250126
3960 E	12 021LAMO5	FTSJ-like protein	681646 IMAGE Consortium	AA237988
3961 G	2 021LAMO5	ESTs	653409 IMAGE Consortium	AA240290
3962 G	4 021LAMO5	Eph receptor A2	680273 IMAGE Consortium	AA237584
3963 G	6 021LAMO5	DNA segment, Chr 1, ERATO Doi 471, expressed	676503 IMAGE Consortium	AI391415
3964 G	8 021LAMO5	calbindin 2	697405 IMAGE Consortium	AA244813
3965 G	10 021LAMO5	ESTs	681313 IMAGE Consortium	AA241939
3966 G	12 021LAMO5	ESTs	680948 IMAGE Consortium	AA242024

3967 A	2 021SAMO6	ESTs	693384 IMAGE Consortium	AA245310
3968 A	4 021SAMO6	ESTs	719168 IMAGE Consortium	AA267124
3969 A	6 021SAMO6	ESTs	719429 IMAGE Consortium	AI464764
3970 A	8 021SAMO6	ESTs	720141 IMAGE Consortium	AA255328
3971 A	10 021SAMO6	ESTs	718360 IMAGE Consortium	AA266167
3972 A	12 021SAMO6	ESTs	717846 IMAGE Consortium	AA250545
3973 C	2 021SAMO6	hydroxysteroid 17-beta dehydrogenase 9	693488 IMAGE Consortium	AI425345
3974 C	4 021SAMO6	ESTs	719193 IMAGE Consortium	AI464728
3975 C	6 021SAMO6	ESTs, Weakly similar to DD10_HUMAN PROBABLE ATP-DEPENDENT RNA HELICASE DDX10 [H.sapiens]	719484 IMAGE Consortium	AI464770
3976 C	8 021SAMO6	Kupffer cell c-type lectin receptor	720272 IMAGE Consortium	AA270880
3977 C	10 021SAMO6	ESTs, Highly similar to BCL6_MOUSE B-CELL LYMPHOMA 6 PROTEIN HOMOLOG [M.musculus]	718376 IMAGE Consortium	AI464594
3978 C	12 021SAMO6	ESTs, Moderately similar to KIAA0240 [H.sapiens]	718647 IMAGE Consortium	AI464635
3979 E	2 021SAMO6	ESTs	693569 IMAGE Consortium	AA239385
3980 E	4 021SAMO6	ESTs	719226 IMAGE Consortium	AI464729
3981 E	6 021SAMO6	ESTs	718807 IMAGE Consortium	AI464667
3982 E	8 021SAMO6	coagulation factor X	720298 IMAGE Consortium	AA270769
3983 E	10 021SAMO6	ESTs	718398 IMAGE Consortium	AI464597
3984 E	12 021SAMO6	ESTs	718658 IMAGE Consortium	AI464636
3985 G	2 021SAMO6	ESTs, Weakly similar to S68418 protein phosphatase 1M chain M110 isoform - rat [R.norvegicus]	694458 IMAGE Consortium	AA218416
3986 G	4 021SAMO6	ESTs	719273 IMAGE Consortium	AI464742
3987 G	6 021SAMO6	ESTs	718865 IMAGE Consortium	AI464674
3988 G	8 021SAMO6	lupus nephritis-associated peptide 1	720411 IMAGE Consortium	AA261440
3989 G	10 021SAMO6	ESTs	718413 IMAGE Consortium	AI464600
3990 G	12 021SAMO6	ESTs	717988 IMAGE Consortium	AA261535
3991 A	2 021ZAMO7	ESTs	699526 IMAGE Consortium	AA245642
3992 A	4 021ZAMO7	ESTs	721070 IMAGE Consortium	AA267383
3993 A	6 021ZAMO7	ESTs	721364 IMAGE Consortium	AA267106
3994 A	8 021ZAMO7			
3995 A	10 021ZAMO7	ESTs	720976 IMAGE Consortium	AA266864
3996 A	12 021ZAMO7	ESTs	734587 IMAGE Consortium	AI529471
3997 C	2 021ZAMO7	ESTs	716430 IMAGE Consortium	AI466825
3998 C	4 021ZAMO7	ESTs	721100 IMAGE Consortium	AI464898
3999 C	6 021ZAMO7	ESTs	721395 IMAGE Consortium	AA267285
4000 C	8 021ZAMO7	ESTs	721571 IMAGE Consortium	AI606595
4001 C	10 021ZAMO7	immunoglobulin-associated beta	720992 IMAGE Consortium	AA266746
4002 C	12 021ZAMO7	ESTs	734652 IMAGE Consortium	AA272714
4003 E	2 021ZAMO7	neuronal guanine nucleotide exchange factor	716546 IMAGE Consortium	AI466833
4004 E	4 021ZAMO7	ESTs	721157 IMAGE Consortium	AI464905
4005 E	6 021ZAMO7	ESTs	722109 IMAGE Consortium	AI608056
4006 E	8 021ZAMO7	ESTs, Highly similar to unknown [H.sapiens]	720855 IMAGE Consortium	AI464858

4007 E	10 021ZAMO7	DNA segment, Chr 4, ERATO Doi 800, expressed	721721 IMAGE Consortium	AI595295
4008 E	12 021ZAMO7	ESTs	734685 IMAGE Consortium	AA261644
4009 G	2 021ZAMO7	ESTs	718025 IMAGE Consortium	AI466932
4010 G	4 021ZAMO7	ESTs	721219 IMAGE Consortium	AI464909
4011 G	6 021ZAMO7	DNA segment, Chr 2, Abbott 2 expressed	722152 IMAGE Consortium	AA268413
4012 G	8 021ZAMO7	ESTs, Weakly similar to RETROVIRUS-RELATED GAG POLYPROTEIN [Hamster intracisternal a-particle (iap-h18)]	720866 IMAGE Consortium	AI464860
4013 G	10 021ZAMO7	Mus musculus mRNA for microtubule associated protein 44 (Mtap44 gene)	721729 IMAGE Consortium	AA260779
4014 G	12 021ZAMO7	ESTs	733967 IMAGE Consortium	AI529445
4015 A	2 0216AMO8	ESTs	723067 IMAGE Consortium	AI463731
4016 A	4 0216AMO8	ESTs	723229 IMAGE Consortium	AI463758
4017 A	6 0216AMO8	ESTs	722855 IMAGE Consortium	AI608142
4018 A	8 0216AMO8	ESTs	749013 IMAGE Consortium	AI644761
4019 A	10 0216AMO8	ESTs, Highly similar to ALPHA-1-INHIBITOR III PRECURSOR [Rattus norvegicus]	747970 IMAGE Consortium	AA260963
4020 A	12 0216AMO8	ESTs, Highly similar to U63408_1 MRV11a protein [M.musculus]	749763 IMAGE Consortium	AA396288
4021 C	2 0216AMO8	ESTs	723080 IMAGE Consortium	AI463733
4022 C	4 0216AMO8	ESTs	722543 IMAGE Consortium	AA254137
4023 C	6 0216AMO8	ESTs, Highly similar to IDN4-GGTR8 [H.sapiens]	722904 IMAGE Consortium	AI608153
4024 C	8 0216AMO8	ESTs	749090 IMAGE Consortium	AA290529
4025 C	10 0216AMO8	ESTs	749437 IMAGE Consortium	AA289581
4026 C	12 0216AMO8	ESTs	749875 IMAGE Consortium	AA389264
4027 E	2 0216AMO8	ESTs	723087 IMAGE Consortium	AI463735
4028 E	4 0216AMO8	ESTs	722563 IMAGE Consortium	AA254161
4029 E	6 0216AMO8	ESTs, Highly similar to class A calcium channel variant riA-I [R.norvegicus]	733370 IMAGE Consortium	AI509417
4030 E	8 0216AMO8	ESTs	749109 IMAGE Consortium	AA288456
4031 E	10 0216AMO8	ESTs	749522 IMAGE Consortium	AI509226
4032 E	12 0216AMO8	ESTs	749894 IMAGE Consortium	AA389157
4033 G	2 0216AMO8	ESTs	722393 IMAGE Consortium	AA253967
4034 G	4 0216AMO8	ESTs	723313 IMAGE Consortium	AI463768
4035 G	6 0216AMO8	ESTs	733458 IMAGE Consortium	AI509421
4036 G	8 0216AMO8	ESTs	749242 IMAGE Consortium	AA289639
4037 G	10 0216AMO8	ESTs	749551 IMAGE Consortium	AA388685
4038 G	12 0216AMO8	ESTs	749971 IMAGE Consortium	AA388807
4039 A	2 021DAMO9	short stature homeobox 2	334541 IMAGE Consortium	W18334
4040 A	4 021DAMO9			
4041 A	6 021DAMO9			
4042 A	8 021DAMO9	ESTs, Moderately similar to unnamed protein product [H.sapiens]	746512 IMAGE Consortium	AI607671

4043 A	10 021DAMO9	ESTs	751277 IMAGE Consortium	AI508699
4044 A	12 021DAMO9	polymerase (DNA directed), mu	750768 IMAGE Consortium	AA289208
4045 C	2 021DAMO9	ESTs	334163 IMAGE Consortium	AI391284
4046 C	4 021DAMO9	ESTs	334553 IMAGE Consortium	W18529
4047 C	6 021DAMO9	ESTs	735705 IMAGE Consortium	AA269814
4048 C	8 021DAMO9	iroquois related homeobox 3 (Drosophila)	746622 IMAGE Consortium	AA268059
4049 C	10 021DAMO9	ESTs	751283 IMAGE Consortium	AA289424
4050 C	12 021DAMO9	ESTs	751538 IMAGE Consortium	AA396036
4051 E	2 021DAMO9	ESTs	335067 IMAGE Consortium	AI390101
4052 E	4 021DAMO9	reticulon 2 (Z-band associated protein)	334677 IMAGE Consortium	W18313
4053 E	6 021DAMO9			
4054 E	8 021DAMO9	sema domain, immunoglobulin domain (Ig), and GPI membrane anchor, (semaphorin) 7A	746763 IMAGE Consortium	AI607683
4055 E	10 021DAMO9	ESTs	750585 IMAGE Consortium	AA472955
4056 E	12 021DAMO9	ESTs	750853 IMAGE Consortium	AA290197
4057 G	2 021DAMO9	small inducible cytokine A21b (serine)	335077 IMAGE Consortium	W17930
4058 G	4 021DAMO9	ESTs	334770 IMAGE Consortium	W36425
4059 G	6 021DAMO9	ESTs	737488 IMAGE Consortium	AI466352
4060 G	8 021DAMO9	lymphotoxin A	751270 IMAGE Consortium	AA288231
4061 G	10 021DAMO9	ESTs	750597 IMAGE Consortium	AI508480
4062 G	12 021DAMO9	signaling lymphocyte activation molecule	750873 IMAGE Consortium	AA289183
4063 A	2 021XAMOA	ESTs, Highly similar to transcription factor MAZR [M.musculus]	750932 IMAGE Consortium	AA289140
4064 A	4 021XAMOA	ESTs	750284 IMAGE Consortium	AI606538
4065 A	6 021XAMOA	ESTs	752202 IMAGE Consortium	AA289072
4066 A	8 021XAMOA	poliovirus receptor-related 1	762124 IMAGE Consortium	AI508995
4067 A	10 021XAMOA	ESTs, Weakly similar to unnamed protein product [H.sapiens]	762641 IMAGE Consortium	AA271838
4068 A	12 021XAMOA	ESTs	764362 IMAGE Consortium	AA277444
4069 C	2 021XAMOA	ESTs	750935 IMAGE Consortium	AI508515
4070 C	4 021XAMOA	TRAF family member-associated Nf-kappa B activator	751006 IMAGE Consortium	AA289404
4071 C	6 021XAMOA	ESTs	752234 IMAGE Consortium	AI508568
4072 C	8 021XAMOA	ESTs	763004 IMAGE Consortium	AA271596
4073 C	10 021XAMOA	ESTs	764137 IMAGE Consortium	AA285830
4074 C	12 021XAMOA	ESTs	762960 IMAGE Consortium	AI595066
4075 E	2 021XAMOA	ESTs	750952 IMAGE Consortium	AA289146
4076 E	4 021XAMOA	ESTs, Weakly similar to I60486 gene trg protein - rat [R.norvegicus]	751015 IMAGE Consortium	AA289502
4077 E	6 021XAMOA	ESTs	752250 IMAGE Consortium	AI508571
4078 E	8 021XAMOA	ESTs	761633 IMAGE Consortium	AI508985
4079 E	10 021XAMOA	ESTs	764165 IMAGE Consortium	AI606557
4080 E	12 021XAMOA	ESTs	764427 IMAGE Consortium	AI606593
4081 G	2 021XAMOA	ESTs	751766 IMAGE Consortium	AI508759

4082 G	4 021XAMOA	CD6 antigen	750331 IMAGE Consortium	AA396636
4083 G	6 021XAMOA	ESTs	752323 IMAGE Consortium	AI508580
4084 G	8 021XAMOA	ESTs, Highly similar to MITOCHONDRIAL LON PROTEASE HOMOLOG PRECURSOR [Homo sapiens]	762385 IMAGE Consortium	AA274011
4085 G	10 021XAMOA	EST	764178 IMAGE Consortium	AI462991
4086 G	12 021XAMOA	schlafen 1	764568 IMAGE Consortium	AA273820
4087 A	2 0214AMOB	ERG-associated protein	778367 IMAGE Consortium	AA413473
4088 A	4 0214AMOB	A kinase anchor protein	778529 IMAGE Consortium	AA413485
4089 A	6 0214AMOB	ESTs, Weakly similar to CIRP [R.norvegicus]	778848 IMAGE Consortium	AA414162
4090 A	8 0214AMOB	ESTs	777345 IMAGE Consortium	AA276927
4091 A	10 0214AMOB	ESTs	765080 IMAGE Consortium	AA273401
4092 A	12 0214AMOB	ESTs	765536 IMAGE Consortium	AA275270
4093 C	2 0214AMOB	ESTs, Moderately similar to HIRA-interacting protein 3 [H.sapiens]	779119 IMAGE Consortium	AA467564
4094 C	4 0214AMOB	ESTs	779307 IMAGE Consortium	AA413985
4095 C	6 0214AMOB	transcription elongation factor A (SII) 1	778905 IMAGE Consortium	AA414169
4096 C	8 0214AMOB	microsomal triglyceride transfer protein	777411 IMAGE Consortium	AA276406
4097 C	10 0214AMOB	ESTs	764446 IMAGE Consortium	AI606592
4098 C	12 0214AMOB			
4099 E	2 0214AMOB	cadherin 3	779189 IMAGE Consortium	AA467584
4100 E	4 0214AMOB	NIMA (never in mitosis gene a)-related expressed kinase 4	779355 IMAGE Consortium	AA413968
4101 E	6 0214AMOB	ESTs	778944 IMAGE Consortium	AA414286
4102 E	8 0214AMOB	ESTs	777493 IMAGE Consortium	AA276636
4103 E	10 0214AMOB	ESTs	765259 IMAGE Consortium	AI510269
4104 E	12 0214AMOB	ESTs	765825 IMAGE Consortium	AA274768
4105 G	2 0214AMOB	prefoldin 5	778480 IMAGE Consortium	AA413511
4106 G	4 0214AMOB	ESTs	778662 IMAGE Consortium	AA413651
4107 G	6 0214AMOB	ESTs, Weakly similar to AF218940_1 formin-2 [M.musculus]	778972 IMAGE Consortium	AA414321
4108 G	8 0214AMOB	ESTs, Highly similar to KS63_MOUSE RIBOSOMAL PROTEIN S6 KINASE II ALPHA 3 [M.musculus]	777538 IMAGE Consortium	AA276748
4109 G	10 0214AMOB	interferon-g induced GTPase	765294 IMAGE Consortium	AA275144
4110 G	12 0214AMOB	ESTs	765934 IMAGE Consortium	AA277945
4111 A	2 021BAMOC	hippocampus abundant gene transcript 1	779029 IMAGE Consortium	AA414014
4112 A	4 021BAMOC	ESTs, Highly similar to HYDROXYMETHYLGLUTARYL- COA SYNTHASE, CYTOPLASMIC [Rattus norvegicus]	779241 IMAGE Consortium	AA466979
4113 A	6 021BAMOC	phosphatidylinositol 3-kinase, catalytic, alpha polypeptide	779408 IMAGE Consortium	AA414173
4114 A	8 021BAMOC	ESTs	779656 IMAGE Consortium	AA466935
4115 A	10 021BAMOC	emerin	779928 IMAGE Consortium	AA414844
4116 A	12 021BAMOC	ESTs, Weakly similar to 2013347A huckebein gene [D.melanogaster]	791102 IMAGE Consortium	AA415189
4117 C	2 021BAMOC	interferon regulatory factor 3	779035 IMAGE Consortium	AA413767



4118 C	4 021BAMOC	DNA segment, Chr 11, Wayne State University 99, expressed	779292 IMAGE Consortium	AA466988
4119 C	6 021BAMOC	ESTs, Weakly similar to DNA-BINDING PROTEIN BMI-1 [Mus musculus]	779427 IMAGE Consortium	AA414199
4120 C	8 021BAMOC	ESTs	779670 IMAGE Consortium	AA413851
4121 C	10 021BAMOC	golgi autoantigen, golgin subfamily a, 5	780020 IMAGE Consortium	AA414933
4122 C	12 021BAMOC	ESTs, Highly similar to S56145 BS69 protein - human [H.sapiens]	791107 IMAGE Consortium	AA415257
4123 E	2 021BAMOC	homeodomain interacting protein kinase 1	779039 IMAGE Consortium	AA413769
4124 E	4 021BAMOC	Mus musculus ubiquitinating enzyme E2-230 kDa mRNA, partial cds	779299 IMAGE Consortium	AA445870
4125 E	6 021BAMOC	DNA segment, Chr 11, ERATO Doi 730, expressed	779440 IMAGE Consortium	AA414450
4126 E	8 021BAMOC	ESTs, Weakly similar to CG13585 gene product [D.melanogaster]	779673 IMAGE Consortium	AA413755
4127 E	10 021BAMOC	abl-interactor 1	780107 IMAGE Consortium	AA414750
4128 E	12 021BAMOC			
4129 G	2 021BAMOC	RAB2, member RAS oncogene family	779061 IMAGE Consortium	AA413796
4130 G	4 021BAMOC	Casitas B-lineage lymphoma	779344 IMAGE Consortium	AA414809
4131 G	6 021BAMOC	stromal cell derived factor receptor 1	779458 IMAGE Consortium	AA414208
4132 G	8 021BAMOC	ESTs, Highly similar to MDM2-like p53-binding protein [M.musculus]	779706 IMAGE Consortium	AA414627
4133 G	10 021BAMOC	ESTs	790235 IMAGE Consortium	AI614783
4134 G	12 021BAMOC	ESTs	791154 IMAGE Consortium	AA415510
4135 A	2 021IAMOD	ESTs, Highly similar to CICL_RAT CHLORIDE CHANNEL PROTEIN CLC-K2 [R.norvegicus]	790543 IMAGE Consortium	AA289182
4136 A	4 021IAMOD	ESTs	790935 IMAGE Consortium	AA415319
4137 A	6 021IAMOD	DNA segment, Chr 7, ERATO Doi 743, expressed	792616 IMAGE Consortium	AA415231
4138 A	8 021IAMOD	ESTs, Moderately similar to AF164799_1 putatative 28 kDa protein [H.sapiens]	792867 IMAGE Consortium	AA415520
4139 A	10 021IAMOD	ribosomal protein S5	793037 IMAGE Consortium	AA444231
4140 A	12 021IAMOD	testis-specific poly(A) polymerase	791952 IMAGE Consortium	AA414244
4141 C	2 021IAMOD	DNA segment, Chr 6, ERATO Doi 109, expressed	791287 IMAGE Consortium	AA420359
4142 C	4 021IAMOD	LIM domain binding 1	790953 IMAGE Consortium	AA415350
4143 C	6 021IAMOD	translocase of inner mitochondrial membrane 23 homolog (yeast)	791901 IMAGE Consortium	AA414425
4144 C	8 021IAMOD	ESTs	792893 IMAGE Consortium	AA415525
4145 C	10 021IAMOD	ESTs, Weakly similar to GOLIATH PROTEIN [Drosophila melanogaster]	793104 IMAGE Consortium	AA437863
4146 C	12 021IAMOD	ESTs, Highly similar to dJ108K11.3 [H.sapiens]	791958 IMAGE Consortium	AA414247
4147 E	2 021IAMOD	ESTs	791300 IMAGE Consortium	AA420355
4148 E	4 021IAMOD	ESTs	793204 IMAGE Consortium	AA437885
4149 E	6 021IAMOD	ESTs, Moderately similar to KIAA0661 protein [H.sapiens]	792645 IMAGE Consortium	AA416433
4150 E	8 021IAMOD	cytoplasmic polyadenylation element binding protein	792916 IMAGE Consortium	AA415529

4151 E	10 021IAMOD	ESTs	793107 IMAGE Consortium	AA437874
4152 E	12 021IAMOD			
4153 G	2 021IAMOD	ESTs, Weakly similar to AF167320_1 zinc finger protein ZFP113 [M.musculus]	790669 IMAGE Consortium	AA388385
4154 G	4 021IAMOD	ESTs, Moderately similar to diphthamide biosynthesis protein-2 [H.sapiens]	792561 IMAGE Consortium	AA416351
4155 G	6 021IAMOD	Mus musculus mRNA for calsyntenin-1 protein	792654 IMAGE Consortium	AA416337
4156 G	8 021IAMOD	Public domain EST	792938 IMAGE Consortium	
4157 G	10 021IAMOD	phosphatidylethanolamine binding protein	793134 IMAGE Consortium	AA437975
4158 G	12 021IAMOD			
4159 A	2 021PAMOE	ELAV (embryonic lethal, abnormal vision, Drosophila)-like 2 (Hu antigen B)	791414 IMAGE Consortium	AA416369
4160 A	4 021PAMOE	endogenous retroviral sequence 4 (with leucine t-RNA primer)	791623 IMAGE Consortium	AA437585
4161 A	6 021PAMOE	thymoma viral proto-oncogene	792852 IMAGE Consortium	AA415535
4162 A	8 021PAMOE	ESTs, Highly similar to KIAA0309 [H.sapiens]	792998 IMAGE Consortium	AA437626
4163 A	10 021PAMOE	endogenous retroviral sequence 4 (with leucine t-RNA primer)	793052 IMAGE Consortium	AA444227
4164 A	12 021PAMOE	cerebellar degeneration-related 2	793192 IMAGE Consortium	AA438015
4165 C	2 021PAMOE	DNA segment, Chr 4, ERATO Doi 786, expressed	791432 IMAGE Consortium	AA416407
4166 C	4 021PAMOE	ESTs, Moderately similar to KIAA1165 protein [H.sapiens]	791672 IMAGE Consortium	AA437603
4167 C	6 021PAMOE	folylpolyglutamyl synthetase	792857 IMAGE Consortium	AA416279
4168 C	8 021PAMOE	ESTs	793755 IMAGE Consortium	AA415711
4169 C	10 021PAMOE	ESTs, Highly similar to zinc transporter ZnT-1 [M.musculus]	793065 IMAGE Consortium	AA444253
4170 C	12 021PAMOE	solute carrier family 30 (zinc transporter), member 3	350388 IMAGE Consortium	W34988
4171 E	2 021PAMOE	ESTs	791447 IMAGE Consortium	AA416421
4172 E	4 021PAMOE	ESTs, Weakly similar to KIAA0542 protein [H.sapiens]	791692 IMAGE Consortium	AA414653
4173 E	6 021PAMOE	Son of sevenless homolog 1, (Drosophila)	792885 IMAGE Consortium	AA415373
4174 E	8 021PAMOE	hypoxia inducible factor 1, alpha subunit	793858 IMAGE Consortium	AA422602
4175 E	10 021PAMOE	peroxisomal membrane protein 20	793074 IMAGE Consortium	AA444248
4176 E	12 021PAMOE	organic cationic transporter-like 1	352949 IMAGE Consortium	W34761
4177 G	2 021PAMOE	ESTs, Moderately similar to COP9 PROTEIN [Arabidopsis thaliana]	791523 IMAGE Consortium	AA437443
4178 G	4 021PAMOE			
4179 G	6 021PAMOE	ESTs	793618 IMAGE Consortium	AA415629
4180 G	8 021PAMOE	ESTs, Weakly similar to P52rIPK [H.sapiens]	793901 IMAGE Consortium	AA422723
4181 G	10 021PAMOE	degenerative spermatocyte homolog (Drosophila)	793115 IMAGE Consortium	AA437876
4182 G	12 021PAMOE	LIM domin binding 3	353911 IMAGE Consortium	W44251
4183 A	2 021WAMOF	ESTs	355186 IMAGE Consortium	W47900
4184 A	4 021WAMOF	ESTs	349934 IMAGE Consortium	AI391160
4185 A	6 021WAMOF	ESTs	350497 IMAGE Consortium	AI450492
4186 A	8 021WAMOF	ESTs, Moderately similar to M phase phosphoprotein 10	352598 IMAGE Consortium	W33952

		[H.sapiens]		
4187 A	10 021WAMOF	ESTs	809050 IMAGE Consortium	AA466043
4188 A	12 021WAMOF	ESTs	808939 IMAGE Consortium	AI596897
4189 C	2 021WAMOF	ESTs	355216 IMAGE Consortium	W47916
4190 C	4 021WAMOF			
4191 C	6 021WAMOF	ESTs	352088 IMAGE Consortium	AI430652
4192 C	8 021WAMOF	ESTs	808049 IMAGE Consortium	AI597441
4193 C	10 021WAMOF	ESTs	807669 IMAGE Consortium	AA466822
4194 C	12 021WAMOF	ESTs	804186 IMAGE Consortium	AA474446
4195 E	2 021WAMOF	ESTs, Highly similar to warts/lats-like kinase [M.musculus]	355235 IMAGE Consortium	W47897
4196 E	4 021WAMOF	downstream of tyrosine kinase 2	350190 IMAGE Consortium	AI391168
4197 E	6 021WAMOF	ESTs	350693 IMAGE Consortium	AI450507
4198 E	8 021WAMOF	ESTs	807433 IMAGE Consortium	AA412939
4199 E	10 021WAMOF	variable group of 2-cell-stage gene family	793677 IMAGE Consortium	AA415841
4200 E	12 021WAMOF	ESTs, Weakly similar to TXTP_RAT TRICARBOXYLATE TRANSPORT PROTEIN PRECURSOR [R.norvegicus]	804238 IMAGE Consortium	AA474460
4201 G	2 021WAMOF	ESTs	353866 IMAGE Consortium	AI427227
4202 G	4 021WAMOF	ESTs, Weakly similar to CARBONYL REDUCTASE [Mus musculus]	350227 IMAGE Consortium	AI322339
4203 G	6 021WAMOF	ESTs, Moderately similar to ZINC FINGER PROTEIN ZFMSA12A [Micropterus salmoides]	352184 IMAGE Consortium	AI893133
4204 G	8 021WAMOF	pyruvate kinase 3	807460 IMAGE Consortium	AA412831
4205 G	10 021WAMOF	ESTs	807717 IMAGE Consortium	AA466836
4206 G	12 021WAMOF			
4207 A	2 0213AMOG	ESTs	803663 IMAGE Consortium	AA422771
4208 A	4 0213AMOG	DNA segment, Chr 5, ERATO Doi 577, expressed	803970 IMAGE Consortium	AA435343
4209 A	6 0213AMOG	ESTs	833435 IMAGE Consortium	AA465951
4210 A	8 0213AMOG	ESTs	835499 IMAGE Consortium	AA422442
4211 A	10 0213AMOG	EST	831667 IMAGE Consortium	AI607423
4212 A	12 0213AMOG	ESTs	818391 IMAGE Consortium	AA434839
4213 C	2 0213AMOG	ESTs, Weakly similar to HC1 ORF [M.musculus]	803675 IMAGE Consortium	AA422835
4214 C	4 0213AMOG	ESTs, Moderately similar to S1-1 protein [R.norvegicus]	807022 IMAGE Consortium	AA404085
4215 C	6 0213AMOG	ESTs, Highly similar to HISTONE MACRO-H2A.1 [Rattus norvegicus]	833510 IMAGE Consortium	AA465981
4216 C	8 0213AMOG			
4217 C	10 0213AMOG	ESTs	831669 IMAGE Consortium	AI549703
4218 C	12 0213AMOG	ESTs	818413 IMAGE Consortium	AI596293
4219 E	2 0213AMOG	germ cell-specific gene 2	803765 IMAGE Consortium	AA427283
4220 E	4 0213AMOG	transcription elongation factor B (SIII), polypeptide 3 (110kD)	833192 IMAGE Consortium	AA467597
4221 E	6 0213AMOG	ESTs, Moderately similar to GTPASE-ACTIVATING PROTEIN 1 [Drosophila melanogaster]	833537 IMAGE Consortium	AA432707
4222 E	8 0213AMOG	ESTs	835808 IMAGE Consortium	AA422609

4223 E	10 0213AMOG	ESTs	831751 IMAGE Consortium	AA445875
4224 E	12 0213AMOG	ESTs, Weakly similar to PPAR gamma coactivator [M.musculus]	818488 IMAGE Consortium	AI596296
4225 G	2 0213AMOG	ESTs, Weakly similar to myosin [M.musculus]	803785 IMAGE Consortium	AA422950
4226 G	4 0213AMOG	inducible IKappaB kinase	832503 IMAGE Consortium	AA423538
4227 G	6 0213AMOG	ESTs, Moderately similar to neuronal voltage-gated sodium channel alpha subunit [M.musculus]	832845 IMAGE Consortium	AI549833
4228 G	8 0213AMOG	ESTs, Highly similar to AF151868_1 CGI-110 protein [H.sapiens]	835822 IMAGE Consortium	AA422611
4229 G	10 0213AMOG	ESTs	831850 IMAGE Consortium	AA444672
4230 G	12 0213AMOG	ESTs, Highly similar to KIAA0903 protein [H.sapiens]	818608 IMAGE Consortium	AA444308
4231 A	2 021AAMOH	ADP-ribosylation-like 4	820282 IMAGE Consortium	AA423230
4232 A	4 021AAMOH	ESTs	848961 IMAGE Consortium	AI552219
4233 A	6 021AAMOH	ESTs	847035 IMAGE Consortium	AI553086
4234 A	8 021AAMOH	ESTs	845866 IMAGE Consortium	AI553016
4235 A	10 021AAMOH			
4236 A	12 021AAMOH	retinoblastoma-binding protein 9	848680 IMAGE Consortium	AA575702
4237 C	2 021AAMOH	ESTs	821048 IMAGE Consortium	AI552333
4238 C	4 021AAMOH	ESTs, Highly similar to PROTEIN TRANSPORT PROTEIN SEC23 [Saccharomyces cerevisiae]	836320 IMAGE Consortium	AA426912
4239 C	6 021AAMOH	ESTs	846329 IMAGE Consortium	AA419710
4240 C	8 021AAMOH	ESTs	847300 IMAGE Consortium	AA420078
4241 C	10 021AAMOH	ESTs, Weakly similar to orphan transporter isoform A10 [M.musculus]	848385 IMAGE Consortium	AA445667
4242 C	12 021AAMOH	ectodysplasin-A	851504 IMAGE Consortium	AA461806
4243 E	2 021AAMOH	ESTs	821108 IMAGE Consortium	AI552341
4244 E	4 021AAMOH	ESTs	836341 IMAGE Consortium	AA426869
4245 E	6 021AAMOH	ESTs	846350 IMAGE Consortium	AA419720
4246 E	8 021AAMOH	ESTs, Weakly similar to actin-filament binding protein Frabin [R.norvegicus]	847339 IMAGE Consortium	AI552157
4247 E	10 021AAMOH	ESTs	846978 IMAGE Consortium	AA433722
4248 E	12 021AAMOH	ESTs, Moderately similar to C321D2.1 [H.sapiens]	849401 IMAGE Consortium	AA457857
4249 G	2 021AAMOH	growth differentiation factor 15	820434 IMAGE Consortium	AA423314
4250 G	4 021AAMOH	cadherin EGF LAG seven-pass G-type receptor	836398 IMAGE Consortium	AA426883
4251 G	6 021AAMOH	ESTs, Moderately similar to AF113539_1 hypothalamus protein HT001 [H.sapiens]	846396 IMAGE Consortium	AI553045
4252 G	8 021AAMOH	ESTs, Weakly similar to putative protein kinase [M.musculus]	846667 IMAGE Consortium	AI553059
4253 G	10 021AAMOH	ESTs	849139 IMAGE Consortium	AI552231
4254 G	12 021AAMOH	ESTs, Weakly similar to polycomb-group protein [R.norvegicus]	849466 IMAGE Consortium	AI552270
4255 A	2 021HAMOI	membrane metallo endopeptidase	849871 IMAGE Consortium	AA450725
4256 A	4 021HAMOI	ESTs	850480 IMAGE Consortium	AA451320

4257 A	6 021HAMOI	EST	860247 IMAGE Consortium	AI481812
4258 A	8 021HAMOI	ESTs, Weakly similar to CD94 [M.musculus]	860632 IMAGE Consortium	AA500171
4259 A	10 021HAMOI	ESTs	875286 IMAGE Consortium	AA475853
4260 A	12 021HAMOI	ESTs	874772 IMAGE Consortium	AA472499
4261 C	2 021HAMOI	recombining binding protein suppressor of hairless (Drosophila)	850016 IMAGE Consortium	AA444538
4262 C	4 021HAMOI	ESTs	851217 IMAGE Consortium	AA462941
4263 C	6 021HAMOI	ESTs	860272 IMAGE Consortium	AA500905
4264 C	8 021HAMOI	ESTs	860864 IMAGE Consortium	AA500239
4265 C	10 021HAMOI	ESTs	873858 IMAGE Consortium	AA471954
4266 C	12 021HAMOI	ESTs	875572 IMAGE Consortium	AA475590
4267 E	2 021HAMOI	ESTs	850136 IMAGE Consortium	AA450692
4268 E	4 021HAMOI	delta-like 4 homolog (Drosophila)	850541 IMAGE Consortium	AA450936
4269 E	6 021HAMOI	ESTs	860342 IMAGE Consortium	AI613599
4270 E	8 021HAMOI	ESTs	849378 IMAGE Consortium	AI552248
4271 E	10 021HAMOI	DNA segment, Chr 6, ERATO Doi 245, expressed	876023 IMAGE Consortium	AA475294
4272 E	12 021HAMOI	ESTs	876409 IMAGE Consortium	AA475884
4273 G	2 021HAMOI	ESTs	850186 IMAGE Consortium	AA451407
4274 G	4 021HAMOI	ESTs	860055 IMAGE Consortium	AA500774
4275 G	6 021HAMOI	ESTs	850974 IMAGE Consortium	AA462386
4276 G	8 021HAMOI	ESTs	874199 IMAGE Consortium	AA472312
4277 G	10 021HAMOI	Mus musculus brain cDNA, clone MNCb-1826	875348 IMAGE Consortium	AA475218
4278 G	12 021HAMOI	ESTs	876436 IMAGE Consortium	AA475897
4279 A	2 021OAMOJ	ESTs, Weakly similar to INTERFERON-GAMMA RECEPTOR BETA CHAIN PRECURSOR [Homo sapiens]	871598 IMAGE Consortium	AA462218
4280 A	4 021OAMOJ	ESTs	871045 IMAGE Consortium	AA462447
4281 A	6 021OAMOJ	ESTs	890432 IMAGE Consortium	AA509560
4282 A	8 021OAMOJ	ESTs, Moderately similar to GTPASE-ACTIVATING PROTEIN [Bos taurus]	876231 IMAGE Consortium	AA475664
4283 A	10 021OAMOJ	intersectin (SH3 domain protein 1A)	888505 IMAGE Consortium	AA497627
4284 A	12 021OAMOJ	slug, chicken homolog	902836 IMAGE Consortium	AA518917
4285 C	2 021OAMOJ	ESTs	873062 IMAGE Consortium	AA466199
4286 C	4 021OAMOJ	RNA polymerase II 1	871109 IMAGE Consortium	AI552902
4287 C	6 021OAMOJ	ESTs	890479 IMAGE Consortium	AA509863
4288 C	8 021OAMOJ	ESTs	890828 IMAGE Consortium	AA510659
4289 C	10 021OAMOJ	ESTs, Weakly similar to F-box protein FBX13 [M.musculus]	888529 IMAGE Consortium	AA498791
4290 C	12 021OAMOJ	ESTs	904353 IMAGE Consortium	AI553232
4291 E	2 021OAMOJ	ESTs	871640 IMAGE Consortium	AA466153
4292 E	4 021OAMOJ	vertebrate homolog of C. elegans Lin-7 type 3	870825 IMAGE Consortium	AA462021
4293 E	6 021OAMOJ	ATPase, Na <sup>+</sup> /K <sup>+</sup> transporting, alpha 4 polypeptide	890575 IMAGE Consortium	AA509743
4294 E	8 021OAMOJ	ESTs	890841 IMAGE Consortium	AA510688
4295 E	10 021OAMOJ	T-box 15	902021 IMAGE Consortium	AA517906

4296 E	12 021OAMOJ	Mus musculus 129 cryptdin 6 gene, complete cds	903673 IMAGE Consortium	AA623172
4297 G	2 021OAMOJ	ESTs	873122 IMAGE Consortium	AA473684
4298 G	4 021OAMOJ	ESTs	870873 IMAGE Consortium	AI552890
4299 G	6 021OAMOJ	ESTs, Weakly similar to similar to citrate lyase beta chain [C.elegans]	891364 IMAGE Consortium	AA510712
4300 G	8 021OAMOJ	ESTs	890972 IMAGE Consortium	AA511209
4301 G	10 021OAMOJ	ESTs, Weakly similar to CG3714 gene product [D.melanogaster]	888713 IMAGE Consortium	AA498776
4302 G	12 021OAMOJ	ESTs	903683 IMAGE Consortium	AA623175
4303 A	2 021VAMOK	ESTs	903781 IMAGE Consortium	AA623237
4304 A	4 021VAMOK	ESTs	904885 IMAGE Consortium	AA519024
4305 A	6 021VAMOK	ESTs, Highly similar to MAN1_RAT ALPHA-MANNOSIDASE [R.norvegicus]	904061 IMAGE Consortium	AA518570
4306 A	8 021VAMOK	ESTs, Highly similar to AF151825_1 CGI-67 protein [H.sapiens]	948596 IMAGE Consortium	AI561842
4307 A	10 021VAMOK	deoxyribonuclease II	959560 IMAGE Consortium	AA546528
4308 A	12 021VAMOK	DNA segment, Chr 15, ERATO Doi 597, expressed	959776 IMAGE Consortium	AA546928
4309 C	2 021VAMOK	ESTs	903863 IMAGE Consortium	AA521580
4310 C	4 021VAMOK	ESTs	905607 IMAGE Consortium	AI553323
4311 C	6 021VAMOK			
4312 C	8 021VAMOK	ESTs	949454 IMAGE Consortium	AA537652
4313 C	10 021VAMOK	ESTs, Weakly similar to casein kinase [M.musculus]	959589 IMAGE Consortium	AA546556
4314 C	12 021VAMOK	tyrosine 3-monooxygenase/tryptophan 5-monooxygenase activation protein, zeta polypeptide	959796 IMAGE Consortium	AA546945
4315 E	2 021VAMOK	ESTs	904626 IMAGE Consortium	AA518115
4316 E	4 021VAMOK	ESTs	904935 IMAGE Consortium	AI553275
4317 E	6 021VAMOK	Mus musculus HSP40-like protein mRNA, partial sequence	902646 IMAGE Consortium	AA518878
4318 E	8 021VAMOK	ESTs, Weakly similar to CD63_MOUSE CD63 ANTIGEN [M.musculus]	949540 IMAGE Consortium	AA537130
4319 E	10 021VAMOK	cell division cycle 2 homolog (S. pombe)-like 2	959628 IMAGE Consortium	AA546645
4320 E	12 021VAMOK	CD20 antigen	959819 IMAGE Consortium	AA546971
4321 G	2 021VAMOK	matrix metalloproteinase 17	904649 IMAGE Consortium	AA518143
4322 G	4 021VAMOK	dentatorubral pallidoluysian atrophy	904961 IMAGE Consortium	AI553277
4323 G	6 021VAMOK	RNA binding motif, single stranded interacting protein 2	903392 IMAGE Consortium	AA518465
4324 G	8 021VAMOK	PH domain containing protein in retina 1	948884 IMAGE Consortium	AA544877
4325 G	10 021VAMOK	ESTs	959668 IMAGE Consortium	AA546740
4326 G	12 021VAMOK	ESTs	959860 IMAGE Consortium	AA547033
4327 A	2 0212AMOL	ESTs	371145 IMAGE Consortium	W53394
4328 A	4 0212AMOL	fibroblast growth factor 12	368515 IMAGE Consortium	AI425567
4329 A	6 0212AMOL	ESTs	922327 IMAGE Consortium	AA512317
4330 A	8 0212AMOL	Williams-Beuren syndrome chromosome region 14 homolog (human)	935589 IMAGE Consortium	AA543589

4331 A	10 0212AMOL	MAD homolog 7 ( <i>Drosophila</i> )	944734 IMAGE Consortium	AA545284
4332 A	12 0212AMOL	NTF2-related export protein	959923 IMAGE Consortium	AA547141
4333 C	2 0212AMOL	ESTs, Weakly similar to ORF YKL030w [ <i>S.cerevisiae</i> ]	355892 IMAGE Consortium	W49172
4334 C	4 0212AMOL	paired-like homeodomain transcription factor 1	368564 IMAGE Consortium	W61694
4335 C	6 0212AMOL	ESTs	921656 IMAGE Consortium	AI550863
4336 C	8 0212AMOL	ESTs, Highly similar to HYPOTHETICAL 18.5 KD PROTEIN C12G12.05C IN CHROMOSOME I [ <i>Schizosaccharomyces pombe</i> ]	944309 IMAGE Consortium	AA545507
4337 C	10 0212AMOL	ESTs	944765 IMAGE Consortium	AA545326
4338 C	12 0212AMOL	RAD17 homolog ( <i>S. pombe</i> )	959937 IMAGE Consortium	AA547157
4339 E	2 0212AMOL	ESTs, Highly similar to CA19_MOUSE COLLAGEN ALPHA 1(IX) CHAIN PRECURSOR [M.musculus]	368032 IMAGE Consortium	W53670
4340 E	4 0212AMOL	ESTs, Highly similar to zyginl [ <i>R.norvegicus</i> ]	368622 IMAGE Consortium	AI425572
4341 E	6 0212AMOL	ESTs	922453 IMAGE Consortium	AI550991
4342 E	8 0212AMOL	ESTs, Highly similar to DCC_MOUSE TUMOR SUPPRESSOR PROTEIN DCC PRECURSOR [M.musculus]	944340 IMAGE Consortium	AA545523
4343 E	10 0212AMOL	peptide N-glycanase homolog ( <i>S.cerevisiae</i> )	944778 IMAGE Consortium	AA545321
4344 E	12 0212AMOL	undifferentiated embryonic cell transcription factor 1	960016 IMAGE Consortium	AA547242
4345 G	2 0212AMOL	ESTs	367405 IMAGE Consortium	AI427388
4346 G	4 0212AMOL	achaete-scute complex homolog-like ( <i>Drosophila</i> )	368696 IMAGE Consortium	W54161
4347 G	6 0212AMOL	RAB11a, member RAS oncogene family	921804 IMAGE Consortium	AI550872
4348 G	8 0212AMOL	Unc-51 like kinase 1 ( <i>C. elegans</i> )	944386 IMAGE Consortium	AA545560
4349 G	10 0212AMOL	ESTs	944794 IMAGE Consortium	AA545397
4350 G	12 0212AMOL	ESTs	960038 IMAGE Consortium	AA547263
4351 A	2 0219AMOM	ESTs	960256 IMAGE Consortium	AA547096
4352 A	4 0219AMOM	ESTs, Highly similar to epsilon isoform of 61kDa regulatory subunit of PP2A [ <i>H.sapiens</i> ]	960465 IMAGE Consortium	AA549314
4353 A	6 0219AMOM	ESTs	960718 IMAGE Consortium	AA549028
4354 A	8 0219AMOM	growth arrest and DNA-damage-inducible, alpha	961868 IMAGE Consortium	AA553242
4355 A	10 0219AMOM	Mus musculus eukaryotic translation elongation factor 1-delta mRNA, complete cds	961189 IMAGE Consortium	AA547446
4356 A	12 0219AMOM	ESTs, Weakly similar to dJ500L14.1 [ <i>H.sapiens</i> ]	1003018 IMAGE Consortium	AA684456
4357 C	2 0219AMOM	ESTs, Weakly similar to contains transmembrane [M.musculus]	960281 IMAGE Consortium	AA549717
4358 C	4 0219AMOM	ESTs, Moderately similar to dJ568C11.2 [ <i>H.sapiens</i> ]	960487 IMAGE Consortium	AA549326
4359 C	6 0219AMOM	ESTs	960741 IMAGE Consortium	AA549056
4360 C	8 0219AMOM	zuotin related factor 2	961888 IMAGE Consortium	AA672310
4361 C	10 0219AMOM	predicted gene ICRFP703B1614Q5.6	961376 IMAGE Consortium	AA547668
4362 C	12 0219AMOM	RNA and export factor binding protein 2	1003028 IMAGE Consortium	AA684459
4363 E	2 0219AMOM	homolog of human MLLT2 unidentified gene	961002 IMAGE Consortium	AA549636
4364 E	4 0219AMOM	ESTs	961256 IMAGE Consortium	AA547535
4365 E	6 0219AMOM	ESTs, Moderately similar to KIAA0242 protein [ <i>H.sapiens</i> ]	960754 IMAGE Consortium	AA549137

4366 E	8 0219AMOM	Mus musculus F2 alpha prostoglandin regulatory protein (FPRP) mRNA, complete cds	963321 IMAGE Consortium	AA671763
4367 E	10 0219AMOM	ubiquitin-conjugating enzyme 7	961400 IMAGE Consortium	AA553292
4368 E	12 0219AMOM	ESTs, Weakly similar to coiled-coil like protein 1 [M.musculus]	1003046 IMAGE Consortium	AA684475
4369 G	2 0219AMOM	ESTs, Moderately similar to T20D3.3 [C.elegans]	960300 IMAGE Consortium	AA549726
4370 G	4 0219AMOM			
4371 G	6 0219AMOM	ESTs	960785 IMAGE Consortium	AA549162
4372 G	8 0219AMOM	ESTs, Highly similar to PTER_MOUSE PHOSPHOTRIESTERASE RELATED PROTEIN [M.musculus]	963362 IMAGE Consortium	AA671947
4373 G	10 0219AMOM	ESTs	961427 IMAGE Consortium	AA549180
4374 G	12 0219AMOM	ESTs	1003049 IMAGE Consortium	AA684484
4375 A	2 021GAMON	DNA segment, Chr 12, ERATO Doi 522, expressed	1003218 IMAGE Consortium	AA684094
4376 A	4 021GAMON	ESTs, Moderately similar to stromelysin PDGF responsive element binding protein transcription factor [M.musculus]	1003439 IMAGE Consortium	AA684253
4377 A	6 021GAMON	epsin 2	1003613 IMAGE Consortium	AA683707
4378 A	8 021GAMON	ESTs, Highly similar to similar to human DNA-binding protein 5. [H.sapiens]	976455 IMAGE Consortium	AA619869
4379 A	10 021GAMON	ESTs, Moderately similar to AC008015_2 unknown [H.sapiens]	976525 IMAGE Consortium	AA619976
4380 A	12 021GAMON	short coiled coil protein	976680 IMAGE Consortium	AA619804
4381 C	2 021GAMON	ESTs, Weakly similar to ADP-RIBOSYLATION FACTOR 2 [Saccharomyces cerevisiae]	1003298 IMAGE Consortium	AA666795
4382 C	4 021GAMON	ESTs, Weakly similar to AF145613_1 BcDNA.GH03108 [D.melanogaster]	1003448 IMAGE Consortium	AA684257
4383 C	6 021GAMON	ESTs, Moderately similar to ATP-DEPENDENT RNA HELICASE A [Homo sapiens]	1002906 IMAGE Consortium	AA792016
4384 C	8 021GAMON			
4385 C	10 021GAMON	B-cell translocation gene 4	976591 IMAGE Consortium	AA619728
4386 C	12 021GAMON	ribose 5-phosphate isomerase A	976685 IMAGE Consortium	AA619811
4387 E	2 021GAMON	ESTs	1003334 IMAGE Consortium	AA666811
4388 E	4 021GAMON	activin receptor interacting protein 1	1003468 IMAGE Consortium	AA684320
4389 E	6 021GAMON	myeloid differentiation primary response gene 116	1002997 IMAGE Consortium	AA684447
4390 E	8 021GAMON	ESTs	976471 IMAGE Consortium	AA619885
4391 E	10 021GAMON	diazepam binding inhibitor	976602 IMAGE Consortium	AA619894
4392 E	12 021GAMON	ESTs	976690 IMAGE Consortium	AA619807
4393 G	2 021GAMON	keratin complex 1, acidic, gene 12	1003372 IMAGE Consortium	AA684158
4394 G	4 021GAMON	ESTs, Moderately similar to ZFP265 [M.musculus]	1003473 IMAGE Consortium	AA684332
4395 G	6 021GAMON	ESTs, Weakly similar to hepatoma-derived growth factor [M.musculus]	976655 IMAGE Consortium	AA619907
4396 G	8 021GAMON	transforming growth factor, beta 2	976483 IMAGE Consortium	AA619890
4397 G	10 021GAMON	ESTs, Weakly similar to nuclear dual-specificity phosphatase [H.sapiens]	976620 IMAGE Consortium	AA619736



4398 G	12 021GAMON	ESTs, Moderately similar to TDT_MOUSE DNA NUCLEOTIDYLEXOTRANSFERASE [M.musculus]	976697 IMAGE Consortium	AA619915
4399 A	2 021NAMOO	ESTs	976735 IMAGE Consortium	AA619838
4400 A	4 021NAMOO	ESTs, Weakly similar to K02B2.3 gene product [C.elegans]	1024964 IMAGE Consortium	AI647202
4401 A	6 021NAMOO	radixin	1003912 IMAGE Consortium	AA674571
4402 A	8 021NAMOO	ESTs	1026070 IMAGE Consortium	AA616357
4403 A	10 021NAMOO	ESTs	1050526 IMAGE Consortium	AA596866
4404 A	12 021NAMOO	ESTs	1052377 IMAGE Consortium	AA839319
4405 C	2 021NAMOO	ESTs, Moderately similar to AF151066_1 HSPC232 [H.sapiens]	976739 IMAGE Consortium	AA619840
4406 C	4 021NAMOO	Mus musculus major histocompatibility locus class III region:butyrophilin-like protein gene, partial cds; Notch4, PBX2, RAGE, lysophatidic acid acyl transferase-alpha, palmitoyl-protein thioesterase 2 (PPT2), CREB-RP, and tenascin	1026625 IMAGE Consortium	AA792170
4407 C	6 021NAMOO	DNA segment, Chr 9, ERATO Doi 85, expressed	1003927 IMAGE Consortium	AA674596
4408 C	8 021NAMOO	ESTs	1025411 IMAGE Consortium	AI647223
4409 C	10 021NAMOO	Bcl2-like 2	1050567 IMAGE Consortium	AA596919
4410 C	12 021NAMOO	ESTs	1050937 IMAGE Consortium	AA592781
4411 E	2 021NAMOO	ubiquitin conjugating enzyme 2e	976746 IMAGE Consortium	AA619927
4412 E	4 021NAMOO	early growth response 1	1004005 IMAGE Consortium	AA674784
4413 E	6 021NAMOO	ESTs	1003961 IMAGE Consortium	AA674713
4414 E	8 021NAMOO	ESTs	1025460 IMAGE Consortium	AI647224
4415 E	10 021NAMOO	stromal interaction molecule 1	1051314 IMAGE Consortium	AI429023
4416 E	12 021NAMOO	chloride channel calcium activated 3	1050998 IMAGE Consortium	AA596289
4417 G	2 021NAMOO	ESTs, Highly similar to UNR PROTEIN [Rattus norvegicus]	976760 IMAGE Consortium	AA619844
4418 G	4 021NAMOO	ESTs	1003535 IMAGE Consortium	AA684418
4419 G	6 021NAMOO	COP9 (constitutive photomorphogenic), subunit 6 (Arabidopsis)	1003985 IMAGE Consortium	AA674732
4420 G	8 021NAMOO	hypothetical protein MNCb-2146	1025469 IMAGE Consortium	AA590700
4421 G	10 021NAMOO	Mus musculus GluR-delta2 specific interacting protein (delta2sip) mRNA, complete cds	1051374 IMAGE Consortium	AI429029
4422 G	12 021NAMOO	ESTs	1053134 IMAGE Consortium	AI594010
4423 A	2 021UAMOP	ESTs	1052650 IMAGE Consortium	AA606291
4424 A	4 021UAMOP	Mus musculus putative CAMP protein (Camp) mRNA, complete cds	1053703 IMAGE Consortium	AI594093
4425 A	6 021UAMOP	ESTs	1067990 IMAGE Consortium	AI552435
4426 A	8 021UAMOP	ESTs	1077800 IMAGE Consortium	AA796471
4427 A	10 021UAMOP	ESTs	1067560 IMAGE Consortium	AA647687
4428 A	12 021UAMOP	ESTs	1066068 IMAGE Consortium	AA673963
4429 C	2 021UAMOP	ESTs	1053444 IMAGE Consortium	AA611409

4430 C	4 021UAMOP	ESTs	1053768 IMAGE Consortium	AA638585
4431 C	6 021UAMOP	interferon-stimulated protein (15 kDa)	1068714 IMAGE Consortium	AA796818
4432 C	8 021UAMOP	ESTs	1077867 IMAGE Consortium	AA796749
4433 C	10 021UAMOP	ESTs	1066981 IMAGE Consortium	AA616189
4434 C	12 021UAMOP	ESTs	1066093 IMAGE Consortium	AA673996
4435 E	2 021UAMOP	phospholipase A2, group X	1053472 IMAGE Consortium	AA611431
4436 E	4 021UAMOP	ESTs	1053922 IMAGE Consortium	AI594110
4437 E	6 021UAMOP	tetranectin (plasminogen-binding protein)	1068748 IMAGE Consortium	AA796641
4438 E	8 021UAMOP	ESTs, Moderately similar to non-receptor tyrosine kinase [H.sapiens]	1077977 IMAGE Consortium	AA815521
4439 E	10 021UAMOP	ESTs	1054879 IMAGE Consortium	AA796126
4440 E	12 021UAMOP	ESTs	1066097 IMAGE Consortium	AA673998
4441 G	2 021UAMOP	ESTs, Moderately similar to unknown [H.sapiens]	1053487 IMAGE Consortium	AA608451
4442 G	4 021UAMOP	ESTs	1053946 IMAGE Consortium	AI594112
4443 G	6 021UAMOP	ESTs	1078210 IMAGE Consortium	AA822922
4444 G	8 021UAMOP	ESTs	1078839 IMAGE Consortium	AI593601
4445 G	10 021UAMOP	ESTs	1064358 IMAGE Consortium	AA615370
4446 G	12 021UAMOP	blood vessel epicardial substance	1065391 IMAGE Consortium	AA615786
4447 A	2 0211AMOQ	Public domain EST	1065627 IMAGE Consortium	
4448 A	4 0211AMOQ	ESTs	1065920 IMAGE Consortium	AI428995
4449 A	6 0211AMOQ	ESTs	1094366 IMAGE Consortium	AA869784
4450 A	8 0211AMOQ	ets homologous factor	1079266 IMAGE Consortium	AI593634
4451 A	10 0211AMOQ	ESTs, Highly similar to GMP REDUCTASE [Homo sapiens]	1095799 IMAGE Consortium	AI641814
4452 A	12 0211AMOQ	ESTs	400263 IMAGE Consortium	W77529
4453 C	2 0211AMOQ	Public domain EST	1067122 IMAGE Consortium	
4454 C	4 0211AMOQ	ESTs	1065935 IMAGE Consortium	AA611769
4455 C	6 0211AMOQ	ESTs, Highly similar to unnamed protein product [H.sapiens]	1079107 IMAGE Consortium	AI593615
4456 C	8 0211AMOQ	Mus musculus anti-HIV-1 reverse transcriptase single-chain variable fragment mRNA, complete cds	1096032 IMAGE Consortium	AA871265
4457 C	10 0211AMOQ	small inducible cytokine B subfamily, member 5	1095873 IMAGE Consortium	AA871143
4458 C	12 0211AMOQ	homeo box D9	400429 IMAGE Consortium	AI893954
4459 E	2 0211AMOQ	ESTs	1066446 IMAGE Consortium	AA615887
4460 E	4 0211AMOQ	carbonic anhydrase 3	1065954 IMAGE Consortium	AA611779
4461 E	6 0211AMOQ	ESTs, Weakly similar to NED4_MOUSE NEDD-4 PROTEIN [M.musculus]	1079125 IMAGE Consortium	AA823450
4462 E	8 0211AMOQ	ESTs	1095313 IMAGE Consortium	AA870728
4463 E	10 0211AMOQ	ESTs, Highly similar to R29515_1 [H.sapiens]	1096631 IMAGE Consortium	AA871670
4464 E	12 0211AMOQ	src homology 2 domain-containing transforming protein D	401189 IMAGE Consortium	AI425730
4465 G	2 0211AMOQ	ESTs	1066455 IMAGE Consortium	AI594395
4466 G	4 0211AMOQ	ESTs	1065990 IMAGE Consortium	AA611946
4467 G	6 0211AMOQ	ESTs, Highly similar to Y143_HUMAN HYPOTHETICAL	1095167 IMAGE Consortium	AI591660

		PROTEIN KIAA0143 [H.sapiens]		
4468 G	8 0211AMOQ	indole 2,3-dioxygenase	1096083 IMAGE Consortium	AA871317
4469 G	10 0211AMOQ	ESTs	1096699 IMAGE Consortium	AA871719
4470 G	12 0211AMOQ	ESTs	400526 IMAGE Consortium	AI425707
4471 A	2 0218AMOR	major urinary protein 3	401418 IMAGE Consortium	AI425738
4472 A	4 0218AMOR	ESTs, Weakly similar to apoptosis signal-regulating kinase 2 [M.musculus]	389745 IMAGE Consortium	W71463
4473 A	6 0218AMOR	ESTs, Moderately similar to unnamed protein product [H.sapiens]	400087 IMAGE Consortium	AI425685
4474 A	8 0218AMOR	inositol-requiring 1 (yeast)	1106865 IMAGE Consortium	AA755189
4475 A	10 0218AMOR	ESTs, Highly similar to PROBABLE UBIQUITIN CARBOXYL-TERMINAL HYDROLASE [Mus musculus]	1107730 IMAGE Consortium	AA623983
4476 A	12 0218AMOR	ESTs	1125347 IMAGE Consortium	AA675645
4477 C	2 0218AMOR	ESTs, Weakly similar to RETINAL-CADHERIN PRECURSOR [Mus musculus]	400759 IMAGE Consortium	AI425726
4478 C	4 0218AMOR	a disintegrin-like and metalloprotease (reprolysin type) with thrombospondin type 1 motif, 2	389987 IMAGE Consortium	W71229
4479 C	6 0218AMOR	ESTs, Weakly similar to RADIAL SPOKE PROTEIN 3 [Chlamydomonas reinhardtii]	1105806 IMAGE Consortium	AA674842
4480 C	8 0218AMOR	kinesin-like 1	1107676 IMAGE Consortium	AA623924
4481 C	10 0218AMOR	Fas apoptotic inhibitory molecule	1107754 IMAGE Consortium	AA623998
4482 C	12 0218AMOR	ESTs	1125088 IMAGE Consortium	AA693262
4483 E	2 0218AMOR	Public domain EST	389562 IMAGE Consortium	
4484 E	4 0218AMOR	ESTs	401932 IMAGE Consortium	AI425755
4485 E	6 0218AMOR	ESTs, Highly similar to histone acetyltransferase [H.sapiens]	1108039 IMAGE Consortium	AA623916
4486 E	8 0218AMOR	speckle-type POZ protein	1107745 IMAGE Consortium	AA624005
4487 E	10 0218AMOR	stromal antigen 2	1107828 IMAGE Consortium	AA623650
4488 E	12 0218AMOR	ESTs, Weakly similar to gag {provirus} [M.musculus]	1125234 IMAGE Consortium	AA693177
4489 G	2 0218AMOR	five SH3 domains	402206 IMAGE Consortium	AI425766
4490 G	4 0218AMOR	ESTs	390092 IMAGE Consortium	AI430171
4491 G	6 0218AMOR	ESTs	1106617 IMAGE Consortium	AA929129
4492 G	8 0218AMOR	ESTs	1107774 IMAGE Consortium	AA624059
4493 G	10 0218AMOR	ESTs, Weakly similar to major sperm fibrous sheath protein Pro-mAKAP82 [M.musculus]	1107850 IMAGE Consortium	AA623665
4494 G	12 0218AMOR	ESTs, Weakly similar to GENERAL NEGATIVE REGULATOR OF TRANSCRIPTION SUBUNIT 2 [Saccharomyces cerevisiae]	1126020 IMAGE Consortium	AA691036
4495 A	2 021FAMOS	ESTs, Weakly similar to HYPOTHETICAL 34.5 KD PROTEIN IN STE50-HIS4 INTERGENIC REGION [Saccharomyces cerevisiae]	1134268 IMAGE Consortium	AA675516
4496 A	4 021FAMOS	ESTs	1134484 IMAGE Consortium	AA681110
4497 A	6 021FAMOS	ESTs	1136125 IMAGE Consortium	AI592831
4498 A	8 021FAMOS	ESTs	1137337 IMAGE Consortium	AA688898

4499 A	10 021FAMOS	ESTs	1138294 IMAGE Consortium	AA821278
4500 A	12 021FAMOS	ESTs	1137898 IMAGE Consortium	AA691363
4501 C	2 021FAMOS	ESTs, Weakly similar to R27328_1 [H.sapiens]	1125575 IMAGE Consortium	AA675296
4502 C	4 021FAMOS	G protein-coupled receptor kinase 5	1134494 IMAGE Consortium	AA681115
4503 C	6 021FAMOS	ESTs	1136183 IMAGE Consortium	AI592835
4504 C	8 021FAMOS	ESTs	1153393 IMAGE Consortium	AA792612
4505 C	10 021FAMOS	ESTs	1138306 IMAGE Consortium	AA821291
4506 C	12 021FAMOS	ESTs, Weakly similar to KIAA0167 protein [H.sapiens]	1139366 IMAGE Consortium	AA667334
4507 E	2 021FAMOS	ESTs, Highly similar to POLYADENYLATE-BINDING PROTEIN [Homo sapiens]	1134300 IMAGE Consortium	AA675536
4508 E	4 021FAMOS	ESTs	1125792 IMAGE Consortium	AA691094
4509 E	6 021FAMOS	matrix metalloproteinase 7	1137047 IMAGE Consortium	AA689037
4510 E	8 021FAMOS	ESTs	1136697 IMAGE Consortium	AA683963
4511 E	10 021FAMOS	ESTs, Weakly similar to S34198 IgE Fc receptor II, low-affinity - rat [R.norvegicus]	1139077 IMAGE Consortium	AA657106
4512 E	12 021FAMOS	estrogen related receptor, alpha	1137956 IMAGE Consortium	AA691441
4513 G	2 021FAMOS	ESTs	1134306 IMAGE Consortium	AA675538
4514 G	4 021FAMOS	Mus musculus ancient conserved domain protein 2 (Acdp2) mRNA, complete cds	1125813 IMAGE Consortium	AA691105
4515 G	6 021FAMOS	ESTs, Moderately similar to GTPASE-ACTIVATING PROTEIN 1 [Drosophila melanogaster]	1153147 IMAGE Consortium	AA644997
4516 G	8 021FAMOS	ESTs	1138112 IMAGE Consortium	AA691720
4517 G	10 021FAMOS	ESTs, Weakly similar to actin-binding protein [M.musculus]	1138417 IMAGE Consortium	AA691628
4518 G	12 021FAMOS	ESTs	1138703 IMAGE Consortium	AA734158
4519 A	2 021MAMOT	ESTs, Weakly similar to 1915177A myosin:SUBUNIT	1166149 IMAGE Consortium	AA710484
4520 A	4 021MAMOT	thrombospondin 2	1154002 IMAGE Consortium	AA796401
4521 A	6 021MAMOT	protein kinase, cGMP-dependent, type II	1162639 IMAGE Consortium	AA771678
4522 A	8 021MAMOT	ESTs	1162977 IMAGE Consortium	AA770942
4523 A	10 021MAMOT	carboxypeptidase X2	1195037 IMAGE Consortium	AA690480
4524 A	12 021MAMOT	ESTs	1195467 IMAGE Consortium	AA690737
4525 C	2 021MAMOT	ESTs, Weakly similar to A56539 giantin - human [H.sapiens]	1166177 IMAGE Consortium	AA710506
4526 C	4 021MAMOT	slit homolog 2 (Drosophila)	1162253 IMAGE Consortium	AA691931
4527 C	6 021MAMOT	ESTs	1161954 IMAGE Consortium	AI642419
4528 C	8 021MAMOT	ESTs	1165109 IMAGE Consortium	AA709668
4529 C	10 021MAMOT	ESTs	1195084 IMAGE Consortium	AA690560
4530 C	12 021MAMOT	ESTs	1195565 IMAGE Consortium	AA711687
4531 E	2 021MAMOT	ESTs	1167044 IMAGE Consortium	AA711170
4532 E	4 021MAMOT			
4533 E	6 021MAMOT	calcium channel, voltage-dependent, gamma subunit 1	1161978 IMAGE Consortium	AA815611
4534 E	8 021MAMOT	ESTs, Moderately similar to hypothetical protein [H.sapiens]	1165151 IMAGE Consortium	AA709999

4535 E	10 021MAMOT	synuclein, gamma	1195168 IMAGE Consortium	AA690669
4536 E	12 021MAMOT	small inducible cytokine A11	1195601 IMAGE Consortium	AA711712
4537 G	2 021MAMOT	ESTs	1165673 IMAGE Consortium	AA710215
4538 G	4 021MAMOT	ESTs	1154283 IMAGE Consortium	AA796234
4539 G	6 021MAMOT	lymphocyte-activation gene 3	1164166 IMAGE Consortium	AA690798
4540 G	8 021MAMOT	ESTs	1165261 IMAGE Consortium	AA709576
4541 G	10 021MAMOT	Mus musculus mRNA for L-specific multifunctional beta-oxidation protein, partial CDS	1195248 IMAGE Consortium	AA718155
4542 G	12 021MAMOT	ESTs	1195630 IMAGE Consortium	AA711758
4543 A	2 021TAMOU	G protein-coupled receptor 83	386074 IMAGE Consortium	W62391
4544 A	4 021TAMOU	syntrophin, acidic 1	373271 IMAGE Consortium	W57275
4545 A	6 021TAMOU	ESTs	374654 IMAGE Consortium	AI430460
4546 A	8 021TAMOU	ESTs	385625 IMAGE Consortium	AI430498
4547 A	10 021TAMOU	ESTs	1178160 IMAGE Consortium	AA718731
4548 A	12 021TAMOU	ESTs	1225036 IMAGE Consortium	AA756333
4549 C	2 021TAMOU	GLI-Kruppel family member GLI	386514 IMAGE Consortium	AI385860
4550 C	4 021TAMOU	ESTs, Weakly similar to RNA-binding protein [M.musculus]	374136 IMAGE Consortium	AI430099
4551 C	6 021TAMOU	ESTs	373937 IMAGE Consortium	AI430089
4552 C	8 021TAMOU	ESTs	386457 IMAGE Consortium	AI430575
4553 C	10 021TAMOU	ESTs, Weakly similar to similar to other protein phosphatases 1, 2A and 2B [C.elegans]	1178274 IMAGE Consortium	AA718881
4554 C	12 021TAMOU	ESTs, Weakly similar to unknown [H.sapiens]	1225044 IMAGE Consortium	AA756337
4555 E	2 021TAMOU	Mus musculus mRNA for S-1, complete cds	386767 IMAGE Consortium	AI430599
4556 E	4 021TAMOU	nuclear factor, erythroid derived 2, 45 kDa	373510 IMAGE Consortium	W59072
4557 E	6 021TAMOU	ESTs, Highly similar to LAMBDA-CRYSTALLIN [Oryctolagus cuniculus]	374698 IMAGE Consortium	AI385417
4558 E	8 021TAMOU	ESTs	385743 IMAGE Consortium	W62237
4559 E	10 021TAMOU	solute carrier family (organic cation transporter), member 4	1177710 IMAGE Consortium	AA718357
4560 E	12 021TAMOU	ESTs	1225070 IMAGE Consortium	AA756360
4561 G	2 021TAMOU	ESTs, Weakly similar to ORF YOL102c [S.cerevisiae]	386948 IMAGE Consortium	AI430610
4562 G	4 021TAMOU	surfeit gene 1	374298 IMAGE Consortium	W65212
4563 G	6 021TAMOU	neuron specific gene family member 2	374991 IMAGE Consortium	W62007
4564 G	8 021TAMOU	ESTs	386516 IMAGE Consortium	AI430582
4565 G	10 021TAMOU	ESTs	1177743 IMAGE Consortium	AA718388
4566 G	12 021TAMOU	citron	1225131 IMAGE Consortium	AA762879
4567 A	2 0210AMOV	3-phosphoinositide dependent protein kinase-1	1225338 IMAGE Consortium	AA764115
4568 A	4 0210AMOV	ESTs	1195689 IMAGE Consortium	AA711863
4569 A	6 0210AMOV	extracellular proteinase inhibitor	1195775 IMAGE Consortium	AA712002
4570 A	8 0210AMOV	ESTs	1195947 IMAGE Consortium	AA716838
4571 A	10 0210AMOV	ESTs	1230674 IMAGE Consortium	AA880332
4572 A	12 0210AMOV	ESTs	1226377 IMAGE Consortium	AA940445

4573 C	2 0210AMOV	ESTs	1196021 IMAGE Consortium	AA716895
4574 C	4 0210AMOV	sialoadhesin	1195699 IMAGE Consortium	AA711868
4575 C	6 0210AMOV	ESTs	1225859 IMAGE Consortium	AA739203
4576 C	8 0210AMOV	ESTs, Weakly similar to transin-1 [M.musculus]	1195964 IMAGE Consortium	AA716836
4577 C	10 0210AMOV	ESTs	1230703 IMAGE Consortium	AA879931
4578 C	12 0210AMOV	ESTs	1227140 IMAGE Consortium	AI425119
4579 E	2 0210AMOV	ESTs	1225428 IMAGE Consortium	AA762068
4580 E	4 0210AMOV	ESTs	1225734 IMAGE Consortium	AI428744
4581 E	6 0210AMOV	RAD51 like 1 (S. cerevisiae)	1225890 IMAGE Consortium	AA739335
4582 E	8 0210AMOV	Mus musculus G7c protein (G7c) mRNA, G7c-d allele, complete cds	1230278 IMAGE Consortium	AA879522
4583 E	10 0210AMOV	laminin, alpha 5	1230752 IMAGE Consortium	AA879966
4584 E	12 0210AMOV	ESTs, Moderately similar to unnamed protein product [H.sapiens]	1243273 IMAGE Consortium	AA763980
4585 G	2 0210AMOV	Public domain EST	1226212 IMAGE Consortium	
4586 G	4 0210AMOV	ESTs	1225763 IMAGE Consortium	AA764259
4587 G	6 0210AMOV	Mus musculus interleukin 2 receptor mRNA, 3'UTR, partial sequence	1195854 IMAGE Consortium	AA716986
4588 G	8 0210AMOV	ESTs	1230347 IMAGE Consortium	AA879701
4589 G	10 0210AMOV	ESTs, Weakly similar to KIAA0377 [H.sapiens]	1230771 IMAGE Consortium	AA880366
4590 G	12 0210AMOV	ESTs, Weakly similar to rho GTPase activating protein 5 [M.musculus]	1226537 IMAGE Consortium	AI466146
4591 A	2 0217AMOW	ESTs	1226720 IMAGE Consortium	AA738602
4592 A	4 0217AMOW	ESTs	1243994 IMAGE Consortium	AA790279
4593 A	6 0217AMOW	ESTs	1263413 IMAGE Consortium	AA874216
4594 A	8 0217AMOW	ESTs	1262894 IMAGE Consortium	AA839604
4595 A	10 0217AMOW	ESTs	1245566 IMAGE Consortium	AA797884
4596 A	12 0217AMOW	ESTs	1245778 IMAGE Consortium	AA822744
4597 C	2 0217AMOW	ESTs	1226755 IMAGE Consortium	AA738637
4598 C	4 0217AMOW	ESTs	1263096 IMAGE Consortium	AA856323
4599 C	6 0217AMOW	ESTs	1248145 IMAGE Consortium	AA762401
4600 C	8 0217AMOW	ESTs	1247611 IMAGE Consortium	AI464994
4601 C	10 0217AMOW	ESTs, Weakly similar to ORF2 [M.musculus]	1245570 IMAGE Consortium	AA797886
4602 C	12 0217AMOW	ESTs	1247240 IMAGE Consortium	AA839453
4603 E	2 0217AMOW	ESTs	1226854 IMAGE Consortium	AA739088
4604 E	4 0217AMOW	ESTs	1263107 IMAGE Consortium	AA856337
4605 E	6 0217AMOW	ESTs, Weakly similar to KIAA0769 protein [H.sapiens]	1248160 IMAGE Consortium	AA762397
4606 E	8 0217AMOW	ESTs	1248365 IMAGE Consortium	AA789636
4607 E	10 0217AMOW	ESTs, Highly similar to ANPC_MOUSE ATRIAL NATRIURETIC PEPTIDE CLEARANCE RECEPTOR PRECURSOR [M.musculus]	1245631 IMAGE Consortium	AA797808
4608 E	12 0217AMOW	ESTs, Highly similar to MYOSIN LIGHT CHAIN KINASE [Dictyostelium discoideum]	1246549 IMAGE Consortium	AA832562

4609 G	2	0217AMOW	ESTs	1244328	IMAGE Consortium	AA790831
4610 G	4	0217AMOW	protein kinase C, zeta	1263199	IMAGE Consortium	AA856421
4611 G	6	0217AMOW	lactotransferrin	1248207	IMAGE Consortium	AA763033
4612 G	8	0217AMOW	ESTs	1247849	IMAGE Consortium	AA959706
4613 G	10	0217AMOW	ESTs	1244946	IMAGE Consortium	AA797632
4614 G	12	0217AMOW	ESTs, Highly similar to RER1 PROTEIN [Saccharomyces cerevisiae]	1245848	IMAGE Consortium	AA822839
4615 A	2	021EAMOX	CD1d1 antigen	1247449	IMAGE Consortium	AA839844
4616 A	4	021EAMOX	ESTs	1244627	IMAGE Consortium	AA797242
4617 A	6	021EAMOX	chemokine orphan receptor 1	1263525	IMAGE Consortium	AA874277
4618 A	8	021EAMOX	ESTs	1265145	IMAGE Consortium	AA880140
4619 A	10	021EAMOX	ESTs, Highly similar to AF173003_1 apoptosis regulator [H.sapiens]	1264612	IMAGE Consortium	AI509629
4620 A	12	021EAMOX	ESTs	1265525	IMAGE Consortium	AA867015
4621 C	2	021EAMOX	thymine glycol DNA glycosylase/AP lyase	1246791	IMAGE Consortium	AA832726
4622 C	4	021EAMOX	ets homologous factor	1244672	IMAGE Consortium	AA797272
4623 C	6	021EAMOX	coatomer protein complex, subunit zeta 1	1264287	IMAGE Consortium	AA881328
4624 C	8	021EAMOX	ESTs	1263726	IMAGE Consortium	AA863680
4625 C	10	021EAMOX	ESTs, Weakly similar to KIAA0328 protein [H.sapiens]	1265355	IMAGE Consortium	AA867162
4626 C	12	021EAMOX	ESTs	1264850	IMAGE Consortium	AI509655
4627 E	2	021EAMOX	ESTs	1244062	IMAGE Consortium	AA790309
4628 E	4	021EAMOX	ESTs, Weakly similar to bA395L14.2 [H.sapiens]	1245478	IMAGE Consortium	AA797865
4629 E	6	021EAMOX	ESTs	1265018	IMAGE Consortium	AA881583
4630 E	8	021EAMOX	ESTs	1264464	IMAGE Consortium	AI509612
4631 E	10	021EAMOX	ESTs	1264641	IMAGE Consortium	AI509636
4632 E	12	021EAMOX	ESTs	1264879	IMAGE Consortium	AI509666
4633 G	2	021EAMOX	ESTs	1245121	IMAGE Consortium	AA797517
4634 G	4	021EAMOX	ESTs	1263254	IMAGE Consortium	AI509561
4635 G	6	021EAMOX	ESTs, Highly similar to T31081 cca3 protein - rat [R.norvegicus]	1264304	IMAGE Consortium	AI509582
4636 G	8	021EAMOX	ESTs	1264483	IMAGE Consortium	AA881375
4637 G	10	021EAMOX	ESTs, Moderately similar to KIAA1495 protein [H.sapiens]	1265380	IMAGE Consortium	AA867173
4638 G	12	021EAMOX	ESTs	1264892	IMAGE Consortium	AI509664
4639 A	2	021LAMOY	tumor necrosis factor	1327679	IMAGE Consortium	AA920586
4640 A	4	021LAMOY				
4641 A	6	021LAMOY	ESTs, Highly similar to hypothetical protein [H.sapiens]	1379599	IMAGE Consortium	AI061004
4642 A	8	021LAMOY	tachykinin 1	1380603	IMAGE Consortium	AI060847
4643 A	10	021LAMOY	Mouse rearranged T-cell receptor beta V14/D1.1/J2.3 gene segment	1365841	IMAGE Consortium	AI020755
4644 A	12	021LAMOY	ESTs	1344869	IMAGE Consortium	AA939711
4645 C	2	021LAMOY	ESTs	1281297	IMAGE Consortium	AA874599
4646 C	4	021LAMOY	islet amyloid polypeptide	1379049	IMAGE Consortium	AI050257
4647 C	6	021LAMOY	ESTs, Highly similar to N155_RAT NUCLEAR PORE	1379662	IMAGE Consortium	AI060336

4648 C	8 021LAMOY	COMPLEX PROTEIN NUP155 [R.norvegicus] Mus musculus mRNA for thrombospondin type 1 domain, complete cds	1365431 IMAGE Consortium	AI020649
4649 C	10 021LAMOY	ESTs	1365968 IMAGE Consortium	AI021230
4650 C	12 021LAMOY	Mouse (clone RAD11) T-cell receptor rearranged delta- chain mRNA V-D-J-C region, 3' end	1345654 IMAGE Consortium	AI154374
4651 E	2 021LAMOY	ESTs, Moderately similar to AF090915_1 PRO0310p1 [H.sapiens]	1282072 IMAGE Consortium	AA920753
4652 E	4 021LAMOY	ESTs	1378442 IMAGE Consortium	AI019266
4653 E	6 021LAMOY	choline/ethanolamine kinase	1379732 IMAGE Consortium	AI049122
4654 E	8 021LAMOY	Mus musculus toll-like receptor 1 (Tlr1) mRNA, complete cds	1364755 IMAGE Consortium	AI019567
4655 E	10 021LAMOY	ESTs	1361208 IMAGE Consortium	AA982338
4656 E	12 021LAMOY	nuclear receptor coactivator 2	1344985 IMAGE Consortium	AA939852
4657 G	2 021LAMOY	ESTs	1281381 IMAGE Consortium	AA874187
4658 G	4 021LAMOY	amyloid beta (A4) precursor protein	1379230 IMAGE Consortium	AI050367
4659 G	6 021LAMOY	ESTs	1379891 IMAGE Consortium	AI049443
4660 G	8 021LAMOY	Public domain EST	1364786 IMAGE Consortium	
4661 G	10 021LAMOY	vascular endothelial zinc finger 1	1344669 IMAGE Consortium	AA939736
4662 G	12 021LAMOY	prostaglandin E receptor EP2 subtype	1347203 IMAGE Consortium	AA958901
4663 A	2 021SAMOZ	ESTs, Highly similar to KIAA0737 protein [H.sapiens]	1347536 IMAGE Consortium	AA981560
4664 A	4 021SAMOZ	ESTs, Weakly similar to T17285 hypothetical protein DKFZp434N0535.1 - human [H.sapiens]	1382098 IMAGE Consortium	AA797957
4665 A	6 021SAMOZ	ESTs	1382407 IMAGE Consortium	AI046527
4666 A	8 021SAMOZ	S100 calcium-binding protein A4	1383390 IMAGE Consortium	AI120885
4667 A	10 021SAMOZ	ESTs	403209 IMAGE Consortium	W81888
4668 A	12 021SAMOZ	Mus musculus Blu mRNA, partial sequence	404450 IMAGE Consortium	AI425321
4669 C	2 021SAMOZ	ESTs	1346997 IMAGE Consortium	AA958733
4670 C	4 021SAMOZ	ESTs	1382121 IMAGE Consortium	AA797990
4671 C	6 021SAMOZ	Sh3 domain YSC-like 1	1382454 IMAGE Consortium	AI036131
4672 C	8 021SAMOZ	mast cell protease 5	1381984 IMAGE Consortium	AA832859
4673 C	10 021SAMOZ	ESTs, Moderately similar to AF161427_1 HSPC309 [H.sapiens]	404184 IMAGE Consortium	AI425306
4674 C	12 021SAMOZ	G-protein coupled receptor 26	404566 IMAGE Consortium	AI425329
4675 E	2 021SAMOZ	regulator of G-protein signaling 14	1348773 IMAGE Consortium	AA981480
4676 E	4 021SAMOZ	ESTs	1382145 IMAGE Consortium	AA798212
4677 E	6 021SAMOZ	decay accelerating factor 1	1383200 IMAGE Consortium	AI120685
4678 E	8 021SAMOZ	guanine nucleotide regulatory protein (oncogene)	1382740 IMAGE Consortium	AI036870
4679 E	10 021SAMOZ	ESTs	404222 IMAGE Consortium	W82565
4680 E	12 021SAMOZ	ESTs, Weakly similar to p190-B [M.musculus]	403917 IMAGE Consortium	W82301
4681 G	2 021SAMOZ	ESTs, Weakly similar to osmosis responsive factor [H.sapiens]	1349829 IMAGE Consortium	AA980896
4682 G	4 021SAMOZ	ESTs, Highly similar to hypothetical protein [H.sapiens]	1382226 IMAGE Consortium	AA798312



4683 G	6 021SAMOZ	CD1d2 antigen	1382522 IMAGE Consortium	AI036466
4684 G	8 021SAMOZ	ESTs, Moderately similar to AF119384_1 putative CAMP protein [M.musculus]	1382786 IMAGE Consortium	AI120198
4685 G	10 021SAMOZ	kinesin light chain 1	403578 IMAGE Consortium	W82109
4686 G	12 021SAMOZ	cellular retinoic acid binding protein I	406272 IMAGE Consortium	W83353
4687 A	2 021SAMP0	ESTs	406471 IMAGE Consortium	W83503
4688 A	4 021SAMP0	ESTs	406782 IMAGE Consortium	W83693
4689 A	6 021SAMP0	ESTs, Weakly similar to HYPOTHETICAL 13.4 KD PROTEIN IN ACS1-GCV3 INTERGENIC REGION [Saccharomyces cerevisiae]	1383858 IMAGE Consortium	AI181708
4690 A	8 021SAMP0	ESTs	1383094 IMAGE Consortium	AI120447
4691 A	10 021SAMP0	NADH ubiquinone oxidoreductase subunit MWFE	1383596 IMAGE Consortium	AI120987
4692 A	12 021SAMP0	ESTs, Weakly similar to similar to YBS4_YEAST [C.elegans]	1430976 IMAGE Consortium	AI180535
4693 C	2 021SAMP0	ESTs	407212 IMAGE Consortium	AA052596
4694 C	4 021SAMP0	matrix gamma-carboxyglutamate (gla) protein	407582 IMAGE Consortium	W88093
4695 C	6 021SAMP0	ESTs	1383865 IMAGE Consortium	AI181735
4696 C	8 021SAMP0	ESTs, Highly similar to similar to human ankyrin 1(S08275) [H.sapiens]	1383143 IMAGE Consortium	AI120792
4697 C	10 021SAMP0	UDP-N-acetyl-alpha-D-galactosamine:polypeptide N-acetylgalactosaminyltransferase 3	1383632 IMAGE Consortium	AI180761
4698 C	12 021SAMP0	ESTs	1434019 IMAGE Consortium	AA833196
4699 E	2 021SAMP0	cofilin 1, non-muscle	407267 IMAGE Consortium	W85144
4700 E	4 021SAMP0	flap structure specific endonuclease 1	407645 IMAGE Consortium	W89311
4701 E	6 021SAMP0	human immunodeficiency virus type I enhancer-binding protein 2	1383996 IMAGE Consortium	AI181636
4702 E	8 021SAMP0	peroxisomal biogenesis factor 11b	1383182 IMAGE Consortium	AI120667
4703 E	10 021SAMP0	serine/threonine kinase 19	1383646 IMAGE Consortium	AI180767
4704 E	12 021SAMP0	ESTs, Weakly similar to C1q-related factor [M.musculus]	1434214 IMAGE Consortium	AA833248
4705 G	2 021SAMP0	ESTs, Weakly similar to P1.11659 5 [H.sapiens]	406612 IMAGE Consortium	W83599
4706 G	4 021SAMP0	insulin-like growth factor binding protein 3	407674 IMAGE Consortium	W89951
4707 G	6 021SAMP0	ESTs	1396033 IMAGE Consortium	AI183059
4708 G	8 021SAMP0	ESTs, Moderately similar to cell growth regulator rCGR19 [R.norvegicus]	1383357 IMAGE Consortium	AI120861
4709 G	10 021SAMP0	ESTs, Weakly similar to open reading frame [M.musculus]	1383710 IMAGE Consortium	AI117381
4710 G	12 021SAMP0	ESTs, Highly similar to AF135493_1 neuronal apoptosis inhibitory protein [M.musculus]	1433663 IMAGE Consortium	AI606608
4711 A	2 021AMSI	Public domain EST	1434676 IMAGE Consortium	
4712 A	4 021AMSI	ESTs	1430029 IMAGE Consortium	AI122253
4713 A	6 021AMSI	ESTs	1400409 IMAGE Consortium	AI117905
4714 A	8 021AMSI	ESTs	1400935 IMAGE Consortium	AI121941
4715 A	10 021AMSI	ESTs	1449402 IMAGE Consortium	AI153921
4716 A	12 021AMSI	ESTs, Weakly similar to MUC1_MOUSE MUCIN 1	1449868 IMAGE Consortium	AI197496

		PRECURSOR [M.musculus]		
4717 C	2 0215AMSI	ESTs	1434779 IMAGE Consortium	AI157015
4718 C	4 0215AMSI	microtubule-associated protein 6	1430217 IMAGE Consortium	AI122516
4719 C	6 0215AMSI	ESTs	1400508 IMAGE Consortium	AI117918
4720 C	8 0215AMSI	angiogenin	1400985 IMAGE Consortium	AI121741
4721 C	10 0215AMSI	guanine nucleotide binding protein (G protein), gamma transducing activity polypeptide 2	1449477 IMAGE Consortium	AI154009
4722 C	12 0215AMSI	ESTs	1449999 IMAGE Consortium	AI197098
4723 E	2 0215AMSI	ESTs, Highly similar to snRNA activating protein complex 50kD subunit [H.sapiens]	1446022 IMAGE Consortium	AI122204
4724 E	4 0215AMSI	adaptor protein complex AP-1, gamma 2 subunit	1430269 IMAGE Consortium	AI181488
4725 E	6 0215AMSI	pleckstrin homology, Sec7 and coiled/coil domains 1	1400576 IMAGE Consortium	AI117872
4726 E	8 0215AMSI	zinc finger protein, subfamily 1A, 4	1429281 IMAGE Consortium	AI153405
4727 E	10 0215AMSI	Max dimerization protein	1448816 IMAGE Consortium	AI573509
4728 E	12 0215AMSI			
4729 G	2 0215AMSI	casein beta	1446079 IMAGE Consortium	AI157135
4730 G	4 0215AMSI	ESTs	1430393 IMAGE Consortium	AI182156
4731 G	6 0215AMSI	ESTs, Weakly similar to I48668 zinc finger protein 51 - mouse [M.musculus]	1398047 IMAGE Consortium	AI118879
4732 G	8 0215AMSI	SWI/SNF related, matrix associated, actin dependent regulator of chromatin, subfamily a, member 3	1429357 IMAGE Consortium	AI153451
4733 G	10 0215AMSI	ESTs	1448861 IMAGE Consortium	AI573559
4734 G	12 0215AMSI			
4735 A	2 0216AMP2	ESTs	1446455 IMAGE Consortium	AI152150
4736 A	4 0216AMP2	suppressor of white apricot homolog 2-pending	1447466 IMAGE Consortium	AI154285
4737 A	6 0216AMP2	Trf-proximal protein	1449285 IMAGE Consortium	AI153977
4738 A	8 0216AMP2	ESTs	1495072 IMAGE Consortium	AI157540
4739 A	10 0216AMP2	ESTs	1477859 IMAGE Consortium	AI155262
4740 A	12 0216AMP2			
4741 C	2 0216AMP2	ESTs	1447242 IMAGE Consortium	AI158475
4742 C	4 0216AMP2	sema domain, immunoglobulin domain (Ig), TM domain, and short cytoplasmic domain	1446785 IMAGE Consortium	AI158284
4743 C	6 0216AMP2			
4744 C	8 0216AMP2	ESTs, Moderately similar to DIAC_RAT DI-N-ACETYLCHITOBIASE PRECURSOR [R.norvegicus]	1477609 IMAGE Consortium	AI155401
4745 C	10 0216AMP2	syntrophin, basic 2	1494615 IMAGE Consortium	AI195906
4746 C	12 0216AMP2	DNA segment, Chr 14, ERATO Doi 453, expressed	1479937 IMAGE Consortium	AI156376
4747 E	2 0216AMP2	ESTs, Weakly similar to DRR1 [H.sapiens]	1447253 IMAGE Consortium	AI158650
4748 E	4 0216AMP2	immunoglobulin heavy chain 3 (serum IgG2b)	1448206 IMAGE Consortium	AI158823
4749 E	6 0216AMP2	ESTs, Weakly similar to A55929 zinc finger protein nocA - fruit fly [D.melanogaster]	1448677 IMAGE Consortium	AI528976
4750 E	8 0216AMP2	ESTs	1478343 IMAGE Consortium	AI152563
4751 E	10 0216AMP2	ring finger protein (C3HC4 type) 19	1494675 IMAGE Consortium	AI197154

4752 E	12 0216AMP2	cystic fibrosis transmembrane conductance regulator homolog	1480003 IMAGE Consortium	AI156185
4753 G	2 0216AMP2	small inducible cytokine B subfamily (Cys-X-Cys), member 10	1446589 IMAGE Consortium	AI158236
4754 G	4 0216AMP2	solute carrier family 4 (anion exchanger), member 1, adapter protein	1448227 IMAGE Consortium	AI173071
4755 G	6 0216AMP2	Mus musculus immunoglobulin scavenger receptor IgSR mRNA, complete cds	1477073 IMAGE Consortium	AI152359
4756 G	8 0216AMP2	ESTs, Highly similar to PROSTAGLANDIN-E2 9-REDUCTASE [Oryctolagus cuniculus]	1477668 IMAGE Consortium	AI155437
4757 G	10 0216AMP2	DNA segment, Chr 18, Abbott 1 expressed	1477949 IMAGE Consortium	AI155369
4758 G	12 0216AMP2	ESTs	1480121 IMAGE Consortium	AI156249
4759 A	2 021DAMP3	ESTs	1749761 IMAGE Consortium	AI047049
4760 A	4 021DAMP3	ESTs	1746306 IMAGE Consortium	AI037657
4761 A	6 021DAMP3	ESTs	1745931 IMAGE Consortium	AI036198
4762 A	8 021DAMP3	ESTs	1510962 IMAGE Consortium	AI227494
4763 A	10 021DAMP3	cadherin-related neuronal receptor 1	1497544 IMAGE Consortium	AI664063
4764 A	12 021DAMP3	ESTs	1497843 IMAGE Consortium	AI664340
4765 C	2 021DAMP3	postmeiotic segregation increased 2 (S. cerevisiae)	1750079 IMAGE Consortium	AI047244
4766 C	4 021DAMP3	ESTs, Weakly similar to fibrousheathin II [H.sapiens]	1746327 IMAGE Consortium	AI037620
4767 C	6 021DAMP3	ESTs	1745968 IMAGE Consortium	AI036204
4768 C	8 021DAMP3	ESTs	1496403 IMAGE Consortium	AI194684
4769 C	10 021DAMP3	OVO homolog-like 1 (Drosophila)	1496850 IMAGE Consortium	AI194439
4770 C	12 021DAMP3			
4771 E	2 021DAMP3	ESTs	1746029 IMAGE Consortium	AI036778
4772 E	4 021DAMP3	ESTs	1746347 IMAGE Consortium	AI037631
4773 E	6 021DAMP3	ESTs	1764194 IMAGE Consortium	AI047703
4774 E	8 021DAMP3	ESTs, Highly similar to elav-type RNA-binding protein [M.musculus]	1496492 IMAGE Consortium	AI197742
4775 E	10 021DAMP3	prolactin-like protein E	1497583 IMAGE Consortium	AI664122
4776 E	12 021DAMP3	brain link protein 1	1616912 IMAGE Consortium	AA967720
4777 G	2 021DAMP3	tsec-2	1746064 IMAGE Consortium	AI036790
4778 G	4 021DAMP3	ESTs, Weakly similar to NSG2_MOUSE NEURON SPECIFIC PROTEIN FAMILY MEMBER 2 [M.musculus]	1763135 IMAGE Consortium	AI132172
4779 G	6 021DAMP3	topoisomerase (DNA) III beta	1749087 IMAGE Consortium	AI047903
4780 G	8 021DAMP3	sulfotransferase family, cytosolic, 2B, member 1	1498071 IMAGE Consortium	AI225394
4781 G	10 021DAMP3	protein kinase C and casein kinase substrate in neurons 1	1617289 IMAGE Consortium	AA982442
4782 G	12 021DAMP3	protein tyrosine phosphatase, receptor type, Z	1617668 IMAGE Consortium	AA968121
4783 A	2 021KAMP4	sine oculis-related homeobox 4 homolog (Drosophila)	1764007 IMAGE Consortium	AI047561
4784 A	4 021KAMP4	ESTs	1763989 IMAGE Consortium	AI047548
4785 A	6 021KAMP4	Cbp/p300-interacting transactivator, with Glu/Asp-rich carboxy-terminal domain, 2	537128 IMAGE Consortium	AA117983
4786 A	8 021KAMP4	ESTs, Moderately similar to putative E1-E2 ATPase	537357 IMAGE Consortium	AA118074

		[M.musculus]		
4787 A	10 021KAMP4	ESTs	537627 IMAGE Consortium	AI550125
4788 A	12 021KAMP4	ESTs, Weakly similar to ANKYRIN, BRAIN VARIANT 2 [Homo sapiens]	538097 IMAGE Consortium	AI325781
4789 C	2 021KAMP4	ESTs	1852376 IMAGE Consortium	AI256607
4790 C	4 021KAMP4	ESTs	1764714 IMAGE Consortium	AI115600
4791 C	6 021KAMP4	ESTs, Highly similar to CASEIN KINASE I, DELTA ISOFORM [Homo sapiens]	537162 IMAGE Consortium	AI550091
4792 C	8 021KAMP4	vacuolar protein sorting 45 (yeast)	537364 IMAGE Consortium	AI324447
4793 C	10 021KAMP4			
4794 C	12 021KAMP4	ESTs	538102 IMAGE Consortium	AI550151
4795 E	2 021KAMP4	ESTs	1764197 IMAGE Consortium	AI047714
4796 E	4 021KAMP4	ESTs, Highly similar to KIAA1469 protein [H.sapiens]	1852949 IMAGE Consortium	AI227034
4797 E	6 021KAMP4	WD40 protein Ciao1	537182 IMAGE Consortium	AA117997
4798 E	8 021KAMP4	endothelial differentiation, lysophosphatidic acid G-protein-coupled receptor 4	537445 IMAGE Consortium	AI550109
4799 E	10 021KAMP4	ESTs, Moderately similar to ERYTHROCYTE BAND 7 INTEGRAL MEMBRANE PROTEIN [Homo sapiens]	536956 IMAGE Consortium	AI325758
4800 E	12 021KAMP4	FK506-binding protein 3 (25kD)	538114 IMAGE Consortium	AA120298
4801 G	2 021KAMP4	potassium voltage-gated channel, subfamily Q, member 1	1852438 IMAGE Consortium	AI789568
4802 G	4 021KAMP4			
4803 G	6 021KAMP4	Public domain EST	537201 IMAGE Consortium	
4804 G	8 021KAMP4	cyclin D1	537489 IMAGE Consortium	AA117547
4805 G	10 021KAMP4	ESTs, Weakly similar to Keap1 [M.musculus]	536961 IMAGE Consortium	AA114654
4806 G	12 021KAMP4	ESTs, Weakly similar to NEURAXIN [Rattus norvegicus]	538116 IMAGE Consortium	AI325784
4807 A	2 021RAMP5	DNA segment, Chr 16, ERATO Doi 454, expressed	538320 IMAGE Consortium	AA116792
4808 A	4 021RAMP5	MAP-1 protein	538578 IMAGE Consortium	AA117404
4809 A	6 021RAMP5	cystein-rich hydrophobic domain 1	538752 IMAGE Consortium	AA123458
4810 A	8 021RAMP5	Mouse mRNA for osteoblast specific factor 2 (OSF-2) {IMAGE:403071}	403071 IMAGE Consortium	W81878
4811 A	10 021RAMP5	collapsin response mediator protein 1	401570 IMAGE Consortium	W78255
4812 A	12 021RAMP5	procollagen, type I, alpha 1	402270 IMAGE Consortium	AI425767
4813 C	2 021RAMP5	alpha-spectrin 1, erythroid	538325 IMAGE Consortium	AA116814
4814 C	4 021RAMP5	ESTs	538588 IMAGE Consortium	AA116700
4815 C	6 021RAMP5	ESTs, Highly similar to similar to hypothetical protein L8167.6 of Saccharomyces cerevisiae. [H.sapiens]	538804 IMAGE Consortium	AA123485
4816 C	8 021RAMP5	Tuberous sclerosis 2 {IMAGE:404321}	404321 IMAGE Consortium	W82751
4817 C	10 021RAMP5	protease, cysteine, 1	403755 IMAGE Consortium	W83121
4818 C	12 021RAMP5	Insulin-like growth factor binding protein 5 {IMAGE:418952}	418952 IMAGE Consortium	W89253
4819 E	2 021RAMP5	ESTs, Highly similar to RABPHILIN-3A [Rattus norvegicus]	538371 IMAGE Consortium	AI325790
4820 E	4 021RAMP5	ESTs	538596 IMAGE Consortium	AA116846

4821 E	6 021RAMP5	laminin, alpha 1	538828 IMAGE Consortium	AA123168
4822 E	8 021RAMP5	ESTs {IMAGE:405779}	405779 IMAGE Consortium	W83025
4823 E	10 021RAMP5	X-ray repair complementing defective repair in Chinese hamster cells 5	404596 IMAGE Consortium	W83084
4824 E	12 021RAMP5	Mus musculus friend of GATA-1 (FOG) mRNA, complete cds {IMAGE:418539}	418539 IMAGE Consortium	W99029
4825 G	2 021RAMP5	Mouse adenine phosphoribosyltransferase (APRT), complete cds	538413 IMAGE Consortium	AA117239
4826 G	4 021RAMP5	ESTs, Highly similar to Sec22 homolog [R.norvegicus]	537899 IMAGE Consortium	AA119813
4827 G	6 021RAMP5	ESTs, Weakly similar to protein kinase [M.musculus] {IMAGE:933770}	933770 IMAGE Consortium	AA545690
4828 G	8 021RAMP5	transcription factor E2a	400175 IMAGE Consortium	W77622
4829 G	10 021RAMP5	developmentally regulated GTP-binding protein 1	406092 IMAGE Consortium	W83272
4830 G	12 021RAMP5	IMAGE EST {IMAGE:418732}	418732 IMAGE Consortium	W96980
4831 A	2 021YAMP6	low density lipoprotein receptor related protein	407600 IMAGE Consortium	W89938
4832 A	4 021YAMP6	Complement component 1, q subcomponent, c polypeptide {IMAGE:426010}	426010 IMAGE Consortium	AA000461
4833 A	6 021YAMP6	matrilin 4	427166 IMAGE Consortium	AA003393
4834 A	8 021YAMP6	transforming growth factor beta 1 induced transcript 1	438541 IMAGE Consortium	AA050948
4835 A	10 021YAMP6	gene rich cluster, C9 gene	443889 IMAGE Consortium	AA014095
4836 A	12 021YAMP6	Very low density lipoprotein receptor {IMAGE:446084}	446084 IMAGE Consortium	AA020307
4837 C	2 021YAMP6			
4838 C	4 021YAMP6	FBJ osteosarcoma oncogene {IMAGE:426070}	426070 IMAGE Consortium	AA002910
4839 C	6 021YAMP6	CD97 antigen	420765 IMAGE Consortium	AI325851
4840 C	8 021YAMP6	Smoothened homolog (Drosophila) {IMAGE:439010}	439010 IMAGE Consortium	AA008222
4841 C	10 021YAMP6	alcohol dehydrogenase 5	438799 IMAGE Consortium	AA008355
4842 C	12 021YAMP6	Mus musculus SMAD3 (mSmad3) mRNA, complete cds {IMAGE:456861}	456861 IMAGE Consortium	AA023641
4843 E	2 021YAMP6	ubiquitin carboxy-terminal hydrolase L1	407703 IMAGE Consortium	AI430255
4844 E	4 021YAMP6	Tumor necrosis factor receptor superfamily, member 1b {IMAGE:437512}	437512 IMAGE Consortium	AA003594
4845 E	6 021YAMP6	C-terminal binding protein 2	426531 IMAGE Consortium	AA002394
4846 E	8 021YAMP6	ESTs, Moderately similar to CPE-receptor [M.musculus] {IMAGE:441695}	441695 IMAGE Consortium	AA013960
4847 E	10 021YAMP6	frizzled homolog 1, (Drosophila)	441720 IMAGE Consortium	AA013810
4848 E	12 021YAMP6	Peripherin {IMAGE:464493}	464493 IMAGE Consortium	AA030447
4849 G	2 021YAMP6	DNA segment, Chr 2, ERATO Doi 554, expressed	407016 IMAGE Consortium	W83757
4850 G	4 021YAMP6	Dlk1-like homolog (Drosophila) {IMAGE:427339}	427339 IMAGE Consortium	AA003303
4851 G	6 021YAMP6	ESTs	438201 IMAGE Consortium	AA008074
4852 G	8 021YAMP6	Tumor necrosis factor induced protein 2 {IMAGE:443884}	443884 IMAGE Consortium	AA014136
4853 G	10 021YAMP6	DNA segment, Chr 10, Wayne State University 136, expressed	441768 IMAGE Consortium	AA013993
4854 G	12 021YAMP6	Keratin complex 1, acidic, gene 19 {IMAGE:464060}	464060 IMAGE Consortium	AA028346

4855 A	2 0215AMP7	hydroxysteroid 17-beta dehydrogenase 2	459718 IMAGE Consortium	AA027607
4856 A	4 0215AMP7	Paired mesoderm homeobox 1 {IMAGE:476293}	476293 IMAGE Consortium	AA050726
4857 A	6 0215AMP7			
4858 A	8 0215AMP7	ESTs	480554 IMAGE Consortium	AA064189
4859 A	10 0215AMP7			
4860 A	12 0215AMP7	keratin complex 1, acidic, gene 13	483466 IMAGE Consortium	AA061729
4861 C	2 0215AMP7	small proline-rich protein 1A	463982 IMAGE Consortium	AA028420
4862 C	4 0215AMP7	Retinoid X receptor beta {IMAGE:475673}	475673 IMAGE Consortium	AA050298
4863 C	6 0215AMP7	cytokine inducible SH2-containing protein 2	479296 IMAGE Consortium	AA048926
4864 C	8 0215AMP7	heterogeneous nuclear ribonucleoprotein D	479057 IMAGE Consortium	AA049003
4865 C	10 0215AMP7	IMAGE EST {IMAGE:482943}	482943 IMAGE Consortium	AA062182
4866 C	12 0215AMP7	pyruvate decarboxylase	482835 IMAGE Consortium	AI323108
4867 E	2 0215AMP7	rhotekin	466059 IMAGE Consortium	AA034678
4868 E	4 0215AMP7	histidine rich calcium binding protein	475063 IMAGE Consortium	AI322629
4869 E	6 0215AMP7	transketolase	479328 IMAGE Consortium	AI385796
4870 E	8 0215AMP7	copper chaperone for superoxide dismutase	477633 IMAGE Consortium	AA048511
4871 E	10 0215AMP7			
4872 E	12 0215AMP7	cysteine rich intestinal protein	482898 IMAGE Consortium	AI385503
4873 G	2 0215AMP7	placental lactogen 2	459622 IMAGE Consortium	AA027530
4874 G	4 0215AMP7	neuron specific gene family member 1	473204 IMAGE Consortium	AI323002
4875 G	6 0215AMP7	Yamaguchi sarcoma viral (v-yes-1) oncogene homolog	479402 IMAGE Consortium	AA049131
4876 G	8 0215AMP7	scavenger receptor class B1	479844 IMAGE Consortium	AA051479
4877 G	10 0215AMP7	solute carrier family 3, member 1	482245 IMAGE Consortium	AI385484
4878 G	12 0215AMP7	Calmodulin 3 {IMAGE:571789}	571789 IMAGE Consortium	AA109041
4879 A	2 021CAMP8	CD3 antigen, epsilon polypeptide	576441 IMAGE Consortium	AI323257
4880 A	4 021CAMP8	serum/glucocorticoid regulated kinase-like	581105 IMAGE Consortium	AA138663
4881 A	6 021CAMP8	leukotriene A4 hydrolase	596365 IMAGE Consortium	AI324333
4882 A	8 021CAMP8	vascular cell adhesion molecule 1	635211 IMAGE Consortium	AA189527
4883 A	10 021CAMP8	nuclear receptor subfamily 2, group F, member 6	657948 IMAGE Consortium	AA198237
4884 A	12 021CAMP8	cleavage stimulation factor, 3' pre-RNA, subunit 3	669834 IMAGE Consortium	AA230454
4885 C	2 021CAMP8	dynactin 3	570659 IMAGE Consortium	AA108719
4886 C	4 021CAMP8	lymphocyte antigen 6 complex, locus E	582679 IMAGE Consortium	AI326722
4887 C	6 021CAMP8	Interferon activated gene 203 {IMAGE:618572}	618572 IMAGE Consortium	AA170251
4888 C	8 021CAMP8	ESTs {IMAGE:642467}	642467 IMAGE Consortium	AA190123
4889 C	10 021CAMP8	ESTs	660439 IMAGE Consortium	AA220400
4890 C	12 021CAMP8	Mus musculus EIG 180 mRNA for ethanol induced gene product, complete cds	671364 IMAGE Consortium	AA222190
4891 E	2 021CAMP8	guanine nucleotide binding protein (G protein), gamma 12	573701 IMAGE Consortium	AA118767
4892 E	4 021CAMP8	prothymosin beta 4	303438 IMAGE Consortium	W09641
4893 E	6 021CAMP8	ral guanine nucleotide dissociation stimulator,-like 2	634926 IMAGE Consortium	AA154819
4894 E	8 021CAMP8	dolichyl-phosphate alpha-N-acetylglucosaminephosphotransferase 2	640474 IMAGE Consortium	AA204168

4895 E	10 021CAMP8	phosphotriesterase related	669815 IMAGE Consortium	AA230452
4896 E	12 021CAMP8	necdin	670072 IMAGE Consortium	AA231099
4897 G	2 021CAMP8	differentially expressed in B16F10 2	570994 IMAGE Consortium	AA108417
4898 G	4 021CAMP8	3-hydroxy-3-methylglutaryl-Coenzyme A lyase	313248 IMAGE Consortium	W11903
4899 G	6 021CAMP8			
4900 G	8 021CAMP8	small nuclear ribonucleoprotein D1	636184 IMAGE Consortium	AA183306
4901 G	10 021CAMP8	kinesin heavy chain member 2	671634 IMAGE Consortium	AA241771
4902 G	12 021CAMP8	phosphomannomutase 2	658481 IMAGE Consortium	AA221696
4903 A	2 021JAMP9	orosomuroid 2	679136 IMAGE Consortium	AA245687
4904 A	4 021JAMP9	Heparin cofactor II {IMAGE:693170}	693170 IMAGE Consortium	AA253700
4905 A	6 021JAMP9	Zinc finger protein 118	697368 IMAGE Consortium	AA242194
4906 A	8 021JAMP9	multiple inositol polyphosphate histidine phosphatase 1	722780 IMAGE Consortium	AA254526
4907 A	10 021JAMP9	flavin containing monooxygenase 5	734267 IMAGE Consortium	AA268913
4908 A	12 021JAMP9	Mus musculus clone L5 uniform group of 2-cell-stage gene family mRNA, complete cds	751180 IMAGE Consortium	AA289585
4909 C	2 021JAMP9	proteasome (prosome, macropain) subunit, beta type 8 (large multifunctional protease 7)	681039 IMAGE Consortium	AA250087
4910 C	4 021JAMP9	Mus musculus semaphorin IV isoform b mRNA, complete cds {IMAGE:695117}	695117 IMAGE Consortium	AA216900
4911 C	6 021JAMP9			
4912 C	8 021JAMP9	procollagen, type XV	720629 IMAGE Consortium	AA261149
4913 C	10 021JAMP9	serum amyloid P-component	720590 IMAGE Consortium	AA261401
4914 C	12 021JAMP9	protein kinase C, beta	751182 IMAGE Consortium	AA289586
4915 E	2 021JAMP9	spinocerebellar ataxia 2 homolog (human)	679715 IMAGE Consortium	AA245190
4916 E	4 021JAMP9	Surfactant associated protein C {IMAGE:695697}	695697 IMAGE Consortium	AA222208
4917 E	6 021JAMP9	Mus musculus Trex2 mRNA, complete sequence	695452 IMAGE Consortium	AA220178
4918 E	8 021JAMP9	dendritic cell protein GA17	720150 IMAGE Consortium	AA254837
4919 E	10 021JAMP9	Mus musculus aquaporin-8 mRNA, complete cds {IMAGE:747401}	747401 IMAGE Consortium	AA272836
4920 E	12 021JAMP9	ERG-associated protein	751905 IMAGE Consortium	AA396251
4921 G	2 021JAMP9	insulin-like growth factor binding protein, acid labile subunit	681320 IMAGE Consortium	AA250132
4922 G	4 021JAMP9	Duffy blood group {IMAGE:718084}	718084 IMAGE Consortium	AA266438
4923 G	6 021JAMP9	adaptor-related protein complex AP-3, delta subunit	699896 IMAGE Consortium	AA958641
4924 G	8 021JAMP9	RNA binding motif protein 4	720918 IMAGE Consortium	AA267187
4925 G	10 021JAMP9	Inositol polyphosphate-5-phosphatase, 75 kDa {IMAGE:749425}	749425 IMAGE Consortium	AA289577
4926 G	12 021JAMP9			
4927 A	2 0213AMPA	5'-3' exoribonuclease 2	737014 IMAGE Consortium	AA272130
4928 A	4 0213AMPA	special AT-rich sequence binding protein 1	765694 IMAGE Consortium	AA275312
4929 A	6 0213AMPA	ESTs, Highly similar to PHOSPHOLIPASE A-2-ACTIVATING PROTEIN [M.musculus] {IMAGE:831959}	831959 IMAGE Consortium	AA444488
4930 A	8 0213AMPA	suppressor of K <sup>+</sup> transport defect 3	821075 IMAGE Consortium	AA646994

4931 A	10 0213AMPA	Mus musculus Xin mRNA, complete cds {IMAGE:860360}	860360 IMAGE Consortium	AA501052
4932 A	12 0213AMPA	ATP-binding cassette, sub-family A (ABC1), member 7	873998 IMAGE Consortium	AA472176
4933 C	2 0213AMPA	Mus musculus mRNA for alpha-albumin protein	747307 IMAGE Consortium	AA274876
4934 C	4 0213AMPA	Mus musculus Ca <sup>2+</sup> -independent phospholipase A2 long form mRNA, complete cds	762922 IMAGE Consortium	AA271866
4935 C	6 0213AMPA	Mus musculus polypeptide GalNAc transferase-T1 (ppGaNTase-T1) mRNA, complete cds {IMAGE:849524}	849524 IMAGE Consortium	AA457927
4936 C	8 0213AMPA	ADP-ribosyltransferase 5	314781 IMAGE Consortium	W12489
4937 C	10 0213AMPA	Mus musculus mRNA for DEDD protein {IMAGE:874041}	874041 IMAGE Consortium	AA471746
4938 C	12 0213AMPA	caspace 2	871077 IMAGE Consortium	AA462472
4939 E	2 0213AMPA	betaine-homocysteine methyltransferase	747383 IMAGE Consortium	AA272831
4940 E	4 0213AMPA	encephalopsin	777428 IMAGE Consortium	AA276584
4941 E	6 0213AMPA	syntaxin 3	850497 IMAGE Consortium	AA451351
4942 E	8 0213AMPA	cytochrome c oxidase, subunit VI a, polypeptide 1	316084 IMAGE Consortium	W08923
4943 E	10 0213AMPA	potassium channel, subfamily K, member 2	851648 IMAGE Consortium	AA500739
4944 E	12 0213AMPA	a disintegrin and metalloproteinase domain 17	860231 IMAGE Consortium	AA500981
4945 G	2 0213AMPA	ESTs {IMAGE:775858}	775858 IMAGE Consortium	AA276003
4946 G	4 0213AMPA	vitamin D receptor	776764 IMAGE Consortium	AA273494
4947 G	6 0213AMPA	Mpv17 transgene, kidney disease mutant	850544 IMAGE Consortium	AA450927
4948 G	8 0213AMPA	heterogeneous nuclear ribonucleoprotein A1	316442 IMAGE Consortium	W11164
4949 G	10 0213AMPA	myosin VI	872172 IMAGE Consortium	AA471862
4950 G	12 0213AMPA	conserved helix-loop-helix ubiquitous kinase	871154 IMAGE Consortium	AA462677
4951 A	2 021AAMPB	Mus musculus protein-tyrosine phosphatase mRNA, complete cds {IMAGE:935557}	935557 IMAGE Consortium	AA543569
4952 A	4 021AAMPB	glutathione S-transferase, theta 2	891492 IMAGE Consortium	AA511089
4953 A	6 021AAMPB	chromobox homolog 3 (Drosophila HP1 gamma)	961342 IMAGE Consortium	AA547637
4954 A	8 021AAMPB	enolase 3, beta muscle	1122937 IMAGE Consortium	AA681602
4955 A	10 021AAMPB	eukaryotic translation initiation factor 2 alpha kinase 4	1195967 IMAGE Consortium	AA716848
4956 A	12 021AAMPB	solute carrier family 22, member 3	1178743 IMAGE Consortium	AA710965
4957 C	2 021AAMPB	solute carrier family 12, member 4	890956 IMAGE Consortium	AA511202
4958 C	4 021AAMPB	E26 avian leukemia oncogene 2, 3' domain	935316 IMAGE Consortium	AA542151
4959 C	6 021AAMPB	triosephosphate isomerase	1065480 IMAGE Consortium	AA611441
4960 C	8 021AAMPB	ATPase, H <sup>+</sup> transporting lysosomal (vacuolar proton pump), 32 kDa	1138971 IMAGE Consortium	AI536399
4961 C	10 021AAMPB	torsin family 3, member A	1180009 IMAGE Consortium	AA791729
4962 C	12 021AAMPB	Mus musculus pcm-1 mRNA for pericentriolar material-1, complete cds	1162868 IMAGE Consortium	AA770808
4963 E	2 021AAMPB	protein kinase, interferon inducible double stranded RNA dependent activator	935506 IMAGE Consortium	AA543453
4964 E	4 021AAMPB	integrin beta 7	963335 IMAGE Consortium	AA671769
4965 E	6 021AAMPB	ankyrin-like repeat protein	1153720 IMAGE Consortium	AA792499
4966 E	8 021AAMPB	protease (prosome, macropain) 28 subunit, alpha	1123011 IMAGE Consortium	AA794081
4967 E	10 021AAMPB	enolase 2, gamma neuronal	1165522 IMAGE Consortium	AA929497



4968 E	12 021AAMPB	protein phosphatase 1A, magnesium dependent, alpha isoform	1162895 IMAGE Consortium	AA770833
4969 G	2 021AAMPB	guanine nucleotide binding protein (G protein), gamma 2 subunit	876696 IMAGE Consortium	AA499295
4970 G	4 021AAMPB	RAB33B, member of RAS oncogene family	1025621 IMAGE Consortium	AA590871
4971 G	6 021AAMPB	ribosomal protein, mitochondrial, L14	1153785 IMAGE Consortium	AA792656
4972 G	8 021AAMPB	Tnf receptor-associated factor 5	1066820 IMAGE Consortium	AA615872
4973 G	10 021AAMPB	stearoyl-Coenzyme A desaturase 1	1195323 IMAGE Consortium	AA754922
4974 G	12 021AAMPB			
4975 A	2 021HAMPC	Rab6 interacting protein 1	1162974 IMAGE Consortium	AI592482
4976 A	4 021HAMPC	general control of amino acid synthesis-like 1 (yeast)	1230868 IMAGE Consortium	AA880081
4977 A	6 021HAMPC	ESTs, Highly similar to protein-serine/threonine kinase [M.musculus]	1265317 IMAGE Consortium	AA867125
4978 A	8 021HAMPC	Public domain EST	1263454 IMAGE Consortium	
4979 A	10 021HAMPC	phosphodiesterase 7B	1364394 IMAGE Consortium	AI006099
4980 A	12 021HAMPC	plasma glutamate carboxypeptidase	1382015 IMAGE Consortium	AA983003
4981 C	2 021HAMPC	benzodiazepine receptor, peripheral	1195602 IMAGE Consortium	AA711705
4982 C	4 021HAMPC	small EDRK-rich factor 2	1247525 IMAGE Consortium	AA763355
4983 C	6 021HAMPC	guanine deaminase	1346255 IMAGE Consortium	AA959127
4984 C	8 021HAMPC	peptidyl arginine deiminase, type II	1329135 IMAGE Consortium	AA929696
4985 C	10 021HAMPC	small inducible cytokine subfamily A20	1362268 IMAGE Consortium	AA990366
4986 C	12 021HAMPC	sialyltransferase 4C (beta-galactosidase alpha-2,3-sialyltransferase)	1382146 IMAGE Consortium	AA798202
4987 E	2 021HAMPC	wingless related MMTV integration site 8b	1225089 IMAGE Consortium	AA756392
4988 E	4 021HAMPC	valosin containing protein	1265872 IMAGE Consortium	AI509818
4989 E	6 021HAMPC	insulin-like 6	1329516 IMAGE Consortium	AA940549
4990 E	8 021HAMPC	forkhead box N1	1329145 IMAGE Consortium	AA929753
4991 E	10 021HAMPC	zinc finger protein 35	1378962 IMAGE Consortium	AI020352
4992 E	12 021HAMPC	protein phosphatase 4, catalytic subunit	1382184 IMAGE Consortium	AA798281
4993 G	2 021HAMPC	frizzled homolog 4 (Drosophila)	1179017 IMAGE Consortium	AA717541
4994 G	4 021HAMPC	cystatin F (leukocystatin)	1314895 IMAGE Consortium	AA881075
4995 G	6 021HAMPC	RAD21 homolog (S. pombe)	1345616 IMAGE Consortium	AI154347
4996 G	8 021HAMPC	mitogen activated protein kinase kinase kinase 7	1345955 IMAGE Consortium	AA967493
4997 G	10 021HAMPC	angiotensin converting enzyme	1382621 IMAGE Consortium	AI036411
4998 G	12 021HAMPC	arachidonate 12-lipoxygenase, pseudogene 2	1379994 IMAGE Consortium	AI050607
4999 A	2 021OAMPD	galactosidase, alpha	1397736 IMAGE Consortium	AI118621
5000 A	4 021OAMPD	ESTs, Highly similar to 60S RIBOSOMAL PROTEIN L35 [Homo sapiens]	1383576 IMAGE Consortium	AI120979
5001 A	6 021OAMPD	ATP-binding cassette, sub-family E (OABP), member 1	1400851 IMAGE Consortium	AI121701
5002 A	8 021OAMPD	sodium channel, voltage-gated, type I, beta polypeptide	1494022 IMAGE Consortium	AI156705
5003 A	10 021OAMPD	protein tyrosine phosphatase, receptor type, D	1448630 IMAGE Consortium	AI528928
5004 A	12 021OAMPD	zinc finger protein 67	332479 IMAGE Consortium	W08617
5005 C	2 021OAMPD	metastasis associated 1-like 1	1383287 IMAGE Consortium	AI120748

5006 C	4 021OAMPD	eleven-nineteen lysine-rich leukemia gene	1399580 IMAGE Consortium	AI121419
5007 C	6 021OAMPD	Pendred syndrome homolog (human)	1478576 IMAGE Consortium	AI156032
5008 C	8 021OAMPD	sporulation protein, meiosis-specific, SPO11 homolog ( <i>S. cerevisiae</i> )	1430293 IMAGE Consortium	AI181574
5009 C	10 021OAMPD	complement component 1, q subcomponent, beta polypeptide	1430527 IMAGE Consortium	AI182838
5010 C	12 021OAMPD	transducin-like enhancer of split 1, homolog of <i>Drosophila</i> E(spl)	318586 IMAGE Consortium	AI536358
5011 E	2 021OAMPD	complement component 1 inhibitor	1383322 IMAGE Consortium	AI120822
5012 E	4 021OAMPD	purine rich element binding protein A	1382922 IMAGE Consortium	AI120282
5013 E	6 021OAMPD	zinc finger protein 1	1447390 IMAGE Consortium	AI154200
5014 E	8 021OAMPD	solute carrier family 12, member 7	1447730 IMAGE Consortium	AI158743
5015 E	10 021OAMPD	budding uninhibited by benzimidazoles 3 homolog ( <i>S. cerevisiae</i> )	318275 IMAGE Consortium	W13647
5016 E	12 021OAMPD	serine protease inhibitor 1-2	332062 IMAGE Consortium	W08430
5017 G	2 021OAMPD	DEAD/H (Asp-Glu-Ala-Asp/His) box polypeptide 19	1384089 IMAGE Consortium	AI181655
5018 G	4 021OAMPD	splicing factor, arginine/serine-rich 5 (SRp40, HRS)	1382973 IMAGE Consortium	AI120315
5019 G	6 021OAMPD	dynamitin 2	1493947 IMAGE Consortium	AI156587
5020 G	8 021OAMPD	hydroxysteroid 11-beta dehydrogenase 1	1495778 IMAGE Consortium	AI789916
5021 G	10 021OAMPD	ribosomal protein L12	318345 IMAGE Consortium	W13748
5022 G	12 021OAMPD	feminization 1 a homolog ( <i>C. elegans</i> )	331435 IMAGE Consortium	W14831
5023 A	2 021VAMPE	a disintegrin and metalloproteinase domain 9 (meltrin gamma)	1749980 IMAGE Consortium	AI047163
5024 A	4 021VAMPE	ESTs, Highly similar to DESTRAIN [ <i>Homo sapiens</i> ; <i>Sus scrofa</i> ] {IMAGE:335112}	335112 IMAGE Consortium	W17549
5025 A	6 021VAMPE	ubiquitin-conjugating enzyme E2B (RAD6 homology)	333483 IMAGE Consortium	W33948
5026 A	8 021VAMPE	histone H1	335232 IMAGE Consortium	W17615
5027 A	10 021VAMPE	dynein, axon, heavy chain 11	332860 IMAGE Consortium	W15606
5028 A	12 021VAMPE	arylsulfatase A	336579 IMAGE Consortium	W36720
5029 C	2 021VAMPE	suppressor of variegation 3-9 homologue 1 ( <i>Drosophila</i> )	1749990 IMAGE Consortium	AI047180
5030 C	4 021VAMPE	FK506-BINDING PROTEIN PRECURSOR {IMAGE:333683}	333683 IMAGE Consortium	W15971
5031 C	6 021VAMPE	glutathione S-transferase like	333818 IMAGE Consortium	W16059
5032 C	8 021VAMPE	ret finger protein	334575 IMAGE Consortium	AI391314
5033 C	10 021VAMPE	P450 (cytochrome) oxidoreductase	335079 IMAGE Consortium	AI322279
5034 C	12 021VAMPE	peptidylprolyl isomerase C-associated protein	337797 IMAGE Consortium	W29311
5035 E	2 021VAMPE	germinal center expressed transcript	2698933 IMAGE Consortium	AW519547
5036 E	4 021VAMPE	DIFF6 PROTEIN {IMAGE:334138}	334138 IMAGE Consortium	W16222
5037 E	6 021VAMPE	cathepsin D	334176 IMAGE Consortium	W16244
5038 E	8 021VAMPE	cell death-inducing DNA fragmentation factor, alpha subunit-like effector A	333124 IMAGE Consortium	W15725
5039 E	10 021VAMPE	hemoglobin, beta adult major chain	332935 IMAGE Consortium	W15809
5040 E	12 021VAMPE	drebrin-like	348161 IMAGE Consortium	W30616

5041 G	2 021VAMPE	cytokine inducible SH2-containing protein 1	1746460 IMAGE Consortium	AI037689
5042 G	4 021VAMPE	Untitled {IMAGE:332480}	332480 IMAGE Consortium	W08488
5043 G	6 021VAMPE	metallothionein 2	334351 IMAGE Consortium	W36474
5044 G	8 021VAMPE	sarcoglycan, alpha (50kDa dystrophin-associated glycoprotein)	334713 IMAGE Consortium	W18272
5045 G	10 021VAMPE	alpha fetoprotein	332979 IMAGE Consortium	AI324392
5046 G	12 021VAMPE	dopachrome tautomerase	349266 IMAGE Consortium	W29342
5047 A	2 0212AMPF	proteasome (prosome, macropain) subunit, alpha type 2	335436 IMAGE Consortium	AI390129
5048 A	4 0212AMPF	demilune cell and parotid protein	335896 IMAGE Consortium	W36921
5049 A	6 0212AMPF	Tial1 cytotoxic granule-associated RNA-binding protein-like 1	353542 IMAGE Consortium	W41637
5050 A	8 0212AMPF			
5051 A	10 0212AMPF	Notch2-like	367362 IMAGE Consortium	W50780
5052 A	12 0212AMPF	valyl-tRNA synthetase 2	387511 IMAGE Consortium	AI893239
5053 C	2 0212AMPF	cytosolic adenylate kinase	337692 IMAGE Consortium	AI391125
5054 C	4 0212AMPF	IMAGE EST {IMAGE:350336}	350336 IMAGE Consortium	W34967
5055 C	6 0212AMPF	immediate early response 5	354281 IMAGE Consortium	W44262
5056 C	8 0212AMPF	procollagen, type IV, alpha 2	368514 IMAGE Consortium	W53787
5057 C	10 0212AMPF	myeloid cell leukemia sequence 1	368211 IMAGE Consortium	W54056
5058 C	12 0212AMPF	sepiapterin reductase	386989 IMAGE Consortium	AI893229
5059 E	2 0212AMPF	growth arrest and DNA-damage-inducible, gamma	335551 IMAGE Consortium	W18822
5060 E	4 0212AMPF	M. musculus mRNA for ribosomal protein S8 {IMAGE:354601}	354601 IMAGE Consortium	W46090
5061 E	6 0212AMPF	Moloney leukemia virus 10	352633 IMAGE Consortium	W34188
5062 E	8 0212AMPF	CD82 antigen	368866 IMAGE Consortium	W53231
5063 E	10 0212AMPF	procollagen, type VI, alpha 2	368442 IMAGE Consortium	W53604
5064 E	12 0212AMPF	gene rich cluster, C2f gene	374897 IMAGE Consortium	W62577
5065 G	2 0212AMPF	selenoprotein W, muscle 1	335583 IMAGE Consortium	W18835
5066 G	4 0212AMPF	M. musculus mRNA for MAP kinase-activated protein kinase 2 {IMAGE:354859}	354859 IMAGE Consortium	W45833
5067 G	6 0212AMPF	glutathione S-transferase, mu 6	353114 IMAGE Consortium	W34862
5068 G	8 0212AMPF	Mus musculus STAT6 (Stat6) gene, partial cds and NAB2 (Nab2) gene, 3'UTR	367815 IMAGE Consortium	W54006
5069 G	10 0212AMPF	WW domain binding protein 1	368496 IMAGE Consortium	W53568
5070 G	12 0212AMPF	hypothetical protein, MNCb-5210	386561 IMAGE Consortium	AI430585
5071 A	2 0211AJ7N	Internal_Control_E	Internal_Control_E	Controls
5072 A	4 0211AJ7N	Internal_Control_F	Internal_Control_F	Controls
5073 A	6 0211AJ7N	Internal_Control_G	Internal_Control_G	Controls
5074 A	8 0211AJ7N	Internal_Control_H	Internal_Control_H	Controls
5075 A	10 0211AJ7N	Internal_Control_I	Internal_Control_I	Controls
5076 A	12 0211AJ7N	Internal_Control_J	Internal_Control_J	Controls
5077 C	2 0211AJ7N	Internal_Control_K	Internal_Control_K	Controls
5078 C	4 0211AJ7N	Internal_Control_L	Internal_Control_L	Controls

5079 C	6 0211AJ7N	Internal_Control_M	Internal_Control_M	Controls
5080 C	8 0211AJ7N	Internal_Control_N	Internal_Control_N	Controls
5081 C	10 0211AJ7N	Internal_Control_O	Internal_Control_O	Controls
5082 C	12 0211AJ7N	Internal_Control_P	Internal_Control_P	Controls
5083 E	2 0211AJ7N	Internal_Control_Q	Internal_Control_Q	Controls
5084 E	4 0211AJ7N	Internal_Control_R	Internal_Control_R	Controls
5085 E	6 0211AJ7N	Internal_Control_S	Internal_Control_S	Controls
5086 E	8 0211AJ7N	Internal_Control_T	Internal_Control_T	Controls
5087 E	10 0211AJ7N	Internal_Control_U	Internal_Control_U	Controls
5088 E	12 0211AJ7N	Internal_Control_V	Internal_Control_V	Controls
5089 G	2 0211AJ7N	Internal_Control_W	Internal_Control_W	Controls
5090 G	4 0211AJ7N	Internal_Control_X	Internal_Control_X	Controls
5091 G	6 0211AJ7N	Internal_Control_Y	Internal_Control_Y	Controls
5092 G	8 0211AJ7N	Internal_Control_Z	Internal_Control_Z	Controls
5093 G	10 0211AJ7N	Internal_Control_AA	Internal_Control_AA	Controls
5094 G	12 0211AJ7N	Internal_Control_AB	Internal_Control_AB	Controls
5101 B	1 0217AJ7T	Control: Ratio 1:3 (Cy3:Cy5)	Ratio_A	Controls
5102 B	3 0217AJ7T	Control: Ratio 3:1 (Cy3:Cy5)	Ratio_B	Controls
5103 B	5 0217AJ7T	Control: Ratio 1:10 (Cy3:Cy5)	Ratio_C	Controls
5104 B	7 0217AJ7T	Control: Ratio 10:1 (Cy3:Cy5)	Ratio_D	Controls
5105 B	9 0217AJ7T	Control: Ratio 1:25 (Cy3:Cy5)	Ratio_E	Controls
5106 B	11 0217AJ7T	Control: Ratio 25:1 (Cy3:Cy5)	Ratio_F	Controls
5107 D	1 0217AJ7T	Control: Sensitivity 2pg	Sensitivity_A	Controls
5108 D	3 0217AJ7T	Control: Sensitivity 20pg	Sensitivity_B	Controls
5109 D	5 0217AJ7T	Control: Sensitivity 200pg	Sensitivity_C	Controls
5110 D	7 0217AJ7T	Control: Sensitivity 2000pg	Sensitivity_D	Controls
5111 D	9 0217AJ7T	Control: Buffer Only	Buffer_Only_Control	Controls
5112 D	11 0217AJ7T	Control: Buffer Only	Buffer_Only_Control	Controls
5113 F	1 0217AJ7T	Control: Internal	Internal_Control_A	Controls
5114 F	3 0217AJ7T	Control: Buffer Only	Buffer_Only_Control	Controls
5115 F	5 0217AJ7T	Control: Internal	Internal_Control_B	Controls
5116 F	7 0217AJ7T	Control: Buffer Only	Buffer_Only_Control	Controls
5117 F	9 0217AJ7T	Control: Internal	Internal_Control_C	Controls
5118 F	11 0217AJ7T	Control: Internal	Internal_Control_D	Controls
5119 H	1 0217AJ7T	Control: Complex Target (Mus musculus)	RNA_Quality_MM	Controls
5120 H	3 0217AJ7T	Control: Complex Target (Mus musculus)	RNA_Quality_MM	Controls
5121 H	5 0217AJ7T	Control: Complex Target (Mus musculus)	RNA_Quality_MM	Controls
5122 H	7 0217AJ7T	Control: Housekeeping (Alpha Tubulin)	Housekeeping_B	Controls
5123 H	9 0217AJ7T	Control: Housekeeping (Ribosomal S9)	Housekeeping_A	Controls
5124 H	11 0217AJ7T	Control: Housekeeping (23kD HBP)	Housekeeping_C	Controls
5125 B	1 021JAMMK	astrotactin	313186 IMAGE Consortium	W10500
5126 B	3 021JAMMK	ESTs	313537 IMAGE Consortium	AI605613

5127 B	5 021JAMMK	calcitonin gene-related peptide-receptor component protein	315404 IMAGE Consortium	AI605726
5128 B	7 021JAMMK	ESTs	315942 IMAGE Consortium	AI605772
5129 B	9 021JAMMK	kinesin family member 5A	419429 IMAGE Consortium	AI893980
5130 B	11 021JAMMK	ESTs	421506 IMAGE Consortium	W99206
5131 D	1 021JAMMK	hairy and enhancer of split 6, ( <i>Drosophila</i> )	303416 IMAGE Consortium	AI627138
5132 D	3 021JAMMK	hepatocyte growth factor activator	314294 IMAGE Consortium	AI327429
5133 D	5 021JAMMK	homeo box C8	316145 IMAGE Consortium	AI327461
5134 D	7 021JAMMK	ESTs, Highly similar to KIAA0453 protein [ <i>H.sapiens</i> ]	316853 IMAGE Consortium	AI607008
5135 D	9 021JAMMK	ESTs	418821 IMAGE Consortium	AI430316
5136 D	11 021JAMMK	frequenin homolog ( <i>Drosophila</i> )	423183 IMAGE Consortium	W97305
5137 F	1 021JAMMK	ESTs	303445 IMAGE Consortium	AI627143
5138 F	3 021JAMMK	ESTs, Weakly similar to fibrousheathin II [ <i>H.sapiens</i> ]	313592 IMAGE Consortium	AI605616
5139 F	5 021JAMMK	phospholipase A2, group IB, pancreas	315430 IMAGE Consortium	AI327450
5140 F	7 021JAMMK	Sequence 4 from Patent WO9957133	316884 IMAGE Consortium	AI536280
5141 F	9 021JAMMK	ISL1 transcription factor, LIM/homeodomain, ( <i>islet-1</i> )	437224 IMAGE Consortium	AA003609
5142 F	11 021JAMMK	ESTs, Moderately similar to bithoraxoid-like protein [ <i>R.norvegicus</i> ]	423549 IMAGE Consortium	W98786
5143 H	1 021JAMMK	ESTs	313060 IMAGE Consortium	W10432
5144 H	3 021JAMMK	crystallin, gamma A	313601 IMAGE Consortium	W10465
5145 H	5 021JAMMK	septin 3	315515 IMAGE Consortium	AI605734
5146 H	7 021JAMMK	ESTs, Highly similar to hepatic nuclear factor 1-beta short form [ <i>M.musculus</i> ]	422998 IMAGE Consortium	AI894004
5147 H	9 021JAMMK	ESTs	437561 IMAGE Consortium	AA003774
5148 H	11 021JAMMK	homeo box D12	424673 IMAGE Consortium	AI894011
5149 B	1 021QAMML	ESTs	539926 IMAGE Consortium	AA120202
5150 B	3 021QAMML	ESTs, Highly similar to PROTEIN HI1715 [ <i>Haemophilus influenzae</i> ]	540481 IMAGE Consortium	AI892530
5151 B	5 021QAMML	ESTs	539519 IMAGE Consortium	AA153231
5152 B	7 021QAMML	upstream transcription factor 1	539108 IMAGE Consortium	AA117357
5153 B	9 021QAMML	<i>Mus musculus</i> unknown protein mRNA, partial cds	539313 IMAGE Consortium	AI550227
5154 B	11 021QAMML	DNA segment, Chr 14, ERATO Doi 209, expressed	539573 IMAGE Consortium	AI550246
5155 D	1 021QAMML	ESTs, Moderately similar to AF220052_1 uncharacterized hematopoietic stem/progenitor cells protein MDS032 [ <i>H.sapiens</i> ]	539996 IMAGE Consortium	AA122677
5156 D	3 021QAMML	ESTs, Weakly similar to F59E12.5 [ <i>C.elegans</i> ]	540497 IMAGE Consortium	AA155311
5157 D	5 021QAMML	ESTs, Highly similar to PROTEIN TRANSLATION FACTOR SUI1 HOMOLOG [ <i>Anopheles gambiae</i> ]	539581 IMAGE Consortium	AI550248
5158 D	7 021QAMML	ESTs	539134 IMAGE Consortium	AA117252
5159 D	9 021QAMML	beta-spectrin 2, non-erythrocytic	539324 IMAGE Consortium	AA117511
5160 D	11 021QAMML	Sec61 alpha isoform 2	538847 IMAGE Consortium	AI550200
5161 F	1 021QAMML	zinc finger protein 106	540229 IMAGE Consortium	AI550554
5162 F	3 021QAMML	ESTs, Highly similar to CEBA_MOUSE	540533 IMAGE Consortium	AA161908

		CCAAT/ENHANCER BINDING PROTEIN ALPHA [M.musculus]		
5163 F	5	021QAMML RNA polymerase I associated factor, 53 kD	539662	IMAGE Consortium AA117502
5164 F	7	021QAMML ESTs	539173	IMAGE Consortium AA116621
5165 F	9	021QAMML TAR (HIV) RNA binding protein 2	539371	IMAGE Consortium AA117478
5166 F	11	021QAMML ESTs, Highly similar to KIAA0657 protein [H.sapiens]	539621	IMAGE Consortium AA123061
5167 H	1	021QAMML Mus musculus e(y)2 homolog mRNA, complete cds	540247	IMAGE Consortium AI550556
5168 H	3	021QAMML ESTs	540650	IMAGE Consortium AI550577
5169 H	5	021QAMML ESTs	539729	IMAGE Consortium AI550270
5170 H	7	021QAMML general transcription factor IIH, polypeptide 2 (44 kDa subunit)	539196	IMAGE Consortium AA116625
5171 H	9	021QAMML ESTs	539398	IMAGE Consortium AA117527
5172 H	11	021QAMML hypothetical protein MNCb-2643	538918	IMAGE Consortium AI550206
5173 B	1	021XAMMM ESTs	541132	IMAGE Consortium AI550608
5174 B	3	021XAMMM mini chromosome maintenance deficient 5 (S. cerevisiae)	540655	IMAGE Consortium AA162741
5175 B	5	021XAMMM ESTs	540910	IMAGE Consortium AI550598
5176 B	7	021XAMMM phosphodiesterase 7A	560009	IMAGE Consortium AA265221
5177 B	9	021XAMMM ESTs, Highly similar to AF151860_1 CGI-102 protein [H.sapiens]	560085	IMAGE Consortium AA265356
5178 B	11	021XAMMM ESTs	560313	IMAGE Consortium AA210243
5179 D	1	021XAMMM ESTs, Highly similar to Bdeight protein [R.norvegicus]	541148	IMAGE Consortium AA139238
5180 D	3	021XAMMM inositol polyphosphate-1-phosphatase	540669	IMAGE Consortium AI324476
5181 D	5	021XAMMM ESTs	540946	IMAGE Consortium AA124301
5182 D	7	021XAMMM ESTs	560068	IMAGE Consortium AA265329
5183 D	9	021XAMMM immunoglobulin (CD79A) binding protein 1	560143	IMAGE Consortium AA210456
5184 D	11	021XAMMM Mus musculus protein tyrosine phosphatase-like protein PTPLB (Ptplb) mRNA, complete cds	560319	IMAGE Consortium AA210263
5185 F	1	021XAMMM ESTs, Weakly similar to ELL2_HUMAN RNA POLYMERASE II ELONGATION FACTOR ELL2□ [H.sapiens]	541164	IMAGE Consortium AA162699
5186 F	3	021XAMMM ESTs, Moderately similar to antigen NY-CO-1 [H.sapiens]	540691	IMAGE Consortium AA155353
5187 F	5	021XAMMM ESTs, Moderately similar to S57447 HPBRIL-7 protein - human□ [H.sapiens]	540972	IMAGE Consortium AI550603
5188 F	7	021XAMMM ESTs, Weakly similar to similar to dynein heavy chain [C.elegans]	560102	IMAGE Consortium AI509376
5189 F	9	021XAMMM forkhead box C1	560170	IMAGE Consortium AA210461
5190 F	11	021XAMMM stem-loop binding protein	560363	IMAGE Consortium AA210416
5191 H	1	021XAMMM breakpoint cluster region protein 1	541226	IMAGE Consortium AA162818
5192 H	3	021XAMMM ESTs, Highly similar to MYOSIN LIGHT CHAIN KINASE [Dictyostelium discoideum]	540775	IMAGE Consortium AA162185
5193 H	5	021XAMMM ESTs, Highly similar to HYPOTHETICAL MYELOID CELL LINE PROTEIN 6 [Homo sapiens]	559816	IMAGE Consortium AA209699
5194 H	7	021XAMMM deoxyguanosine kinase	560118	IMAGE Consortium AA265337

5195 H	9 021XAMMM	ESTs, Highly similar to PROTEIN DISULFIDE ISOMERASE PRECURSOR [Medicago sativa]	560178 IMAGE Consortium	AA210531
5196 H	11 021XAMMM	ESTs	560367 IMAGE Consortium	AA210418
5197 B	1 0214AMMN	ESTs	560479 IMAGE Consortium	AA210488
5198 B	3 0214AMMN	ESTs	560902 IMAGE Consortium	AA212148
5199 B	5 0214AMMN	ESTs, Moderately similar to AAC-RICH MRNA CLONE AAC3 PROTEIN [Dictyostelium discoideum]	822627 IMAGE Consortium	AI549661
5200 B	7 0214AMMN	nuclear localization signal protein absent in velo-cardio-facial patients	822914 IMAGE Consortium	AA438136
5201 B	9 0214AMMN	ESTs	822416 IMAGE Consortium	AI549635
5202 B	11 0214AMMN	pituitary tumor-transforming 1	822510 IMAGE Consortium	AI549649
5203 D	1 0214AMMN	DNA segment, Chr 3, Wayne State University 133, expressed	560499 IMAGE Consortium	AI509378
5204 D	3 0214AMMN	ESTs, Moderately similar to T17302 hypothetical protein DKFZp564D177.1 - human [H.sapiens]	806035 IMAGE Consortium	AA399805
5205 D	5 0214AMMN	ESTs, Highly similar to ISOLEUCYL-TRNA SYNTHETASE, CYTOPLASMIC [Homo sapiens]	822680 IMAGE Consortium	AI549668
5206 D	7 0214AMMN	ribonuclease, RNase A family 4	822932 IMAGE Consortium	AI549690
5207 D	9 0214AMMN	ESTs, Moderately similar to dJ69E11.3 [H.sapiens]	822421 IMAGE Consortium	AI549637
5208 D	11 0214AMMN	protein tyrosine phosphatase, non-receptor type 2	822540 IMAGE Consortium	AI549652
5209 F	1 0214AMMN	ESTs	560537 IMAGE Consortium	AA212083
5210 F	3 0214AMMN	ESTs	806050 IMAGE Consortium	AA399845
5211 F	5 0214AMMN	transforming growth factor beta 1 induced transcript 4	822692 IMAGE Consortium	AI549669
5212 F	7 0214AMMN	ESTs, Highly similar to P300_HUMAN E1A-ASSOCIATED PROTEIN P300 [H.sapiens]	822939 IMAGE Consortium	AI549694
5213 F	9 0214AMMN	zinc finger protein 207	822430 IMAGE Consortium	AA474132
5214 F	11 0214AMMN	ESTs, Moderately similar to unnamed protein product [H.sapiens]	821858 IMAGE Consortium	AA413026
5215 H	1 0214AMMN	ESTs, Highly similar to PUTATIVE ADENOSINE KINASE [Saccharomyces cerevisiae]	560555 IMAGE Consortium	AA209792
5216 H	3 0214AMMN	ESTs, Moderately similar to Y247_HUMAN HYPOTHETICAL PROTEIN KIAA0247 [H.sapiens]	806093 IMAGE Consortium	AA432487
5217 H	5 0214AMMN	DNA segment, Chr 1, Wayne State University 40, expressed	822737 IMAGE Consortium	AA474881
5218 H	7 0214AMMN	Public domain EST	822947 IMAGE Consortium	
5219 H	9 0214AMMN	lysosomal acid lipase 1	822450 IMAGE Consortium	AI549639
5220 H	11 0214AMMN	ESTs	821865 IMAGE Consortium	AI549624
5221 B	1 021BAMMO	ESTs, Weakly similar to testis ubiquitin specific processing protease [R.norvegicus]	821353 IMAGE Consortium	AA438035
5222 B	3 021BAMMO	ESTs, Weakly similar to scaffold attachment factor B [R.norvegicus]	821688 IMAGE Consortium	AI552364
5223 B	5 021BAMMO	ESTs, Highly similar to HSPC004 [H.sapiens]	821810 IMAGE Consortium	AA474172
5224 B	7 021BAMMO	WW domain binding protein 5	865121 IMAGE Consortium	AA469724
5225 B	9 021BAMMO	claudin 6	865184 IMAGE Consortium	AA450511

5226 B	11 021BAMMO	apurinic/apyrimidinic endonuclease	865239 IMAGE Consortium	AA450772
5227 D	1 021BAMMO	ESTs	821367 IMAGE Consortium	AA438040
5228 D	3 021BAMMO	hypothetical protein MNCb-1213	806387 IMAGE Consortium	AA432845
5229 D	5 021BAMMO	ESTs, Moderately similar to unknown [H.sapiens]	865406 IMAGE Consortium	AA444595
5230 D	7 021BAMMO	Down syndrome critical region homolog 1 (human)	865140 IMAGE Consortium	AA450437
5231 D	9 021BAMMO	ESTs, Moderately similar to HYPOTHETICAL 63.5 KD PROTEIN ZK353.1 IN CHROMOSOME III [Caenorhabditis elegans]	865186 IMAGE Consortium	AA450512
5232 D	11 021BAMMO	Y box protein 1	865253 IMAGE Consortium	AA450785
5233 F	1 021BAMMO	apolipoprotein CI	821372 IMAGE Consortium	AI552353
5234 F	3 021BAMMO	Mus musculus zinc finger protein 289 (Zfp289) mRNA, complete cds	806397 IMAGE Consortium	AI597542
5235 F	5 021BAMMO	choline kinase	865040 IMAGE Consortium	AA469630
5236 F	7 021BAMMO	Mus musculus mRNA for erythroid differentiation regulator, partial	865142 IMAGE Consortium	AA450438
5237 F	9 021BAMMO	epidermal growth factor receptor pathway substrate 15	865189 IMAGE Consortium	AA450519
5238 F	11 021BAMMO	RAB10, member RAS oncogene family	865266 IMAGE Consortium	AA450782
5239 H	1 021BAMMO	ESTs	821382 IMAGE Consortium	AA438045
5240 H	3 021BAMMO	ESTs	806422 IMAGE Consortium	AA470281
5241 H	5 021BAMMO	DNA segment, Chr 5, Wayne State University 31, expressed	865047 IMAGE Consortium	AI614737
5242 H	7 021BAMMO	solute carrier family 35 (CMP-sialic acid transporter), member 1	865150 IMAGE Consortium	AA450441
5243 H	9 021BAMMO	cyclin B1, related sequence 1	865199 IMAGE Consortium	AA450523
5244 H	11 021BAMMO	inosine 5'-phosphate dehydrogenase 2	865271 IMAGE Consortium	AA450789
5245 B	1 021IAMMP	ESTs, Highly similar to unnamed protein product [H.sapiens]	865319 IMAGE Consortium	AA450862
5246 B	3 021IAMMP	CDC like kinase 4	865387 IMAGE Consortium	AA444586
5247 B	5 021IAMMP	ESTs, Weakly similar to I49636 DNA-binding protein - mouse [M.musculus]	865786 IMAGE Consortium	AA458440
5248 B	7 021IAMMP	F-box protein FBA	906318 IMAGE Consortium	AA529559
5249 B	9 021IAMMP	lamin B1	908563 IMAGE Consortium	AA591536
5250 B	11 021IAMMP	protein tyrosine phosphatase, non-receptor type substrate 1	907260 IMAGE Consortium	AA591654
5251 D	1 021IAMMP	poly(A) binding protein, nuclear 1	865328 IMAGE Consortium	AA450901
5252 D	3 021IAMMP	calpain 12	865398 IMAGE Consortium	AI643198
5253 D	5 021IAMMP	DNA segment, Chr 14, ERATO Doi 231, expressed	906012 IMAGE Consortium	AI553349
5254 D	7 021IAMMP	proliferation-associated protein 1	906398 IMAGE Consortium	AA529626
5255 D	9 021IAMMP	ESTs, Moderately similar to KIAA0280 [H.sapiens]	908639 IMAGE Consortium	AA555700
5256 D	11 021IAMMP	Mus musculus collagen type IX alpha 3 mRNA, partial cds	907298 IMAGE Consortium	AA671019
5257 F	1 021IAMMP	thyroid hormone receptor interactor 4	865332 IMAGE Consortium	AI643192
5258 F	3 021IAMMP	ESTs, Moderately similar to AF151028_1 HSPC194 [H.sapiens]	865421 IMAGE Consortium	AI643203



5259 F	5 021IAMMP	RNA binding motif, single stranded interacting protein 1	865832 IMAGE Consortium	AA511835
5260 F	7 021IAMMP	ESTs, Highly similar to unnamed protein product [H.sapiens]	906470 IMAGE Consortium	AA529739
5261 F	9 021IAMMP	methyltransferase	908780 IMAGE Consortium	AA591498
5262 F	11 021IAMMP	ESTs, Moderately similar to HYPOTHETICAL 29.4 KD PROTEIN IN STE6-LOS1 INTERGENIC REGION [Saccharomyces cerevisiae]	906632 IMAGE Consortium	AA522301
5263 H	1 021IAMMP			
5264 H	3 021IAMMP	DNA segment, Chr 11, ERATO Doi 768, expressed	865442 IMAGE Consortium	AI643204
5265 H	5 021IAMMP	Mus musculus LEK1 mRNA, partial cds	906048 IMAGE Consortium	AA529656
5266 H	7 021IAMMP	ESTs	905777 IMAGE Consortium	AA529300
5267 H	9 021IAMMP	frequently rearranged in advanced T-cell lymphomas	907020 IMAGE Consortium	AA591714
5268 H	11 021IAMMP			
5269 B	1 021PAMMQ	pyruvate kinase 3	906766 IMAGE Consortium	AA522342
5270 B	3 021PAMMQ	ESTs	906967 IMAGE Consortium	AA591875
5271 B	5 021PAMMQ	centrin 1	513751 IMAGE Consortium	AI326525
5272 B	7 021PAMMQ	testis specific gene A2	514683 IMAGE Consortium	AA064586
5273 B	9 021PAMMQ	Mus musculus mRNA for MDM2 Binding protein (Mtbp gene)	514275 IMAGE Consortium	AA065551
5274 B	11 021PAMMQ	ESTs, Weakly similar to HS44_MOUSE HEAT SHOCK 40 KDA PROTEIN 4 [R.norvegicus]	515177 IMAGE Consortium	AI595594
5275 D	1 021PAMMQ	ESTs, Moderately similar to putative spliceosome associated protein [H.sapiens]	907523 IMAGE Consortium	AA589125
5276 D	3 021PAMMQ	ESTs	906998 IMAGE Consortium	AI551955
5277 D	5 021PAMMQ	phospholipase A2, group IIC	513783 IMAGE Consortium	AA061920
5278 D	7 021PAMMQ	preproacrosin	513983 IMAGE Consortium	AI326527
5279 D	9 021PAMMQ	ESTs, Highly similar to NADH-UBIQUINONE OXIDOREDUCTASE 13 KD-B SUBUNIT [Bos taurus]	514319 IMAGE Consortium	AI595138
5280 D	11 021PAMMQ	testis specific gene A8	515223 IMAGE Consortium	AA064212
5281 F	1 021PAMMQ	ESTs	907593 IMAGE Consortium	AA590533
5282 F	3 021PAMMQ	gastrulation brain homeobox 2	908481 IMAGE Consortium	AA589157
5283 F	5 021PAMMQ	acetylcholine receptor epsilon	514537 IMAGE Consortium	AI385975
5284 F	7 021PAMMQ	expressed sequence tag mouse EST 25	514803 IMAGE Consortium	AI595257
5285 F	9 021PAMMQ	Mus musculus eukaryotic translation elongation factor 1-delta mRNA, complete cds	514368 IMAGE Consortium	AA061091
5286 F	11 021PAMMQ	DNA segment, Chr 9, ERATO Doi 393, expressed	515240 IMAGE Consortium	AA064214
5287 H	1 021PAMMQ	ESTs	906875 IMAGE Consortium	AA522418
5288 H	3 021PAMMQ	ESTs, Highly similar to containing human KIAA0714 protein [H.sapiens]	908644 IMAGE Consortium	AA555702
5289 H	5 021PAMMQ	ESTs, Moderately similar to transcription factor-like 5 [H.sapiens]	513822 IMAGE Consortium	AI595248
5290 H	7 021PAMMQ	ESTs, Highly similar to NADH-UBIQUINONE OXIDOREDUCTASE B9 SUBUNIT [Bos taurus]	514016 IMAGE Consortium	AA062281
5291 H	9 021PAMMQ	ESTs	514386 IMAGE Consortium	AI595145

5292 H	11 021PAMMQ	ESTs	514544 IMAGE Consortium	AA066063
5293 B	1 021WAMMR	neighbor of A-kinase anchoring protein 95	514620 IMAGE Consortium	AA066102
5294 B	3 021WAMMR	proacrosin binding protein	515524 IMAGE Consortium	AA065355
5295 B	5 021WAMMR	ESTs, Highly similar to T00377 KIAA0642 protein - human [H.sapiens]	515043 IMAGE Consortium	AI595285
5296 B	7 021WAMMR	RAS-related C3 botulinum substrate 1, guanine nucleotide exchange factor 1	516082 IMAGE Consortium	AI606688
5297 B	9 021WAMMR	cytochrome c oxidase, subunit VIIIc	515735 IMAGE Consortium	AA065490
5298 B	11 021WAMMR	ESTs, Moderately similar to proteasome inhibitor hPI31 subunit [H.sapiens]	515950 IMAGE Consortium	AA062408
5299 D	1 021WAMMR	ESTs	515377 IMAGE Consortium	AI595618
5300 D	3 021WAMMR	deformed epidermal autoregulatory factor 1 (Drosophila)	515533 IMAGE Consortium	AA065379
5301 D	5 021WAMMR	2,3-bisphosphoglycerate mutase	515091 IMAGE Consortium	AI326531
5302 D	7 021WAMMR	ESTs	516152 IMAGE Consortium	AI606703
5303 D	9 021WAMMR	protein kinase, cAMP dependent regulatory, type II alpha	515810 IMAGE Consortium	AI326534
5304 D	11 021WAMMR	integrin, alpha E, epithelial-associated	515982 IMAGE Consortium	AI326538
5305 F	1 021WAMMR	ornithine decarboxylase antizyme 3	515386 IMAGE Consortium	AA064494
5306 F	3 021WAMMR	Public domain EST	514823 IMAGE Consortium	
5307 F	5 021WAMMR			
5308 F	7 021WAMMR	TATA box binding protein	516268 IMAGE Consortium	AI386012
5309 F	9 021WAMMR	ESTs, Highly similar to AF151018_1 HSPC184 [H.sapiens]	515824 IMAGE Consortium	AI595526
5310 F	11 021WAMMR	SH3-domain GRB2-like B1 (endophilin)	517025 IMAGE Consortium	AA062457
5311 H	1 021WAMMR	ESTs	515404 IMAGE Consortium	AI595626
5312 H	3 021WAMMR			
5313 H	5 021WAMMR	receptor (calcitonin) activity modifying protein 3	515470 IMAGE Consortium	AA065905
5314 H	7 021WAMMR	tubulin, beta 5	516294 IMAGE Consortium	AI386014
5315 H	9 021WAMMR	ESTs	515835 IMAGE Consortium	AA066037
5316 H	11 021WAMMR	feminization 1 b homolog (C. elegans)	517112 IMAGE Consortium	AA062503
5317 B	1 0213AMMS	protein kinase, interferon inducible double stranded RNA dependent inhibitor	516569 IMAGE Consortium	AA105298
5318 B	3 0213AMMS	transforming growth factor, beta receptor II	517428 IMAGE Consortium	AI324400
5319 B	5 0213AMMS	ESTs, Moderately similar to PTD015 [H.sapiens]	517500 IMAGE Consortium	AA065573
5320 B	7 0213AMMS			
5321 B	9 0213AMMS	ESTs	517700 IMAGE Consortium	AA066484
5322 B	11 0213AMMS	E74-like factor 4 (ets domain transcription factor)	519082 IMAGE Consortium	AI597326
5323 D	1 0213AMMS	Fc receptor, IgG, low affinity III	516594 IMAGE Consortium	AA105303
5324 D	3 0213AMMS	peptide N-glycanase homolog (S.cerevisiae)	516714 IMAGE Consortium	AA108210
5325 D	5 0213AMMS	EST AA792894	516809 IMAGE Consortium	AI596622
5326 D	7 0213AMMS	ESTs, Highly similar to GATM_RAT GLYCINE AMIDINOTRANSFERASE PRECURSOR [R.norvegicus]	517556 IMAGE Consortium	AA066615
5327 D	9 0213AMMS	mago-nashi homolog, proliferation-associated (Drosophila)	517745 IMAGE Consortium	AA066510

5328 D	11 0213AMMS	ESTs, Moderately similar to TYROSINE-PROTEIN KINASE FRK [Homo sapiens]	519153 IMAGE Consortium	AA105999
5329 F	1 0213AMMS	quaking	517326 IMAGE Consortium	AA062440
5330 F	3 0213AMMS	ESTs, Weakly similar to CG5676 gene product [D.melanogaster]	517448 IMAGE Consortium	AA065566
5331 F	5 0213AMMS	ESTs	516834 IMAGE Consortium	AI596623
5332 F	7 0213AMMS	ESTs, Weakly similar to T00351 hypothetical protein KIAA0707 - human [H.sapiens]	517569 IMAGE Consortium	AI597110
5333 F	9 0213AMMS	beclin 1 (coiled-coil, myosin-like BCL2-interacting protein)	517749 IMAGE Consortium	AA066511
5334 F	11 0213AMMS	ESTs, Highly similar to hypothetical protein [H.sapiens]	519175 IMAGE Consortium	AA108165
5335 H	1 0213AMMS			
5336 H	3 0213AMMS	Public domain EST	517456 IMAGE Consortium	
5337 H	5 0213AMMS	solute carrier family 25 (mitochondrial carrier; peroxisomal membrane protein, 34 kDa), member 17	516869 IMAGE Consortium	AA062372
5338 H	7 0213AMMS	ESTs, Highly similar to ELONGATION FACTOR TU, MITOCHONDRIAL PRECURSOR [Bos taurus]	517589 IMAGE Consortium	AI892239
5339 H	9 0213AMMS	lectin, galactose binding, soluble 3	517763 IMAGE Consortium	AA066647
5340 H	11 0213AMMS	Mus musculus narf mRNA for neural activity-related ring finger protein, complete cds	519234 IMAGE Consortium	AI597336
5341 B	1 021AAMMT	phenylalanine hydroxylase	518706 IMAGE Consortium	AA105958
5342 B	3 021AAMMT	thymopoietin	518690 IMAGE Consortium	AA105756
5343 B	5 021AAMMT	hexosaminidase B	518913 IMAGE Consortium	AA106623
5344 B	7 021AAMMT	ESTs	519576 IMAGE Consortium	AA107109
5345 B	9 021AAMMT	reduced expression 3	520150 IMAGE Consortium	AA107928
5346 B	11 021AAMMT	crystallin, zeta	520244 IMAGE Consortium	AA107690
5347 D	1 021AAMMT	glutathione S-transferase, alpha 4	518754 IMAGE Consortium	AA105866
5348 D	3 021AAMMT	hnRNP-associated with lethal yellow	518737 IMAGE Consortium	AA105774
5349 D	5 021AAMMT	ESTs, Highly similar to GUANINE NUCLEOTIDE-BINDING PROTEIN G(I)/G(S)/G(O) GAMMA-5 SUBUNIT [Bos taurus; Rattus norvegicus]	518931 IMAGE Consortium	AA106171
5350 D	7 021AAMMT	guanylate cyclase activator 2b (retina)	519711 IMAGE Consortium	AA107035
5351 D	9 021AAMMT	growth arrest specific 2	520189 IMAGE Consortium	AA117554
5352 D	11 021AAMMT	acyl-CoA thioesterase 1, cytosolic	520295 IMAGE Consortium	AI552677
5353 F	1 021AAMMT	ESTs, Highly similar to unnamed protein product [H.sapiens]	518895 IMAGE Consortium	AA106618
5354 F	3 021AAMMT	glutathione peroxidase 3	518767 IMAGE Consortium	AI892736
5355 F	5 021AAMMT	ESTs, Highly similar to PUTATIVE 40S RIBOSOMAL PROTEIN YHR148W [Saccharomyces cerevisiae]	518977 IMAGE Consortium	AA106070
5356 F	7 021AAMMT	ESTs	519912 IMAGE Consortium	AA109539
5357 F	9 021AAMMT	ESTs	520198 IMAGE Consortium	AI552670
5358 F	11 021AAMMT	ESTs, Highly similar to TADA1 protein [H.sapiens]	521019 IMAGE Consortium	AA066757
5359 H	1 021AAMMT	X transporter protein 2	518916 IMAGE Consortium	AA106621
5360 H	3 021AAMMT			

5361 H	5 021AAMMT	kidney expressed gene 1	518981 IMAGE Consortium	AA106072
5362 H	7 021AAMMT	ESTs	519969 IMAGE Consortium	AI552654
5363 H	9 021AAMMT	ESTs	520283 IMAGE Consortium	AA110886
5364 H	11 021AAMMT	hemoglobin, beta adult major chain	521046 IMAGE Consortium	AA066763
5365 B	1 021HAMMU	trypsin 4	521185 IMAGE Consortium	AI386046
5366 B	3 021HAMMU	DNA segment, Chr 13, ERATO Doi 372, expressed	520714 IMAGE Consortium	AA066392
5367 B	5 021HAMMU	ESTs	442475 IMAGE Consortium	AA015108
5368 B	7 021HAMMU	ESTs	444903 IMAGE Consortium	AA016790
5369 B	9 021HAMMU	hydroxysteroid 17-beta dehydrogenase 4	523155 IMAGE Consortium	AA067285
5370 B	11 021HAMMU	small nuclear ribonucleoprotein B	522717 IMAGE Consortium	AA087675
5371 D	1 021HAMMU	ESTs, Highly similar to TUBULIN BETA CHAIN [Sus scrofa]	521222 IMAGE Consortium	AI386049
5372 D	3 021HAMMU	small inducible cytokine subfamily B (Cys-X-Cys), member 14	520740 IMAGE Consortium	AA066282
5373 D	5 021HAMMU	ESTs	443273 IMAGE Consortium	AA015506
5374 D	7 021HAMMU	ESTs	439237 IMAGE Consortium	AA003923
5375 D	9 021HAMMU	ESTs	523159 IMAGE Consortium	AA067295
5376 D	11 021HAMMU	cleavage and polyadenylation specificity factor 3	522766 IMAGE Consortium	AA087685
5377 F	1 021HAMMU	D-amino acid oxidase	520508 IMAGE Consortium	AA108226
5378 F	3 021HAMMU	NADPH oxidase 4	520782 IMAGE Consortium	AI552705
5379 F	5 021HAMMU	glutamine fructose-6-phosphate transaminase 1	443380 IMAGE Consortium	AI322480
5380 F	7 021HAMMU	syntaxin binding protein 1	439318 IMAGE Consortium	AA004089
5381 F	9 021HAMMU	ESTs	523179 IMAGE Consortium	AA067861
5382 F	11 021HAMMU	ESTs, Moderately similar to P120 PROTEIN [Mus musculus]	522797 IMAGE Consortium	AA087440
5383 H	1 021HAMMU			
5384 H	3 021HAMMU	protein that interacts with C kinase 1	520876 IMAGE Consortium	AI386037
5385 H	5 021HAMMU	prolactin-like protein F	443457 IMAGE Consortium	AA013842
5386 H	7 021HAMMU	meiosis-specific nuclear structural protein 1	439503 IMAGE Consortium	AA004193
5387 H	9 021HAMMU	ESTs	523183 IMAGE Consortium	AA067863
5388 H	11 021HAMMU	DNA segment, Chr 18, ERATO Doi 240, expressed	522799 IMAGE Consortium	AA087441
5389 B	1 021OAMMV	solute carrier family 22, member 1	522987 IMAGE Consortium	AA066406
5390 B	3 021OAMMV			
5391 B	5 021OAMMV	Ras suppressor protein 1	521255 IMAGE Consortium	AA066778
5392 B	7 021OAMMV	peroxiredoxin 3	521641 IMAGE Consortium	AA066332
5393 B	9 021OAMMV	ESTs, Highly similar to S57447 HPBR11-7 protein - human [H.sapiens]	523309 IMAGE Consortium	AI614602
5394 B	11 021OAMMV	seven in absentia 1A	523455 IMAGE Consortium	AA080210
5395 D	1 021OAMMV	topoisomerase (DNA) II beta	522994 IMAGE Consortium	AA066404
5396 D	3 021OAMMV	ESTs, Weakly similar to salivary proline-rich protein [R.norvegicus]	521844 IMAGE Consortium	AI552745
5397 D	5 021OAMMV	S-adenosylmethionine decarboxylase 1	522029 IMAGE Consortium	AI552765
5398 D	7 021OAMMV	carbonic anhydrase 3	521648 IMAGE Consortium	AI386062

5399 D	9 021OAMMV	ESTs, Highly similar to MYELIN P0 PROTEIN PRECURSOR [Mus musculus]	523321 IMAGE Consortium	AA080181
5400 D	11 021OAMMV	syntaxin binding protein 3	523484 IMAGE Consortium	AA097241
5401 F	1 021OAMMV	ESTs, Weakly similar to POSSIBLE GLOBAL TRANSCRIPTION ACTIVATOR SNF2L [Caenorhabditis elegans]	523016 IMAGE Consortium	AI325620
5402 F	3 021OAMMV	cytochrome P450, 2c29	521856 IMAGE Consortium	AA106162
5403 F	5 021OAMMV	plasminogen	522078 IMAGE Consortium	AA106793
5404 F	7 021OAMMV	glucose-6-phosphatase, catalytic	523120 IMAGE Consortium	AI326575
5405 F	9 021OAMMV	ESTs, Weakly similar to Unknown [H.sapiens]	523369 IMAGE Consortium	AI614610
5406 F	11 021OAMMV	tumor necrosis factor receptor superfamily, member 1a	523585 IMAGE Consortium	AA097343
5407 H	1 021OAMMV	ESTs	522343 IMAGE Consortium	AI552778
5408 H	3 021OAMMV	ESTs, Weakly similar to HP33 [R.norvegicus]	521860 IMAGE Consortium	AA106164
5409 H	5 021OAMMV	phosphoenolpyruvate carboxykinase 1, cytosolic	522080 IMAGE Consortium	AA106128
5410 H	7 021OAMMV	histocompatibility 2, T region locus 10	523150 IMAGE Consortium	AA067069
5411 H	9 021OAMMV	ESTs	523399 IMAGE Consortium	AA080065
5412 H	11 021OAMMV	ESTs	522911 IMAGE Consortium	AA067214
5413 B	1 021VAMMW	DNA segment, Chr 10, ERATO Doi 667, expressed	523000 IMAGE Consortium	AA067217
5414 B	3 021VAMMW	ESTs, Highly similar to NEUROLYSIN PRECURSOR [Rattus norvegicus]	524208 IMAGE Consortium	AI386253
5415 B	5 021VAMMW	ESTs	523718 IMAGE Consortium	AA067317
5416 B	7 021VAMMW	ESTs, Highly similar to ENOYL-COA HYDRATASE, MITOCHONDRIAL PRECURSOR [Rattus norvegicus]	523990 IMAGE Consortium	AA067736
5417 B	9 021VAMMW	high mobility group nucleosomal binding domain 2	524656 IMAGE Consortium	AA068490
5418 B	11 021VAMMW	ESTs	524882 IMAGE Consortium	AA097074
5419 D	1 021VAMMW	ESTs	523054 IMAGE Consortium	AI552859
5420 D	3 021VAMMW	ESTs, Weakly similar to CG8009 gene product [D.melanogaster]	524237 IMAGE Consortium	AA065669
5421 D	5 021VAMMW	FK506 binding protein 1a (12 kDa)	523746 IMAGE Consortium	AA067353
5422 D	7 021VAMMW	DNA segment, Chr 6, Wayne State University 157, expressed	524445 IMAGE Consortium	AI614939
5423 D	9 021VAMMW	FK506 binding protein 9	524686 IMAGE Consortium	AA068499
5424 D	11 021VAMMW	RAS p21 protein activator 3	524952 IMAGE Consortium	AA089049
5425 F	1 021VAMMW			
5426 F	3 021VAMMW	Mus musculus BAC 171m12 MESDC1 (Mesdc1) and MESDC2 (Mesdc2) genes, complete cds	524258 IMAGE Consortium	AA065676
5427 F	5 021VAMMW	activator of S phase kinase	523810 IMAGE Consortium	AA067144
5428 F	7 021VAMMW	nuclear factor I/B	524471 IMAGE Consortium	AA068096
5429 F	9 021VAMMW	solute carrier family 30 (zinc transporter), member 4	524721 IMAGE Consortium	AA067699
5430 F	11 021VAMMW	pantophysin	524978 IMAGE Consortium	AI325652
5431 H	1 021VAMMW	kidney cell line derived transcript 1	524020 IMAGE Consortium	AA072021
5432 H	3 021VAMMW	ESTs, Weakly similar to FGD1_MOUSE PUTATIVE RHO/RAC GUANINE NUCLEOTIDE EXCHANGE	524262 IMAGE Consortium	AA065678

		FACTOR [M.musculus]		
5433 H	5 021VAMMW	high mobility group AT-hook 1	523831 IMAGE Consortium	AA067083
5434 H	7 021VAMMW	ESTs, Moderately similar to AF151841_1 CGI-83 protein [H.sapiens]	524509 IMAGE Consortium	AI614942
5435 H	9 021VAMMW	proteolipid protein (myelin)	524747 IMAGE Consortium	AA067778
5436 H	11 021VAMMW	ESTs	524057 IMAGE Consortium	AA066245
5437 B	1 0212AMMX	coatamer protein complex, subunit zeta 2	524158 IMAGE Consortium	AA066137
5438 B	3 0212AMMX	amyloid beta (A4) precursor protein	524351 IMAGE Consortium	AI325647
5439 B	5 0212AMMX	palate, lung, and nasal epithelium expressed transcript	524664 IMAGE Consortium	AA068625
5440 B	7 0212AMMX	ESTs, Weakly similar to ER72_MOUSE PROTEIN DISULFIDE ISOMERASE-RELATED PROTEIN ERP-72 PRECURSOR [M.musculus]	533239 IMAGE Consortium	AA068784
5441 B	9 0212AMMX	ribosomal protein L5	524892 IMAGE Consortium	AI892334
5442 B	11 0212AMMX	ESTs, Highly similar to KIAA0157 gene product is novel. [H.sapiens]	533276 IMAGE Consortium	AI551980
5443 D	1 0212AMMX			
5444 D	3 0212AMMX	AXL receptor tyrosine kinase	524391 IMAGE Consortium	AA065764
5445 D	5 0212AMMX	ESTs	524713 IMAGE Consortium	AA068198
5446 D	7 0212AMMX	ESTs, Moderately similar to 2ACA_HUMAN SERINE/THREONINE PROTEIN PHOSPHATASE 2A, 72/130 KDA REGULATORY SUBUNIT B [H.sapiens]	532703 IMAGE Consortium	AA068827
5447 D	9 0212AMMX	antigen identified by monoclonal antibodies 4F2	524901 IMAGE Consortium	AA088974
5448 D	11 0212AMMX	potassium large conductance pH-sensitive channel, subfamily M, alpha member 3	534033 IMAGE Consortium	AA072586
5449 F	1 0212AMMX	ESTs, Moderately similar to hypothetical protein [H.sapiens]	524225 IMAGE Consortium	AA065657
5450 F	3 0212AMMX			
5451 F	5 0212AMMX	ESTs	524737 IMAGE Consortium	AA068204
5452 F	7 0212AMMX	kynurenine aminotransferase II	524821 IMAGE Consortium	AA096815
5453 F	9 0212AMMX			
5454 F	11 0212AMMX			
5455 H	1 0212AMMX	U1 small nuclear ribonucleoprotein 70 kDa polypeptide A	524255 IMAGE Consortium	AA067888
5456 H	3 0212AMMX	phosphatidylinositol transfer protein	524427 IMAGE Consortium	AA067802
5457 H	5 0212AMMX	ESTs	524761 IMAGE Consortium	AA068211
5458 H	7 0212AMMX	ESTs, Weakly similar to CG3337 gene product [D.melanogaster]	524825 IMAGE Consortium	AA088931
5459 H	9 0212AMMX	tissue specific transplantation antigen P35B	524947 IMAGE Consortium	AA096862
5460 H	11 0212AMMX			
5461 B	1 0219AMMY			
5462 B	3 0219AMMY	lipopolysaccharide binding protein	533799 IMAGE Consortium	AA071848
5463 B	5 0219AMMY	ESTs	550855 IMAGE Consortium	AA098600
5464 B	7 0219AMMY	ESTs	536664 IMAGE Consortium	AI550053
5465 B	9 0219AMMY	ESTs, Weakly similar to CG14130 gene product	535538 IMAGE Consortium	AA087378

		[D.melanogaster]		
5466 B	11 0219AMMY	crystallin, alpha 2	535934 IMAGE Consortium	AI325730
5467 D	1 0219AMMY	Bruton agammaglobulinemia tyrosine kinase	533494 IMAGE Consortium	AA105092
5468 D	3 0219AMMY	protein-tyrosine sulfotransferase 2	533826 IMAGE Consortium	AA072292
5469 D	5 0219AMMY	ESTs	536374 IMAGE Consortium	AA073925
5470 D	7 0219AMMY	ESTs	535025 IMAGE Consortium	AA073305
5471 D	9 0219AMMY	lymphocyte antigen 6 complex	534886 IMAGE Consortium	AI325697
5472 D	11 0219AMMY	thymidylate synthase	477222 IMAGE Consortium	AI323028
5473 F	1 0219AMMY	lysophosphatidic acid phosphatase	534227 IMAGE Consortium	AA072749
5474 F	3 0219AMMY	cyclin F	533922 IMAGE Consortium	AI326604
5475 F	5 0219AMMY	phosphoprotein enriched in astrocytes 15	536520 IMAGE Consortium	AA073869
5476 F	7 0219AMMY	IQ motif containing GTPase activating protein 1	535030 IMAGE Consortium	AI552087
5477 F	9 0219AMMY	ESTs, Moderately similar to NADH-UBIQUINONE OXIDOREDUCTASE KFYI SUBUNIT PRECURSOR [Bos taurus]	535634 IMAGE Consortium	AA110846
5478 F	11 0219AMMY	ESTs	475646 IMAGE Consortium	AA050315
5479 H	1 0219AMMY	ESTs, Moderately similar to unnamed protein product [H.sapiens]	534253 IMAGE Consortium	AA072758
5480 H	3 0219AMMY	cyclin-dependent kinase inhibitor 1A (P21)	533961 IMAGE Consortium	AI326605
5481 H	5 0219AMMY	ESTs	536534 IMAGE Consortium	AA073778
5482 H	7 0219AMMY	ESTs	535051 IMAGE Consortium	AA073331
5483 H	9 0219AMMY	ribosomal protein, large, P1	534918 IMAGE Consortium	AA073266
5484 H	11 0219AMMY	hypothetical protein, MNCb-1301	477180 IMAGE Consortium	AI510023
5485 B	1 021GAMMZ	guanine nucleotide binding protein, beta 5	478224 IMAGE Consortium	AI323039
5486 B	3 021GAMMZ	epimorphin	479304 IMAGE Consortium	AA048759
5487 B	5 021GAMMZ	origin recognition complex, subunit 4	572595 IMAGE Consortium	AA110785
5488 B	7 021GAMMZ	solute carrier family 7 (cationic amino acid transporter, y+ system), member 9	572009 IMAGE Consortium	AI893687
5489 B	9 021GAMMZ	suppressor of clear, C. elegans, homolog of	573245 IMAGE Consortium	AA119314
5490 B	11 021GAMMZ	ESTs	480682 IMAGE Consortium	AA064133
5491 D	1 021GAMMZ	intercellular adhesion molecule 2	477574 IMAGE Consortium	AI323033
5492 D	3 021GAMMZ	ESTs	478690 IMAGE Consortium	AI510089
5493 D	5 021GAMMZ	ESTs	492430 IMAGE Consortium	AI595180
5494 D	7 021GAMMZ	ESTs, Highly similar to KIAA0960 protein [H.sapiens]	571314 IMAGE Consortium	AI587761
5495 D	9 021GAMMZ	GrpE-like 2, mitochondrial	482996 IMAGE Consortium	AA060861
5496 D	11 021GAMMZ	cytochrome c oxidase, subunit VI a, polypeptide 2	480732 IMAGE Consortium	AI892132
5497 F	1 021GAMMZ	ESTs, Weakly similar to integral inner nuclear membrane protein MAN1 [H.sapiens]	478373 IMAGE Consortium	AI510071
5498 F	3 021GAMMZ	splicing factor 3a, subunit 2, 66kD	478715 IMAGE Consortium	AI323047
5499 F	5 021GAMMZ	ESTs	572815 IMAGE Consortium	AI587855
5500 F	7 021GAMMZ	proviral integration site 1	571396 IMAGE Consortium	AA108991
5501 F	9 021GAMMZ	nephronophthisis 1 (juvenile) homolog (human)	483702 IMAGE Consortium	AA061347
5502 F	11 021GAMMZ	ESTs, Moderately similar to SENR [R.norvegicus]	482207 IMAGE Consortium	AI385474

5503 H	1 021GAMMZ			
5504 H	3 021GAMMZ	tachykinin 2	478732 IMAGE Consortium	AA051164
5505 H	5 021GAMMZ	musculin	493457 IMAGE Consortium	AI595213
5506 H	7 021GAMMZ	ESTs	571457 IMAGE Consortium	AA105367
5507 H	9 021GAMMZ	somatostatin	480070 IMAGE Consortium	AA051655
5508 H	11 021GAMMZ	cyclin-dependent kinase 5	480815 IMAGE Consortium	AI385450
5509 B	1 021GAMNO	ESTs	482482 IMAGE Consortium	AA063830
5510 B	3 021GAMNO	intestinal cell kinase	574243 IMAGE Consortium	AI530567
5511 B	5 021GAMNO	ESTs	573869 IMAGE Consortium	AA119196
5512 B	7 021GAMNO	ESTs	575526 IMAGE Consortium	AA125010
5513 B	9 021GAMNO	Mouse complement factor H-related protein mRNA, complete cds, clone 9C4	987791 IMAGE Consortium	AA570965
5514 B	11 021GAMNO	4-hydroxyphenylpyruvic acid dioxygenase	1001085 IMAGE Consortium	AA771396
5515 D	1 021GAMNO	hypothetical protein, MNCb-0385	482552 IMAGE Consortium	AA063845
5516 D	3 021GAMNO	ESTs, Moderately similar to 6-pyruvoyl-tetrahydropterin synthase [M.musculus]	575013 IMAGE Consortium	AI530537
5517 D	5 021GAMNO	ESTs	576027 IMAGE Consortium	AI530741
5518 D	7 021GAMNO	ESTs	574814 IMAGE Consortium	AI530505
5519 D	9 021GAMNO	angiotensinogen	987842 IMAGE Consortium	AA571053
5520 D	11 021GAMNO	troponin C, fast skeletal	1001275 IMAGE Consortium	AA771566
5521 F	1 021GAMNO	N-acetylglucosamine kinase	482594 IMAGE Consortium	AA061835
5522 F	3 021GAMNO	ESTs	574315 IMAGE Consortium	AI530580
5523 F	5 021GAMNO	transformation related protein 53	576063 IMAGE Consortium	AI323249
5524 F	7 021GAMNO	ESTs	576341 IMAGE Consortium	AI551483
5525 F	9 021GAMNO	ESTs	987869 IMAGE Consortium	AA571152
5526 F	11 021GAMNO	ESTs, Weakly similar to DIA1_MOUSE DIAPHANOUS PROTEIN HOMOLOG 1 [M.musculus]	1002051 IMAGE Consortium	AA684309
5527 H	1 021GAMNO	ESTs, Weakly similar to unnamed protein product [H.sapiens]	483301 IMAGE Consortium	AI594964
5528 H	3 021GAMNO	ESTs	575096 IMAGE Consortium	AI530552
5529 H	5 021GAMNO	ESTs, Weakly similar to I49636 DNA-binding protein - mouse [M.musculus]	575358 IMAGE Consortium	AI530661
5530 H	7 021GAMNO	ESTs, Weakly similar to Ras-related protein [R.norvegicus]	576372 IMAGE Consortium	AI551485
5531 H	9 021GAMNO	caseinolytic protease X (E.coli)	987445 IMAGE Consortium	AA572634
5532 H	11 021GAMNO	ESTs, Weakly similar to AF101779_1 Ring1 interactor RYBP [M.musculus]	1001366 IMAGE Consortium	AA771616
5533 B	1 021NAMN1	ESTs	1001468 IMAGE Consortium	AI587949
5534 B	3 021NAMN1	ESTs	1001718 IMAGE Consortium	AI587984
5535 B	5 021NAMN1	ESTs	1002621 IMAGE Consortium	AA596768
5536 B	7 021NAMN1	ESTs	1021283 IMAGE Consortium	AA636198
5537 B	9 021NAMN1	ESTs	1020816 IMAGE Consortium	AA645769
5538 B	11 021NAMN1	ESTs, Highly similar to dJ487J7.1.1 [H.sapiens]	1002200 IMAGE Consortium	AA608118



5539 D	1	021NAMN1	ESTs	1001475	IMAGE Consortium	AI587952
5540 D	3	021NAMN1	ESTs	1001733	IMAGE Consortium	AI587986
5541 D	5	021NAMN1	B-cell translocation gene 2, anti-proliferative	1002668	IMAGE Consortium	AA596779
5542 D	7	021NAMN1	ESTs	1020565	IMAGE Consortium	AA647876
5543 D	9	021NAMN1	ESTs, Highly similar to AF175407_1 selenocysteine lyase SCLY [M.musculus]	1020846	IMAGE Consortium	AI594179
5544 D	11	021NAMN1	ESTs, Moderately similar to S12207 hypothetical protein [M.musculus]	1002209	IMAGE Consortium	AA596431
5545 F	1	021NAMN1	heterogeneous nuclear ribonucleoprotein C	1001531	IMAGE Consortium	AI587961
5546 F	3	021NAMN1	ESTs, Highly similar to ADIPOCYTE P27 PROTEIN [Mus musculus]	1001768	IMAGE Consortium	AA666697
5547 F	5	021NAMN1	ESTs	1002731	IMAGE Consortium	AA607467
5548 F	7	021NAMN1	thymine DNA glycosylase	1020592	IMAGE Consortium	AI594148
5549 F	9	021NAMN1	ESTs	1020877	IMAGE Consortium	AA645931
5550 F	11	021NAMN1	ESTs, Moderately similar to KIAA0007 [H.sapiens]	1002230	IMAGE Consortium	AA596602
5551 H	1	021NAMN1	ESTs, Moderately similar to hypothetical protein [H.sapiens]	1001567	IMAGE Consortium	AI587963
5552 H	3	021NAMN1	ESTs	1001839	IMAGE Consortium	AA683979
5553 H	5	021NAMN1	proteoglycan, secretory granule	1002759	IMAGE Consortium	AA607483
5554 H	7	021NAMN1	ESTs, Moderately similar to dJ102H19.3 [H.sapiens]	1020635	IMAGE Consortium	AI594156
5555 H	9	021NAMN1	ESTs, Highly similar to KIAA1181 protein [H.sapiens]	1020934	IMAGE Consortium	AA646192
5556 H	11	021NAMN1	ESTs	1002266	IMAGE Consortium	AA596625
5557 B	1	021UAMN2	interferon gamma induced GTPase	1002441	IMAGE Consortium	AA596514
5558 B	3	021UAMN2	ESTs	1002699	IMAGE Consortium	AA607449
5559 B	5	021UAMN2	antigen identified by monoclonal antibody Ki 67	1038700	IMAGE Consortium	AA671317
5560 B	7	021UAMN2	ESTs	1023308	IMAGE Consortium	AA673522
5561 B	9	021UAMN2	neoplastic progression 3	1023863	IMAGE Consortium	AA866949
5562 B	11	021UAMN2	platelet derived growth factor receptor, beta polypeptide	1038854	IMAGE Consortium	AA624552
5563 D	1	021UAMN2	ESTs	1002458	IMAGE Consortium	AA596518
5564 D	3	021UAMN2	ESTs	1002745	IMAGE Consortium	AA607478
5565 D	5	021UAMN2	selenoprotein R	1038730	IMAGE Consortium	AA671340
5566 D	7	021UAMN2	ESTs	1023338	IMAGE Consortium	AA646795
5567 D	9	021UAMN2	ESTs, Weakly similar to thyroid hormone receptor-associated protein complex component TRAP150 [H.sapiens]	1023948	IMAGE Consortium	AA798799
5568 D	11	021UAMN2	SRY-box containing gene 4	1038880	IMAGE Consortium	AA624572
5569 F	1	021UAMN2	ESTs, Highly similar to p53 regulated PA26-T2 nuclear protein [H.sapiens]	1002471	IMAGE Consortium	AA596529
5570 F	3	021UAMN2	TGFB inducible early growth response	1021027	IMAGE Consortium	AA646509
5571 F	5	021UAMN2	actin, alpha, vascular smooth muscle	1038801	IMAGE Consortium	AA624460
5572 F	7	021UAMN2	ESTs	1023512	IMAGE Consortium	AA647195
5573 F	9	021UAMN2	DNA segment, Chr 19, Wayne State University 162, expressed	1039128	IMAGE Consortium	AA671440

5574 F	11 021UAMN2	solute carrier family 4 (anion exchanger), member 1	1039627 IMAGE Consortium	AA792268
5575 H	1 021UAMN2	nucleoredoxin	1002490 IMAGE Consortium	AA596534
5576 H	3 021UAMN2	solute carrier family 8 (sodium/calcium exchanger), member 1	1021099 IMAGE Consortium	AA645954
5577 H	5 021UAMN2	ESTs, Weakly similar to Yhr116wp [ <i>S.cerevisiae</i> ]	1038842 IMAGE Consortium	AA624547
5578 H	7 021UAMN2	ESTs	1023591 IMAGE Consortium	AI647179
5579 H	9 021UAMN2	ESTs	1039178 IMAGE Consortium	AA624615
5580 H	11 021UAMN2	adenylosuccinate synthetase 1, muscle	1038959 IMAGE Consortium	AA624853
5581 B	1 0211AMN3	homeo box, msh-like 1	1040034 IMAGE Consortium	AA612115
5582 B	3 0211AMN3			
5583 B	5 0211AMN3	troponin I, skeletal, fast 2	1077551 IMAGE Consortium	AA821941
5584 B	7 0211AMN3	NAD(P) dependent steroid dehydrogenase-like	1077729 IMAGE Consortium	AA822050
5585 B	9 0211AMN3	ESTs, Weakly similar to MSI [ <i>M.musculus</i> ]	1077485 IMAGE Consortium	AI594825
5586 B	11 0211AMN3	ubiquitin-conjugating enzyme E2H	1040225 IMAGE Consortium	AA792298
5587 D	1 0211AMN3	ESTs, Moderately similar to unnamed protein product [ <i>H.sapiens</i> ]	1040045 IMAGE Consortium	AA615213
5588 D	3 0211AMN3	cytochrome P450, steroid inducible 3a11	1091786 IMAGE Consortium	AA647336
5589 D	5 0211AMN3	c-src tyrosine kinase	1077565 IMAGE Consortium	AA822071
5590 D	7 0211AMN3	mitogen-activated protein kinase 8 interacting protein 3	1077738 IMAGE Consortium	AA822046
5591 D	9 0211AMN3	tubulin alpha 4	1077532 IMAGE Consortium	AA821924
5592 D	11 0211AMN3	ESTs, Highly similar to unnamed protein product [ <i>H.sapiens</i> ]	1040323 IMAGE Consortium	AA612147
5593 F	1 0211AMN3	small protein effector 1 of Cdc42	1040115 IMAGE Consortium	AA615252
5594 F	3 0211AMN3	syndecan 1	1077399 IMAGE Consortium	AA733479
5595 F	5 0211AMN3	ESTs	1077587 IMAGE Consortium	AI507907
5596 F	7 0211AMN3	cytochrome P450, 17	1077759 IMAGE Consortium	AA822113
5597 F	9 0211AMN3	epoxide hydrolase 1, microsomal	1077557 IMAGE Consortium	AA822067
5598 F	11 0211AMN3	Mus musculus putative chitinase precursor (YNL) mRNA, partial cds	1040515 IMAGE Consortium	AA612413
5599 H	1 0211AMN3	cytokine inducible SH2-containing protein	1040166 IMAGE Consortium	AA615275
5600 H	3 0211AMN3	vesicle-associated membrane protein, associated protein B and C	1077423 IMAGE Consortium	AA733495
5601 H	5 0211AMN3	Fc receptor, IgG, alpha chain transporter	1077599 IMAGE Consortium	AA821960
5602 H	7 0211AMN3	transthyretin	1077765 IMAGE Consortium	AA822116
5603 H	9 0211AMN3	glutaryl-Coenzyme A dehydrogenase	1077580 IMAGE Consortium	AA821944
5604 H	11 0211AMN3	tuftelin 1	1040522 IMAGE Consortium	AA612327
5605 B	1 0218AMN4	keratin complex 2, basic, gene 19	1040670 IMAGE Consortium	AA606310
5606 B	3 0218AMN4	murinoglobulin 1	1093220 IMAGE Consortium	AA675398
5607 B	5 0218AMN4	phosphatidic acid phosphatase 2a	1092160 IMAGE Consortium	AA670943
5608 B	7 0218AMN4	calpastatin	1139847 IMAGE Consortium	AA646105
5609 B	9 0218AMN4	ESTs, Highly similar to SKD1 PROTEIN [ <i>Mus musculus</i> ]	1140210 IMAGE Consortium	AA645634
5610 B	11 0218AMN4	Niemann Pick type C1	1148532 IMAGE Consortium	AA672765
5611 D	1 0218AMN4	ESTs	1040703 IMAGE Consortium	AA606340

5612 D	3 0218AMN4	ESTs	1092651 IMAGE Consortium	AA674176
5613 D	5 0218AMN4	Fas death domain-associated protein	1139624 IMAGE Consortium	AA646003
5614 D	7 0218AMN4	T-cell receptor beta, variable 13	1148698 IMAGE Consortium	AA791518
5615 D	9 0218AMN4	ESTs	1140294 IMAGE Consortium	AA645878
5616 D	11 0218AMN4	hypothetical protein 425O18-1	1148536 IMAGE Consortium	AA672767
5617 F	1 0218AMN4	ESTs	1040719 IMAGE Consortium	AA606427
5618 F	3 0218AMN4	ESTs, Weakly similar to similar to vacuolar biogenesis protein [C.elegans]	1092727 IMAGE Consortium	AA674326
5619 F	5 0218AMN4	inter-alpha trypsin inhibitor, heavy chain 1	1092207 IMAGE Consortium	AA755277
5620 F	7 0218AMN4	ESTs	1139971 IMAGE Consortium	AA646434
5621 F	9 0218AMN4	Public domain EST	1139604 IMAGE Consortium	
5622 F	11 0218AMN4	eukaryotic translation initiation factor 4E	1148547 IMAGE Consortium	AA672783
5623 H	1 0218AMN4	polymerase, gamma	1040750 IMAGE Consortium	AA606723
5624 H	3 0218AMN4	major urinary protein 2	1092779 IMAGE Consortium	AA674270
5625 H	5 0218AMN4	ESTs	1092223 IMAGE Consortium	AA755374
5626 H	7 0218AMN4	alcohol dehydrogenase family 1, subfamily A2	1148727 IMAGE Consortium	AA791665
5627 H	9 0218AMN4	ESTs, Moderately similar to I38909 damage-specific DNA binding protein 2 - human [H.sapiens]	1139673 IMAGE Consortium	AI592870
5628 H	11 0218AMN4	isocitrate dehydrogenase 1 (NADP+), soluble	1148559 IMAGE Consortium	AA672788
5629 B	1 021FAMN5			
5630 B	3 021FAMN5			
5631 B	5 021FAMN5			
5632 B	7 021FAMN5	ESTs	1148887 IMAGE Consortium	AI644881
5633 B	9 021FAMN5			
5634 B	11 021FAMN5	ESTs, Weakly similar to ankyrin 3 [M.musculus]	1149155 IMAGE Consortium	AI644933
5635 D	1 021FAMN5			
5636 D	3 021FAMN5			
5637 D	5 021FAMN5	DnaJ-like protein 1	1149747 IMAGE Consortium	AA667623
5638 D	7 021FAMN5	Sel1 (suppressor of lin-12) 1 homolog (C. elegans)	1148899 IMAGE Consortium	AA655874
5639 D	9 021FAMN5	ESTs, Weakly similar to unnamed protein product [H.sapiens]	1149068 IMAGE Consortium	AI644912
5640 D	11 021FAMN5	ESTs, Moderately similar to putative [C.elegans]	1149163 IMAGE Consortium	AA655810
5641 F	1 021FAMN5			
5642 F	3 021FAMN5			
5643 F	5 021FAMN5	ESTs	1149792 IMAGE Consortium	AA672740
5644 F	7 021FAMN5	ubiquitin-conjugating enzyme E2I	1148947 IMAGE Consortium	AA655553
5645 F	9 021FAMN5	Mus musculus synembryon mRNA, complete cds	1149083 IMAGE Consortium	AA655904
5646 F	11 021FAMN5	tryptophan hydroxylase	1149189 IMAGE Consortium	AA655597
5647 H	1 021FAMN5			
5648 H	3 021FAMN5			
5649 H	5 021FAMN5	lymphocyte antigen 84	1149822 IMAGE Consortium	AA795621
5650 H	7 021FAMN5	ESTs, Moderately similar to AF161518_1 HSPC169 [H.sapiens]	1148957 IMAGE Consortium	AA655568

5651 H	9 021FAMN5	ESTs	1149090 IMAGE Consortium	AI644918
5652 H	11 021FAMN5	ESTs, Highly similar to PROBABLE RIBOSOMAL PROTEIN B0303.15 IN CHROMOSOME III [Caenorhabditis elegans]	1149589 IMAGE Consortium	AA655922
5653 B	1 021MAMN6	ESTs	1149790 IMAGE Consortium	AA674114
5654 B	3 021MAMN6	ubiquitin specific protease 2	1150462 IMAGE Consortium	AA930268
5655 B	5 021MAMN6	transcriptional regulator, SIN3B (yeast)	1193024 IMAGE Consortium	AA711605
5656 B	7 021MAMN6	Mus musculus unknown mRNA	1193455 IMAGE Consortium	AA726858
5657 B	9 021MAMN6	vascular endothelial growth factor C	1193606 IMAGE Consortium	AA726944
5658 B	11 021MAMN6	ESTs	1193725 IMAGE Consortium	AA727067
5659 D	1 021MAMN6	ESTs, Moderately similar to KIAA1398 protein [H.sapiens]	1149877 IMAGE Consortium	AI645058
5660 D	3 021MAMN6	peroxisomal assembly protein 3	1150532 IMAGE Consortium	AA930433
5661 D	5 021MAMN6	guanosine diphosphate (GDP) dissociation inhibitor 1	1193032 IMAGE Consortium	AA711609
5662 D	7 021MAMN6	phosphatidylinositol 3-kinase, regulatory subunit, polypeptide 1 (p85 alpha)	1193479 IMAGE Consortium	AA726880
5663 D	9 021MAMN6	ESTs	1193626 IMAGE Consortium	AA727038
5664 D	11 021MAMN6	DNA segment, Chr 11, ERATO Doi 636, expressed	1209194 IMAGE Consortium	AA726101
5665 F	1 021MAMN6	ESTs, Moderately similar to DEOXYCYTIDYLATE DEAMINASE [Homo sapiens]	1149947 IMAGE Consortium	AA880054
5666 F	3 021MAMN6	POZ/zinc finger transcription factor	1150564 IMAGE Consortium	AA930334
5667 F	5 021MAMN6	ESTs, Highly similar to phosphodiesterase 3B [M.musculus]	1193062 IMAGE Consortium	AA711628
5668 F	7 021MAMN6	signal transducer and activator of transcription 1	1193486 IMAGE Consortium	AA726873
5669 F	9 021MAMN6	myeloid leukemia factor 1	1193633 IMAGE Consortium	AA727051
5670 F	11 021MAMN6	ESTs	1193735 IMAGE Consortium	AA727070
5671 H	1 021MAMN6	actin-related protein 1 homolog A (yeast)	1150065 IMAGE Consortium	AA791769
5672 H	3 021MAMN6	ESTs, Weakly similar to CG10978 gene product [D.melanogaster]	1150613 IMAGE Consortium	AA930455
5673 H	5 021MAMN6	SKAP55 homologue	1193180 IMAGE Consortium	AA718433
5674 H	7 021MAMN6	immediate early response 2	1193520 IMAGE Consortium	AA727017
5675 H	9 021MAMN6	ESTs, Weakly similar to I55595 splicing factor - human [H.sapiens]	1193652 IMAGE Consortium	AA726962
5676 H	11 021MAMN6	ESTs, Weakly similar to JAK3_MOUSE TYROSINE-PROTEIN KINASE JAK3 [M.musculus]	1209243 IMAGE Consortium	AA726152
5677 B	1 021TAMN7	ESTs, Highly similar to ALANINE AMINOTRANSFERASE [Rattus norvegicus]	1196506 IMAGE Consortium	AA794294
5678 B	3 021TAMN7	ESTs, Weakly similar to PARATHYMOSIN [Rattus norvegicus]	1197437 IMAGE Consortium	AA794319
5679 B	5 021TAMN7	ESTs, Weakly similar to KE4 [M.musculus]	1209153 IMAGE Consortium	AA726092
5680 B	7 021TAMN7	ESTs, Moderately similar to spindle pole body protein spc97 homolog GCP2 [H.sapiens]	1209972 IMAGE Consortium	AA727429
5681 B	9 021TAMN7	ESTs	1210300 IMAGE Consortium	AA727388
5682 B	11 021TAMN7	ESTs, Highly similar to AF250859_1 UDP-GlcNAc:a-3-D-mannoside b1,2-N-acetylglucosaminyltransferase I.2	1223192 IMAGE Consortium	AA734124

		[H.sapiens]		
5683 D	1 021TAMN7	Public domain EST	1196523 IMAGE Consortium	
5684 D	3 021TAMN7	ESTs	1198187 IMAGE Consortium	AA725941
5685 D	5 021TAMN7			
5686 D	7 021TAMN7	cadherin EGF LAG seven-pass G-type receptor 2	1210024 IMAGE Consortium	AA727503
5687 D	9 021TAMN7	ESTs	1210369 IMAGE Consortium	AA727485
5688 D	11 021TAMN7	uridine monophosphate synthetase	1223250 IMAGE Consortium	AA734206
5689 F	1 021TAMN7	ESTs	1198006 IMAGE Consortium	AA727098
5690 F	3 021TAMN7	ESTs	1196773 IMAGE Consortium	AA717080
5691 F	5 021TAMN7	matrix metalloproteinase 24	1209283 IMAGE Consortium	AA726203
5692 F	7 021TAMN7	interferon-induced protein with tetratricopeptide repeats 3	1210036 IMAGE Consortium	AA727519
5693 F	9 021TAMN7	MYC-associated zinc finger protein (purine-binding transcription factor)	1210473 IMAGE Consortium	AA727641
5694 F	11 021TAMN7	homeo box B13	1223369 IMAGE Consortium	AA734350
5695 H	1 021TAMN7	lymphoid blast crisis-like 1	1198038 IMAGE Consortium	AA725965
5696 H	3 021TAMN7	ESTs	1196814 IMAGE Consortium	AA733502
5697 H	5 021TAMN7	ESTs	1209307 IMAGE Consortium	AA726262
5698 H	7 021TAMN7	poly A binding protein, cytoplasmic 1	1210045 IMAGE Consortium	AA727531
5699 H	9 021TAMN7	mannoside acetylglucosaminyltransferase 1	1223023 IMAGE Consortium	AA733854
5700 H	11 021TAMN7	ESTs, Weakly similar to CG11900 gene product [D.melanogaster]	1223475 IMAGE Consortium	AA795793
5701 B	1 0210AMN8	endothelin receptor type B	1224368 IMAGE Consortium	AA755763
5702 B	3 0210AMN8	ESTs, Weakly similar to contains similarity to Oryctolagus cuniculus sarcolemmal associated protein-3 [C.elegans]	1210910 IMAGE Consortium	AA795744
5703 B	5 0210AMN8	myomesin 1	1222968 IMAGE Consortium	AA795463
5704 B	7 0210AMN8	ESTs	597022 IMAGE Consortium	AA138758
5705 B	9 0210AMN8	ESTs	597285 IMAGE Consortium	AA138293
5706 B	11 0210AMN8	ESTs	596799 IMAGE Consortium	AA146572
5707 D	1 0210AMN8	ESTs, Weakly similar to PRP3_MOUSE PROLINE-RICH PROTEIN MP-3 [M.musculus]	1224388 IMAGE Consortium	AA755781
5708 D	3 0210AMN8	neural proliferation, differentiation and control gene 1	1211659 IMAGE Consortium	AI425175
5709 D	5 0210AMN8	ESTs	1210343 IMAGE Consortium	AA727423
5710 D	7 0210AMN8	lymphocyte antigen 86	597065 IMAGE Consortium	AA138913
5711 D	9 0210AMN8	ESTs	597310 IMAGE Consortium	AA138485
5712 D	11 0210AMN8	ESTs	596846 IMAGE Consortium	AA145093
5713 F	1 0210AMN8	ESTs, Highly similar to T17237 hypothetical protein DKFZp434P106.1 - human [H.sapiens]	1223739 IMAGE Consortium	AA815761
5714 F	3 0210AMN8	troponin T1, skeletal, slow	1211707 IMAGE Consortium	AA656407
5715 F	5 0210AMN8	ESTs	1210415 IMAGE Consortium	AA727556
5716 F	7 0210AMN8	ESTs	597078 IMAGE Consortium	AI447910
5717 F	9 0210AMN8	ESTs	596629 IMAGE Consortium	AA146523
5718 F	11 0210AMN8			
5719 H	1 0210AMN8	DNA segment, Chr 11, ERATO Doi 99, expressed	1223764 IMAGE Consortium	AA815772

5720 H	3 0210AMN8	ESTs	1223064 IMAGE Consortium	AA733932
5721 H	5 0210AMN8	eukaryotic translation initiation factor 4E binding protein 2	1211217 IMAGE Consortium	AA655382
5722 H	7 0210AMN8	ESTs	596366 IMAGE Consortium	AA138861
5723 H	9 0210AMN8	ESTs	596632 IMAGE Consortium	AA138388
5724 H	11 0210AMN8	ESTs, Weakly similar to CG5676 gene product [D.melanogaster]	596956 IMAGE Consortium	AA145294
5725 B	1 0217AMN9	insulin-like growth factor 2 receptor	1224549 IMAGE Consortium	AA755961
5726 B	3 0217AMN9	cysteine-rich protein 3	1224764 IMAGE Consortium	AA756171
5727 B	5 0217AMN9	procollagen, type XVII, alpha 1	1227239 IMAGE Consortium	AA759995
5728 B	7 0217AMN9	ESTs	1227517 IMAGE Consortium	AA738860
5729 B	9 0217AMN9	ESTs, Weakly similar to CDNC_HUMAN CYCLIN-DEPENDENT KINASE INHIBITOR 1C [H.sapiens]	1227787 IMAGE Consortium	AA760220
5730 B	11 0217AMN9	ESTs	1228169 IMAGE Consortium	AA760260
5731 D	1 0217AMN9	ESTs, Weakly similar to AF148447_1 ubiquitin C-terminal hydrolase UCH37 [M.musculus]	1224595 IMAGE Consortium	AA756040
5732 D	3 0217AMN9	ubiquitous nuclear protein	1227221 IMAGE Consortium	AA759991
5733 D	5 0217AMN9	glycosylphosphatidylinositol 1 homolog (human)	1227296 IMAGE Consortium	AA756773
5734 D	7 0217AMN9	claudin 1	1227538 IMAGE Consortium	AA738911
5735 D	9 0217AMN9	epithelial membrane protein 3	1227794 IMAGE Consortium	AA760223
5736 D	11 0217AMN9	DNA segment, Chr 2, ERATO Doi 695, expressed	1228220 IMAGE Consortium	AA760323
5737 F	1 0217AMN9	ESTs	1224619 IMAGE Consortium	AA756099
5738 F	3 0217AMN9	splicing factor, arginine/serine-rich 3 (SRp20)	1227270 IMAGE Consortium	AA756753
5739 F	5 0217AMN9	Mus musculus Sep2 mRNA, partial cds	1227324 IMAGE Consortium	AA760001
5740 F	7 0217AMN9	ESTs, Highly similar to GAP JUNCTION ALPHA-1 PROTEIN [Mus musculus]	1227544 IMAGE Consortium	AA738914
5741 F	9 0217AMN9	fatty acid synthase	1227828 IMAGE Consortium	AA760287
5742 F	11 0217AMN9	ESTs	1228231 IMAGE Consortium	AI429838
5743 H	1 0217AMN9	myosin light chain, alkali, cardiac atria	1224644 IMAGE Consortium	AA756112
5744 H	3 0217AMN9	capping protein alpha 1	1227319 IMAGE Consortium	AA760009
5745 H	5 0217AMN9	lipoprotein lipase	1227378 IMAGE Consortium	AA739040
5746 H	7 0217AMN9	mutS homolog 2 (E. coli)	1227569 IMAGE Consortium	AA738946
5747 H	9 0217AMN9	ESTs, Weakly similar to A45841 T-complex-associated-testes-expressed-1 protein - mouse [M.musculus]	1227846 IMAGE Consortium	AA760305
5748 H	11 0217AMN9	decorin	1228255 IMAGE Consortium	AA755007
5749 B	1 021RAMNA	ESTs	1229593 IMAGE Consortium	AA763290
5750 B	3 021RAMNA	pleckstrin homology-like domain, family A, member 3	1230140 IMAGE Consortium	AA798939
5751 B	5 021RAMNA	Mus musculus Fas activated serine/threonine FAST kinase mRNA, complete cds	1260007 IMAGE Consortium	AA839998
5752 B	7 021RAMNA	ESTs, Highly similar to TRICARBOXYLATE TRANSPORT PROTEIN PRECURSOR [Rattus norvegicus]	1261007 IMAGE Consortium	AA840565
5753 B	9 021RAMNA	solute carrier family 12, member 2	1261456 IMAGE Consortium	AA856063
5754 B	11 021RAMNA	SNF1-like kinase	1260296 IMAGE Consortium	AA855274

5755 D	1	021RAMNA	ESTs	1229516	IMAGE Consortium	AA763211
5756 D	3	021RAMNA	ESTs	1230151	IMAGE Consortium	AA798960
5757 D	5	021RAMNA	DNA segment, Chr 6, ERATO Doi 286, expressed	1242913	IMAGE Consortium	AA790506
5758 D	7	021RAMNA	Mus musculus mRNA for Sil1 protein	1261117	IMAGE Consortium	AA838920
5759 D	9	021RAMNA	small optic lobes homolog (Drosophila)	1261483	IMAGE Consortium	AA856084
5760 D	11	021RAMNA	ESTs, Moderately similar to 2204390A synaptojanin [R.norvegicus]	1261036	IMAGE Consortium	AA838847
5761 F	1	021RAMNA	ESTs, Highly similar to HYPOTHETICAL MYELOID CELL LINE PROTEIN 5 [Homo sapiens]	1229586	IMAGE Consortium	AA763271
5762 F	3	021RAMNA	ESTs, Weakly similar to PTPL1-associated RhoGAP [H.sapiens]	1230170	IMAGE Consortium	AA798970
5763 F	5	021RAMNA	melastatin 1	1229490	IMAGE Consortium	AA763189
5764 F	7	021RAMNA	solute carrier family 4 (anion exchanger), member 3	1261294	IMAGE Consortium	AA855923
5765 F	9	021RAMNA	odd Oz/ten-m homolog 3 (Drosophila)	1260848	IMAGE Consortium	AA840175
5766 F	11	021RAMNA	ESTs	1260347	IMAGE Consortium	AI463946
5767 H	1	021RAMNA	ESTs, Moderately similar to /prediction	1229749	IMAGE Consortium	AA791119
5768 H	3	021RAMNA	smoothelin	1230178	IMAGE Consortium	AA798974
5769 H	5	021RAMNA	ESTs	1261528	IMAGE Consortium	AI465689
5770 H	7	021RAMNA	ESTs	1262007	IMAGE Consortium	AA874070
5771 H	9	021RAMNA	programmed cell death 6	1259879	IMAGE Consortium	AA855573
5772 H	11	021RAMNA	ESTs, Highly similar to KIAA0763 protein [H.sapiens]	1260374	IMAGE Consortium	AI463993
5773 B	1	021YAMNB	pleiotropic regulator 1, PRL1 homolog (Arabidopsis)	1260742	IMAGE Consortium	AA855784
5774 B	3	021YAMNB	2-hydroxyphytanoyl-CoA lyase	1262735	IMAGE Consortium	AA839264
5775 B	5	021YAMNB	purinergic receptor P2X, ligand-gated ion channel 4	1277150	IMAGE Consortium	AA882035
5776 B	7	021YAMNB	ESTs, Weakly similar to lama gene product [D.melanogaster]	1277607	IMAGE Consortium	AA880303
5777 B	9	021YAMNB	ESTs	329885	IMAGE Consortium	W14153
5778 B	11	021YAMNB	ESTs	330537	IMAGE Consortium	AI390824
5779 D	1	021YAMNB				
5780 D	3	021YAMNB				
5781 D	5	021YAMNB	cytochrome P450, 2c37	1277188	IMAGE Consortium	AA881013
5782 D	7	021YAMNB	calponin 1	1277623	IMAGE Consortium	AA880322
5783 D	9	021YAMNB	butyrate response factor 2	330120	IMAGE Consortium	AI327503
5784 D	11	021YAMNB	ESTs, Moderately similar to AF224508_1 sodium bicarbonate cotransporter isoform 3 kNBC-3 [M.musculus]	331470	IMAGE Consortium	W13854
5785 F	1	021YAMNB	polyglutamine binding protein 1	1262252	IMAGE Consortium	AA920161
5786 F	3	021YAMNB	keratin-associated protein 14	1262838	IMAGE Consortium	AI036534
5787 F	5	021YAMNB	ESTs, Weakly similar to EMR1_MOUSE CELL SURFACE GLYCOPROTEIN EMR1 PRECURSOR [M.musculus]	1277212	IMAGE Consortium	AA881032
5788 F	7	021YAMNB	ESTs	1277755	IMAGE Consortium	AA880559
5789 F	9	021YAMNB	ESTs	331036	IMAGE Consortium	AI390859
5790 F	11	021YAMNB	rad and gem related GTP-binding protein	330806	IMAGE Consortium	AI390842
5791 H	1	021YAMNB	ESTs	1262334	IMAGE Consortium	AA880501

5792 H	3 021YAMNB	ESTs, Weakly similar to AF148145_1 putative thymic stromal co-transporter TSCOT [M.musculus]	1276802 IMAGE Consortium	AA880757
5793 H	5 021YAMNB	ESTs, Weakly similar to CG11237 gene product [D.melanogaster]	1277368 IMAGE Consortium	AI466466
5794 H	7 021YAMNB	ESTs	1277834 IMAGE Consortium	AA882195
5795 H	9 021YAMNB	ESTs	331081 IMAGE Consortium	AI390862
5796 H	11 021YAMNB	ESTs, Weakly similar to heat shock factor binding protein 1 HSBP1 [H.sapiens]	330877 IMAGE Consortium	AI390850
5797 B	1 0215AMNC	ESTs	317745 IMAGE Consortium	W12952
5798 B	3 0215AMNC			
5799 B	5 0215AMNC			
5800 B	7 0215AMNC			
5801 B	9 0215AMNC			
5802 B	11 0215AMNC	zinc finger protein 260	1281179 IMAGE Consortium	AA799270
5803 D	1 0215AMNC	natriuretic peptide precursor type B	318539 IMAGE Consortium	W12084
5804 D	3 0215AMNC			
5805 D	5 0215AMNC			
5806 D	7 0215AMNC			
5807 D	9 0215AMNC			
5808 D	11 0215AMNC	ESTs, Weakly similar to syntaxin 7 [M.musculus]	1280541 IMAGE Consortium	AA789908
5809 F	1 0215AMNC	ESTs, Weakly similar to CABA_MOUSE CARG-BINDING FACTOR-A [M.musculus]	317892 IMAGE Consortium	W12955
5810 F	3 0215AMNC			
5811 F	5 0215AMNC			
5812 F	7 0215AMNC			
5813 F	9 0215AMNC			
5814 F	11 0215AMNC	expressed sequence tag mouse EST 34	1280674 IMAGE Consortium	AA790564
5815 H	1 0215AMNC	ESTs, Weakly similar to PROTEIN-GLUTAMINE GLUTAMYLTRANSFERASE E3 PRECURSOR [Homo sapiens]	317915 IMAGE Consortium	W12962
5816 H	3 0215AMNC			
5817 H	5 0215AMNC			
5818 H	7 0215AMNC			
5819 H	9 0215AMNC	ESTs	1279701 IMAGE Consortium	AA896036
5820 H	11 0215AMNC	src homology three (SH3) and cysteine rich domain	1280710 IMAGE Consortium	AA790586
5821 B	1 021CAMND	ESTs	1278113 IMAGE Consortium	AA882448
5822 B	3 021CAMND	ESTs	1278435 IMAGE Consortium	AA895221
5823 B	5 021CAMND	ESTs	1278704 IMAGE Consortium	AA895512
5824 B	7 021CAMND	glutathione S-transferase, alpha 3	1350040 IMAGE Consortium	AA982227
5825 B	9 021CAMND	ESTs	598846 IMAGE Consortium	AA162779
5826 B	11 021CAMND	molybdenum cofactor synthesis 2	598351 IMAGE Consortium	AI592114
5827 D	1 021CAMND	ESTs	1278163 IMAGE Consortium	AA882491
5828 D	3 021CAMND	ESTs	1279180 IMAGE Consortium	AA981343



5829 D	5 021CAMND	ESTs	1278738 IMAGE Consortium	AA895539
5830 D	7 021CAMND	N-myc (and STAT) interactor	1350110 IMAGE Consortium	AA982280
5831 D	9 021CAMND	ESTs	598882 IMAGE Consortium	AI592197
5832 D	11 021CAMND	ESTs	597626 IMAGE Consortium	AI592029
5833 F	1 021CAMND	ESTs, Weakly similar to acid ceramidase [M.musculus]	1278248 IMAGE Consortium	AA880821
5834 F	3 021CAMND	hyaluronan mediated motility receptor (RHAMM)	1279214 IMAGE Consortium	AA981371
5835 F	5 021CAMND			
5836 F	7 021CAMND	retinol binding protein 4, plasma	1350209 IMAGE Consortium	AA967219
5837 F	9 021CAMND	ESTs	598072 IMAGE Consortium	AA265210
5838 F	11 021CAMND	ESTs	598360 IMAGE Consortium	AA154872
5839 H	1 021CAMND	ESTs	1279000 IMAGE Consortium	AA896641
5840 H	3 021CAMND	proteinase 3	1279267 IMAGE Consortium	AA982151
5841 H	5 021CAMND			
5842 H	7 021CAMND	Mus musculus glutathione peroxidase (Gpx2) gene, complete cds	1350332 IMAGE Consortium	AA980349
5843 H	9 021CAMND	ESTs	598130 IMAGE Consortium	AA265440
5844 H	11 021CAMND	ESTs	598397 IMAGE Consortium	AI592123
5845 B	1 021JAMNE	ESTs, Weakly similar to E6-AP ubiquitin protein ligase [M.musculus]	597752 IMAGE Consortium	AI592044
5846 B	3 021JAMNE	ESTs	598648 IMAGE Consortium	AI592157
5847 B	5 021JAMNE	MAD homolog 1 (Drosophila)	775204 IMAGE Consortium	AA386935
5848 B	7 021JAMNE	ESTs, Weakly similar to Y182_HUMAN HYPOTHETICAL PROTEIN KIAA0182 [H.sapiens]	775388 IMAGE Consortium	AA386910
5849 B	9 021JAMNE	protein tyrosine phosphatase, non-receptor type 13	775515 IMAGE Consortium	AA388296
5850 B	11 021JAMNE	ESTs, Weakly similar to cDNA EST yk194a10.3 comes from this gene [C.elegans]	790151 IMAGE Consortium	AA388206
5851 D	1 021JAMNE	ESTs, Highly similar to AMIDOPHOSPHORIBOSYLTRANSFERASE PRECURSOR [Rattus norvegicus]	598497 IMAGE Consortium	AA161828
5852 D	3 021JAMNE	ESTs, Weakly similar to HRX [H.sapiens]	775073 IMAGE Consortium	AA387189
5853 D	5 021JAMNE	ESTs	775213 IMAGE Consortium	AA387204
5854 D	7 021JAMNE	ESTs, Highly similar to HYPOTHETICAL 13.5 KD PROTEIN C45G9.7 IN CHROMOSOME III [Caenorhabditis elegans]	775393 IMAGE Consortium	AA387336
5855 D	9 021JAMNE	ESTs, Weakly similar to CG7611 gene product [D.melanogaster]	775545 IMAGE Consortium	AA388315
5856 D	11 021JAMNE	block of proliferation 1	805066 IMAGE Consortium	AA397220
5857 F	1 021JAMNE	ESTs	597779 IMAGE Consortium	AA163807
5858 F	3 021JAMNE	cyclin-dependent kinase 7 (homolog of Xenopus MO15 cdk-activating kinase)	775078 IMAGE Consortium	AA387188
5859 F	5 021JAMNE	ESTs, Moderately similar to AF083217_1 WD repeat protein WDR3 [H.sapiens]	775221 IMAGE Consortium	AA387224
5860 F	7 021JAMNE	ESTs, Weakly similar to B0432.8 [C.elegans]	775399 IMAGE Consortium	AA387337

5861 F	9 021JAMNE	ESTs, Weakly similar to DHAR_RAT ALDEHYDE DEHYDROGENASE, CYTOSOLIC 2 [R.norvegicus]	775554 IMAGE Consortium	AA387284
5862 F	11 021JAMNE	ESTs	805119 IMAGE Consortium	AA397274
5863 H	1 021JAMNE	ESTs	597808 IMAGE Consortium	AI592051
5864 H	3 021JAMNE	ESTs, Moderately similar to KIAA1310 protein [H.sapiens]	775139 IMAGE Consortium	AA386870
5865 H	5 021JAMNE	ESTs, Weakly similar to CG4447 gene product [D.melanogaster]	775234 IMAGE Consortium	AA387207
5866 H	7 021JAMNE	ESTs	775403 IMAGE Consortium	AA387339
5867 H	9 021JAMNE	ESTs	774928 IMAGE Consortium	AI606754
5868 H	11 021JAMNE			
5869 B	1 021QAMNF	ESTs	805254 IMAGE Consortium	AI551641
5870 B	3 021QAMNF	growth factor receptor bound protein 2	804801 IMAGE Consortium	AA472836
5871 B	5 021QAMNF	ESTs	805803 IMAGE Consortium	AA437473
5872 B	7 021QAMNF	thymoma viral proto-oncogene 2	819872 IMAGE Consortium	AA437947
5873 B	9 021QAMNF	ESTs, Weakly similar to AF119384_1 putative CAMP protein [M.musculus]	820039 IMAGE Consortium	AI551582
5874 B	11 021QAMNF	C-terminal binding protein 1	805714 IMAGE Consortium	AA437639
5875 D	1 021QAMNF	ESTs	804548 IMAGE Consortium	AA474488
5876 D	3 021QAMNF	ESTs	804978 IMAGE Consortium	AI551693
5877 D	5 021QAMNF	GATA-binding protein 2	805864 IMAGE Consortium	AA399858
5878 D	7 021QAMNF			
5879 D	9 021QAMNF	ESTs, Highly similar to INORGANIC PYROPHOSPHATASE [Bos taurus]	820086 IMAGE Consortium	AI551585
5880 D	11 021QAMNF	Tcra enhancer-binding factor interacting protein 1	805740 IMAGE Consortium	AA437646
5881 F	1 021QAMNF	RNA polymerase 1-1 (40 kDa subunit)	804639 IMAGE Consortium	AA474735
5882 F	3 021QAMNF	ESTs	789847 IMAGE Consortium	AA387979
5883 F	5 021QAMNF	SMC (segregation of mitotic chromosomes 1)-like 1 (yeast)	805887 IMAGE Consortium	AA437510
5884 F	7 021QAMNF	thymus expressed acidic protein	819912 IMAGE Consortium	AA437755
5885 F	9 021QAMNF	ESTs	805172 IMAGE Consortium	AI551706
5886 F	11 021QAMNF	ESTs	805771 IMAGE Consortium	AA437745
5887 H	1 021QAMNF	ESTs, Moderately similar to OVCA1	805400 IMAGE Consortium	AA473397
5888 H	3 021QAMNF	PHD finger protein 2	789877 IMAGE Consortium	AA387885
5889 H	5 021QAMNF	FK506 binding protein 6 (65 kDa)	819722 IMAGE Consortium	AA437512
5890 H	7 021QAMNF	aplysia ras-related homolog B (RhoB)	819960 IMAGE Consortium	AA437783
5891 H	9 021QAMNF	adenylyl cyclase-associated CAP protein homolog 1 (S. cerevisiae, S. Pombe)	805411 IMAGE Consortium	AA473213
5892 H	11 021QAMNF	ESTs, Highly similar to H<BETA>58 PROTEIN [Mus musculus]	834128 IMAGE Consortium	AA412893
5893 B	1 021XAMNG	ESTs, Moderately similar to RL32_HUMAN 60S RIBOSOMAL PROTEIN L32 [M.musculus]	833561 IMAGE Consortium	AA470267
5894 B	3 021XAMNG	ESTs	833900 IMAGE Consortium	AI596508
5895 B	5 021XAMNG	cytokine inducible SH2-containing protein 1	819853 IMAGE Consortium	AA437942

5896 B	7 021XAMNG	ESTs	819999 IMAGE Consortium	AA437812
5897 B	9 021XAMNG	nuclear factor of kappa light chain gene enhancer in B-cells inhibitor, alpha	894295 IMAGE Consortium	AA517353
5898 B	11 021XAMNG	DNA segment, Chr 12, Wayne State University 118, expressed	894417 IMAGE Consortium	AA517501
5899 D	1 021XAMNG	zinc finger protein 162	833695 IMAGE Consortium	AI596499
5900 D	3 021XAMNG	fibulin 5	819734 IMAGE Consortium	AA437518
5901 D	5 021XAMNG	myogenic differentiation 1	819856 IMAGE Consortium	AA437694
5902 D	7 021XAMNG	ESTs	820021 IMAGE Consortium	AI551576
5903 D	9 021XAMNG	ESTs	894309 IMAGE Consortium	AA517359
5904 D	11 021XAMNG	ESTs	894435 IMAGE Consortium	AA517518
5905 F	1 021XAMNG	ESTs	833722 IMAGE Consortium	AI596500
5906 F	3 021XAMNG	ESTs	819742 IMAGE Consortium	AA437662
5907 F	5 021XAMNG	ESTs, Weakly similar to unnamed protein product [H.sapiens]	819909 IMAGE Consortium	AA437705
5908 F	7 021XAMNG	ESTs	820031 IMAGE Consortium	AI551577
5909 F	9 021XAMNG	zinc finger protein 90	894331 IMAGE Consortium	AA517408
5910 F	11 021XAMNG	DNA primase, p58 subunit	894463 IMAGE Consortium	AA517535
5911 H	1 021XAMNG	HIRA-interacting protein 5	834512 IMAGE Consortium	AA466741
5912 H	3 021XAMNG	ESTs	819764 IMAGE Consortium	AI595387
5913 H	5 021XAMNG	ESTs	819927 IMAGE Consortium	AI467499
5914 H	7 021XAMNG	ESTs	820071 IMAGE Consortium	AA444220
5915 H	9 021XAMNG	MAD2 (mitotic arrest deficient, homolog)-like 1 (yeast)	894340 IMAGE Consortium	AA517411
5916 H	11 021XAMNG	B-cell receptor-associated protein 29	894470 IMAGE Consortium	AA517532
5917 B	1 0214AMNH	upstream transcription factor 2	932717 IMAGE Consortium	AA538409
5918 B	3 0214AMNH	ESTs	835205 IMAGE Consortium	AI551715
5919 B	5 0214AMNH	X-ray repair complementing defective repair in Chinese hamster cells 1	893835 IMAGE Consortium	AA517184
5920 B	7 0214AMNH	ESTs, Highly similar to AF132970_1 CGI-36 protein [H.sapiens]	934097 IMAGE Consortium	AA544586
5921 B	9 0214AMNH	thymidine kinase 1	933772 IMAGE Consortium	AA545691
5922 B	11 0214AMNH	xeroderma pigmentosum, complementation group C	934267 IMAGE Consortium	AA543950
5923 D	1 0214AMNH	adaptor-related protein complex AP-3, sigma 1 subunit	932829 IMAGE Consortium	AA538552
5924 D	3 0214AMNH	ESTs	894176 IMAGE Consortium	AA517038
5925 D	5 0214AMNH	ESTs	893978 IMAGE Consortium	AA516644
5926 D	7 0214AMNH	DEAD/H (Asp-Glu-Ala-Asp/His) box polypeptide 15 (RNA helicase A)	934141 IMAGE Consortium	AA543848
5927 D	9 0214AMNH	Mus musculus X chromosome: L1cam locus	933868 IMAGE Consortium	AA544392
5928 D	11 0214AMNH	ESTs	934285 IMAGE Consortium	AA543967
5929 F	1 0214AMNH	peroxiredoxin 1	932882 IMAGE Consortium	AA541869
5930 F	3 0214AMNH	ESTs	892736 IMAGE Consortium	AI614566
5931 F	5 0214AMNH	ESTs, Weakly similar to AF151067_1 HSPC233 [H.sapiens]	836803 IMAGE Consortium	AA426848

5932 F	7 0214AMNH	Eph receptor A7	933489 IMAGE Consortium	AA542347
5933 F	9 0214AMNH	thioredoxin	933897 IMAGE Consortium	AA544415
5934 F	11 0214AMNH	ESTs, Highly similar to ER transmembrane protein [R.norvegicus]	934399 IMAGE Consortium	AA545032
5935 H	1 0214AMNH	Public domain EST	932997 IMAGE Consortium	
5936 H	3 0214AMNH	ESTs, Weakly similar to C1TC_RAT C-1-TETRAHYDROFOLATE SYNTHASE, CYTOPLASMIC [R.norvegicus]	835341 IMAGE Consortium	AI551722
5937 H	5 0214AMNH	mitogen activated protein kinase kinase kinase 4	933010 IMAGE Consortium	AA538061
5938 H	7 0214AMNH	ESTs, Highly similar to KIAA1194 protein [H.sapiens]	933526 IMAGE Consortium	AA542363
5939 H	9 0214AMNH	dihydroorotate dehydrogenase	933929 IMAGE Consortium	AA544433
5940 H	11 0214AMNH	ESTs, Weakly similar to CG7053 gene product [D.melanogaster]	934415 IMAGE Consortium	AA545037
5941 B	1 021BAMNI	glutathione synthetase	934599 IMAGE Consortium	AA536773
5942 B	3 021BAMNI	TEA domain family member 2	934812 IMAGE Consortium	AA617316
5943 B	5 021BAMNI	ESTs	598949 IMAGE Consortium	AI592213
5944 B	7 021BAMNI	ESTs	616791 IMAGE Consortium	AI640067
5945 B	9 021BAMNI	ESTs	599126 IMAGE Consortium	AI592237
5946 B	11 021BAMNI	ESTs	947483 IMAGE Consortium	AA588967
5947 D	1 021BAMNI	ESTs, Weakly similar to 50S RIBOSOMAL PROTEIN L13 [Staphylococcus carnosus]	934641 IMAGE Consortium	AA536840
5948 D	3 021BAMNI	ESTs, Highly similar to ANTI-SILENCING PROTEIN 1 [Saccharomyces cerevisiae]	934829 IMAGE Consortium	AA617354
5949 D	5 021BAMNI	ESTs, Weakly similar to AF151066_1 HSPC232 [H.sapiens]	617015 IMAGE Consortium	AI640084
5950 D	7 021BAMNI	ESTs	617518 IMAGE Consortium	AI639733
5951 D	9 021BAMNI	ESTs	599244 IMAGE Consortium	AI592253
5952 D	11 021BAMNI	ESTs	947509 IMAGE Consortium	AA589092
5953 F	1 021BAMNI	stimulated by retinoic acid 14	934650 IMAGE Consortium	AA536836
5954 F	3 021BAMNI	ESTs	934836 IMAGE Consortium	AA617350
5955 F	5 021BAMNI	ESTs	617207 IMAGE Consortium	AI640122
5956 F	7 021BAMNI	ESTs	617558 IMAGE Consortium	AA174870
5957 F	9 021BAMNI	ESTs	946371 IMAGE Consortium	AA591929
5958 F	11 021BAMNI	ubiquitin conjugating enzyme E3A	962904 IMAGE Consortium	AA561616
5959 H	1 021BAMNI	meiotic recombination 11 homolog A (S. cerevisiae)	934673 IMAGE Consortium	AA536853
5960 H	3 021BAMNI	ESTs, Weakly similar to AF145632_1 BcDNA.GH06032 [D.melanogaster]	934846 IMAGE Consortium	AA617435
5961 H	5 021BAMNI	ESTs	617277 IMAGE Consortium	AI639704
5962 H	7 021BAMNI	ESTs	617574 IMAGE Consortium	AA174799
5963 H	9 021BAMNI	X-linked lymphocyte-regulated 4	947247 IMAGE Consortium	AA592023
5964 H	11 021BAMNI	ESTs, Weakly similar to T08675 hypothetical protein DKFZp564F0522.1 - human [H.sapiens]	962906 IMAGE Consortium	AA561617
5965 B	1 021IAMNJ	ESTs	946276 IMAGE Consortium	AA591758

5966 B	3 021IAMNJ	annexin A7	935159 IMAGE Consortium	AA616992
5967 B	5 021IAMNJ	ESTs	945048 IMAGE Consortium	AA537308
5968 B	7 021IAMNJ	ESTs	945526 IMAGE Consortium	AA545932
5969 B	9 021IAMNJ	ESTs	962946 IMAGE Consortium	AA619655
5970 B	11 021IAMNJ	ESTs, Highly similar to DNA-DIRECTED RNA POLYMERASE II 14.4 KD POLYPEPTIDE [Homo sapiens; Cricetulus griseus]	976809 IMAGE Consortium	AA563333
5971 D	1 021IAMNJ	ESTs, Moderately similar to tumor necrosis factor receptor type 1 associated protein [H.sapiens]	945604 IMAGE Consortium	AA546092
5972 D	3 021IAMNJ	ras-GTPase-activating protein (GAP<120>) SH3-domain-binding protein 2	935173 IMAGE Consortium	AA617613
5973 D	5 021IAMNJ			
5974 D	7 021IAMNJ	MORF-related gene X	945534 IMAGE Consortium	AA545943
5975 D	9 021IAMNJ	ESTs	977016 IMAGE Consortium	AI645409
5976 D	11 021IAMNJ	DNA segment, Chr 19, Wayne State University 55, expressed	976844 IMAGE Consortium	AA563439
5977 F	1 021IAMNJ	ESTs	945643 IMAGE Consortium	AA546122
5978 F	3 021IAMNJ	calcium binding protein A11 (calgizzarin)	935181 IMAGE Consortium	AA616998
5979 F	5 021IAMNJ	ESTs	934952 IMAGE Consortium	AA617469
5980 F	7 021IAMNJ	ESTs	945584 IMAGE Consortium	AA546079
5981 F	9 021IAMNJ	ESTs	977139 IMAGE Consortium	AA563239
5982 F	11 021IAMNJ	ESTs	977597 IMAGE Consortium	AI645441
5983 H	1 021IAMNJ	melanoma antigen, family A, 2	945651 IMAGE Consortium	AI585424
5984 H	3 021IAMNJ	ribosomal protein, mitochondrial, L12	935203 IMAGE Consortium	AA617002
5985 H	5 021IAMNJ	ESTs	945134 IMAGE Consortium	AA537410
5986 H	7 021IAMNJ	membrane protein, palmitoylated (55 kDa)	962724 IMAGE Consortium	AA562453
5987 H	9 021IAMNJ	Mus musculus prostaglandin transporter PGT mRNA, complete cds	977205 IMAGE Consortium	AA563370
5988 H	11 021IAMNJ	glutathione peroxidase 1	976879 IMAGE Consortium	AA562552
5989 B	1 021PAMNK	cofilin 1, non-muscle	976930 IMAGE Consortium	AA562691
5990 B	3 021PAMNK	ESTs	989119 IMAGE Consortium	AA571342
5991 B	5 021PAMNK	ectonucleoside triphosphate diphosphohydrolase 6	988648 IMAGE Consortium	AA638277
5992 B	7 021PAMNK	ESTs	988950 IMAGE Consortium	AA591081
5993 B	9 021PAMNK	expressed in non-metastatic cells 6, protein (nucleoside diphosphate kinase)	991006 IMAGE Consortium	AA590168
5994 B	11 021PAMNK	ESTs	992309 IMAGE Consortium	AA589343
5995 D	1 021PAMNK			
5996 D	3 021PAMNK	guanine nucleotide binding protein, beta-2, related sequence 1	988421 IMAGE Consortium	AA571755
5997 D	5 021PAMNK	ESTs	988663 IMAGE Consortium	AA638300
5998 D	7 021PAMNK	DNA segment, Chr 8, ERATO Doi 233, expressed	978036 IMAGE Consortium	AA563376
5999 D	9 021PAMNK	protein phosphatase 2a, catalytic subunit, beta isoform	991138 IMAGE Consortium	AI604561
6000 D	11 021PAMNK	ESTs	991639 IMAGE Consortium	AI604605

6001 F	1	021PAMNK	EST AI115388	976937	IMAGE Consortium	AA562703
6002 F	3	021PAMNK	ESTs, Moderately similar to 2012303A SP-H antigen [H.sapiens]	989185	IMAGE Consortium	AA571790
6003 F	5	021PAMNK	fascin homolog 1 (actin bundling protein, Strongylocentrotus purpuratus)	988708	IMAGE Consortium	AA638321
6004 F	7	021PAMNK	ESTs, Highly similar to sodium-dependent multi-vitamin transporter [R.norvegicus]	978067	IMAGE Consortium	AA563404
6005 F	9	021PAMNK	DNA segment, Chr 5, Wayne State University 152, expressed	991383	IMAGE Consortium	AI604578
6006 F	11	021PAMNK	ESTs	991657	IMAGE Consortium	AI604607
6007 H	1	021PAMNK	RNA binding motif protein, X chromosome retogene	976965	IMAGE Consortium	AA562722
6008 H	3	021PAMNK	ATP synthase, H <sup>+</sup> transporting, mitochondrial F0 complex subunit F	988520	IMAGE Consortium	AA638190
6009 H	5	021PAMNK	ESTs, Highly similar to JC6523 26s proteasom p55 protein - human [H.sapiens]	988745	IMAGE Consortium	AI586347
6010 H	7	021PAMNK	galactokinase	978228	IMAGE Consortium	AA620113
6011 H	9	021PAMNK	ESTs, Moderately similar to Dyrk3 protein [H.sapiens]	992145	IMAGE Consortium	AA589241
6012 H	11	021PAMNK	transcription factor 1	991667	IMAGE Consortium	AA574603
6013 B	1	021WAMNL				
6014 B	3	021WAMNL	ESTs	990046	IMAGE Consortium	AI604488
6015 B	5	021WAMNL	ESTs	990960	IMAGE Consortium	AA590024
6016 B	7	021WAMNL	ESTs, Moderately similar to 60S RIBOSOMAL PROTEIN L39 [Rattus norvegicus]	1004267	IMAGE Consortium	AA600661
6017 B	9	021WAMNL	NPC derived proline rich protein 1	1004424	IMAGE Consortium	AA791966
6018 B	11	021WAMNL	ESTs	1004720	IMAGE Consortium	AI503045
6019 D	1	021WAMNL	ESTs, Highly similar to snurportin1 [H.sapiens]	989580	IMAGE Consortium	AA571557
6020 D	3	021WAMNL	DNA segment, Chr 3, ERATO Doi 801, expressed	990136	IMAGE Consortium	AI604492
6021 D	5	021WAMNL	claudin 7	1004063	IMAGE Consortium	AA600553
6022 D	7	021WAMNL	ESTs	1004278	IMAGE Consortium	AA600656
6023 D	9	021WAMNL	DNA segment, Chr 2, Wayne State University 101, expressed	1004486	IMAGE Consortium	AA791990
6024 D	11	021WAMNL	ESTs	1004744	IMAGE Consortium	AI594233
6025 F	1	021WAMNL	ESTs, Highly similar to I38863 E1B 19K/Bcl-2-interacting protein Nip1 - human [H.sapiens]	989694	IMAGE Consortium	AA571665
6026 F	3	021WAMNL	fibrillarlin	990178	IMAGE Consortium	AI604500
6027 F	5	021WAMNL	ESTs, Weakly similar to B3AT_MOUSE BAND 3 ANION EXCHANGE PROTEIN [M.musculus]	1004086	IMAGE Consortium	AA600562
6028 F	7	021WAMNL	thioredoxin nuclear gene encoding mitochondrial protein	1004301	IMAGE Consortium	AA600673
6029 F	9	021WAMNL	ESTs, Highly similar to T12523 hypothetical protein DKFZp434K063.1 - human [H.sapiens]	1004510	IMAGE Consortium	AA792001
6030 F	11	021WAMNL	myeloid-associated differentiation marker	1004798	IMAGE Consortium	AA606711
6031 H	1	021WAMNL	glutamyl aminopeptidase	989776	IMAGE Consortium	AA793433
6032 H	3	021WAMNL	ESTs, Weakly similar to AAC82480 BAT2 [M.musculus]	990214	IMAGE Consortium	AA589966
6033 H	5	021WAMNL	DEAD/H (Asp-Glu-Ala-Asp/His) box polypeptide 13 (RNA	1004105	IMAGE Consortium	AA600577

		helicase A)		
6034 H	7	021WAMNL		
6035 H	9	021WAMNL	ESTs, Highly similar to PRE-MRNA SPLICING FACTOR SF2, P33 SUBUNIT [Homo sapiens]	1004627 IMAGE Consortium AI606396
6036 H	11	021WAMNL	ESTs	992579 IMAGE Consortium AA607939
6037 B	1	0213AMNM	neogenin	1005030 IMAGE Consortium AA608182
6038 B	3	0213AMNM	ESTs, Weakly similar to AF161429_1 HSPC311 [H.sapiens]	1005115 IMAGE Consortium AA606595
6039 B	5	0213AMNM	proteasome (prosome, macropain) subunit, beta type 3	1004778 IMAGE Consortium AA606697
6040 B	7	0213AMNM	ATP-binding cassette, sub-family B (MDR/TAP), member 8	1004884 IMAGE Consortium AA606983
6041 B	9	0213AMNM	ESTs, Weakly similar to nuclear respiratory factor-1 [M.musculus]	1004955 IMAGE Consortium AI594252
6042 B	11	0213AMNM	DNA segment, Chr 5, ERATO Doi 591, expressed	1006235 IMAGE Consortium AA624171
6043 D	1	0213AMNM	expressed in non-metastatic cells 2, protein (NM23B) (nucleoside diphosphate kinase)	1005045 IMAGE Consortium AA608196
6044 D	3	0213AMNM	ESTs, Highly similar to dithiolethione-inducible gene-1 [R.norvegicus]	1005129 IMAGE Consortium AA606610
6045 D	5	0213AMNM	dynactin 1	1004802 IMAGE Consortium AA606713
6046 D	7	0213AMNM	ELKL motif kinase	1004888 IMAGE Consortium AA606985
6047 D	9	0213AMNM	ESTs, Weakly similar to scaffold attachment factor B [R.norvegicus]	1004964 IMAGE Consortium AA607193
6048 D	11	0213AMNM	ESTs	1005594 IMAGE Consortium AI594275
6049 F	1	0213AMNM	activating transcription factor 4	1005055 IMAGE Consortium AA608326
6050 F	3	0213AMNM	ESTs, Highly similar to hD54+ins2 isoform [H.sapiens]	1005132 IMAGE Consortium AA606603
6051 F	5	0213AMNM	ESTs	1004816 IMAGE Consortium AI594239
6052 F	7	0213AMNM	ESTs	1004901 IMAGE Consortium AI594247
6053 F	9	0213AMNM	solute carrier family 15 (H+/peptide transporter), member 2	1004969 IMAGE Consortium AA607205
6054 F	11	0213AMNM	Mus musculus clone BAC126c8 Rsp29-like protein (Rsp29) and Als splice variant 2 (Als) genes, partial cds; Als splice variant 1 (Als), TCE2 (Tce2), NDK3-like protein (Ndk3), and TCE4 (Tce4) genes, complete cds; and TCE5 (Tce5) gene	1005613 IMAGE Consortium AA606928
6055 H	1	0213AMNM	DNA segment, Chr 9, Wayne State University 149, expressed	1005059 IMAGE Consortium AA608328
6056 H	3	0213AMNM	importin beta	1005136 IMAGE Consortium AA606604
6057 H	5	0213AMNM	plasmacytoma variant translocation 1	1004823 IMAGE Consortium AA606860
6058 H	7	0213AMNM	pericentrin	1004915 IMAGE Consortium AA607089
6059 H	9	0213AMNM	ESTs, Highly similar to TRAM PROTEIN [Canis familiaris]	1004980 IMAGE Consortium AA607209
6060 H	11	0213AMNM	peroxisomal biogenesis factor 14	1005622 IMAGE Consortium AA606923
6061 B	1	021AAMNN	cytochrome c oxidase, subunit IV	1005717 IMAGE Consortium AA607141
6062 B	3	021AAMNN	Public domain EST	1005838 IMAGE Consortium
6063 B	5	021AAMNN	neural precursor cell expressed, developmentally down-	1005949 IMAGE Consortium AA606845

		regulated gene 1		
6064 B	7 021AAMNN	serine/threonine kinase 16	1005393 IMAGE Consortium	AA608208
6065 B	9 021AAMNN	ESTs	1005606 IMAGE Consortium	AI594276
6066 B	11 021AAMNN	mitogen activated protein kinase kinase kinase 8	620757 IMAGE Consortium	AA174638
6067 D	1 021AAMNN	MYB binding protein (P160) 1a	1005736 IMAGE Consortium	AA607245
6068 D	3 021AAMNN	eIF3 p66	1005852 IMAGE Consortium	AA606687
6069 D	5 021AAMNN	E74-like factor 5	1005961 IMAGE Consortium	AA607032
6070 D	7 021AAMNN	RAB2, member RAS oncogene family	1005404 IMAGE Consortium	AA608203
6071 D	9 021AAMNN	matrin 3	1005648 IMAGE Consortium	AA607011
6072 D	11 021AAMNN	ESTs, Weakly similar to A26882 pIL2 hypothetical protein - rat [R.norvegicus]	620055 IMAGE Consortium	AI604172
6073 F	1 021AAMNN	DNA methyltransferase 3A	1005791 IMAGE Consortium	AA606936
6074 F	3 021AAMNN	ribosomal protein S3a	1005855 IMAGE Consortium	AA606691
6075 F	5 021AAMNN	fatty acid binding protein 3, muscle and heart	1005968 IMAGE Consortium	AA607030
6076 F	7 021AAMNN	ESTs, Weakly similar to C43H8.1 [C.elegans]	1005407 IMAGE Consortium	AI594266
6077 F	9 021AAMNN	ESTs	1005651 IMAGE Consortium	AI594280
6078 F	11 021AAMNN	ESTs	620100 IMAGE Consortium	AA176034
6079 H	1 021AAMNN	branched chain aminotransferase 1, cytosolic	1005806 IMAGE Consortium	AA606666
6080 H	3 021AAMNN	cytochrome c oxidase, subunit VIIIa	1005861 IMAGE Consortium	AA606950
6081 H	5 021AAMNN	ESTs, Moderately similar to unnamed protein product [H.sapiens]	1005972 IMAGE Consortium	AI594296
6082 H	7 021AAMNN	ESTs	1005419 IMAGE Consortium	AA608224
6083 H	9 021AAMNN	ESTs	1005744 IMAGE Consortium	AA607255
6084 H	11 021AAMNN	ESTs	620175 IMAGE Consortium	AI604192
6085 B	1 021HAMNO	ESTs	621109 IMAGE Consortium	AI592272
6086 B	3 021HAMNO	ESTs	620768 IMAGE Consortium	AI604265
6087 B	5 021HAMNO	ESTs	1037717 IMAGE Consortium	AA638851
6088 B	7 021HAMNO	ESTs, Highly similar to KIAA0121 protein [H.sapiens]	1007095 IMAGE Consortium	AA637080
6089 B	9 021HAMNO	ESTs	1006598 IMAGE Consortium	AA624363
6090 B	11 021HAMNO	ESTs, Highly similar to ANKYRIN, BRAIN VARIANT 2 [Homo sapiens]	1038028 IMAGE Consortium	AA624418
6091 D	1 021HAMNO	ESTs	621129 IMAGE Consortium	AI592278
6092 D	3 021HAMNO	ESTs	621513 IMAGE Consortium	AI592335
6093 D	5 021HAMNO	ESTs, Weakly similar to /prediction	1037763 IMAGE Consortium	AA638885
6094 D	7 021HAMNO	ESTs	1006441 IMAGE Consortium	AA624262
6095 D	9 021HAMNO	ESTs, Weakly similar to veli 2 [M.musculus]	1005998 IMAGE Consortium	AA607044
6096 D	11 021HAMNO	ESTs	1038033 IMAGE Consortium	AA624428
6097 F	1 021HAMNO	ESTs	621152 IMAGE Consortium	AI592284
6098 F	3 021HAMNO	ESTs	620812 IMAGE Consortium	AI604270
6099 F	5 021HAMNO	ESTs, Weakly similar to KIAA0402 [H.sapiens]	1022071 IMAGE Consortium	AI048432
6100 F	7 021HAMNO	ESTs	1021754 IMAGE Consortium	AI047988
6101 F	9 021HAMNO	neuroblastoma myc-related oncogene 1	1022857 IMAGE Consortium	AA623387
6102 F	11 021HAMNO	cornichon homolog (Drosophila)	1038036 IMAGE Consortium	AA624421



6103 H	1 021HAMNO	ESTs	620451 IMAGE Consortium	AA178165
6104 H	3 021HAMNO	ESTs	620865 IMAGE Consortium	AI604277
6105 H	5 021HAMNO	cell division cycle 25 homolog C ( <i>S. cerevisiae</i> )	1022189 IMAGE Consortium	AA623344
6106 H	7 021HAMNO	ESTs, Weakly similar to S14234 hypothetical protein - mouse [M.musculus]	1006476 IMAGE Consortium	AI594323
6107 H	9 021HAMNO	ESTs	1006855 IMAGE Consortium	AI594128
6108 H	11 021HAMNO	ESTs, Highly similar to KIAA0017 protein [H.sapiens]	1038080 IMAGE Consortium	AA624518
6109 B	1 021OAMNP	fibroblast growth factor regulated protein	1038144 IMAGE Consortium	AA624690
6110 B	3 021OAMNP			
6111 B	5 021OAMNP	proliferating cell nuclear antigen	1037829 IMAGE Consortium	AA636924
6112 B	7 021OAMNP	ESTs	1038218 IMAGE Consortium	AA671098
6113 B	9 021OAMNP	cystatin B	1038303 IMAGE Consortium	AA671231
6114 B	11 021OAMNP	tropomyosin 5	1038416 IMAGE Consortium	AA624485
6115 D	1 021OAMNP	ethanol induced 1	1038180 IMAGE Consortium	AA624810
6116 D	3 021OAMNP	Williams-Beuren syndrome chromosome region 1 homolog (human)	1037710 IMAGE Consortium	AA638805
6117 D	5 021OAMNP	caudal type homeo box 1	1037845 IMAGE Consortium	AA638924
6118 D	7 021OAMNP	heat shock protein, 110 kDa	1038241 IMAGE Consortium	AA671121
6119 D	9 021OAMNP	erythroblast macrophage protein	1038309 IMAGE Consortium	AA671232
6120 D	11 021OAMNP	silica-induced gene 41	1038420 IMAGE Consortium	AA624487
6121 F	1 021OAMNP	ESTs, Highly similar to HYPOTHETICAL 16.5 KD PROTEIN IN PAS8-EGT2 INTERGENIC REGION [Saccharomyces cerevisiae]	1038208 IMAGE Consortium	AA624825
6122 F	3 021OAMNP	ESTs	1037729 IMAGE Consortium	AI642260
6123 F	5 021OAMNP	ESTs, Moderately similar to KIAA1291 protein [H.sapiens]	1037850 IMAGE Consortium	AA638918
6124 F	7 021OAMNP	RAB7, member RAS oncogene family	1038263 IMAGE Consortium	AA671131
6125 F	9 021OAMNP			
6126 F	11 021OAMNP	prothymosin alpha	1038424 IMAGE Consortium	AA624488
6127 H	1 021OAMNP	ESTs	1037577 IMAGE Consortium	AA623587
6128 H	3 021OAMNP	Mus musculus eIF4E-like protein 4E-LP mRNA, complete cds	1037767 IMAGE Consortium	AA638887
6129 H	5 021OAMNP	poly(rC) binding protein 2	1037871 IMAGE Consortium	AA638938
6130 H	7 021OAMNP	ESTs	1038266 IMAGE Consortium	AA671194
6131 H	9 021OAMNP	Mus musculus dUTPase mRNA, complete cds	1038333 IMAGE Consortium	AA671302
6132 H	11 021OAMNP	RNA polymerase II transcriptional coactivator	1038443 IMAGE Consortium	AA624511
6133 B	1 021VAMNQ	ESTs, Highly similar to bK223H9.2 [H.sapiens]	1038485 IMAGE Consortium	AA624607
6134 B	3 021VAMNQ	NIMA (never in mitosis gene a)-related expressed kinase 6	1037879 IMAGE Consortium	AA638941
6135 B	5 021VAMNQ	secretory carrier membrane protein 3	1109659 IMAGE Consortium	AA822797
6136 B	7 021VAMNQ	lectin, galactose binding, soluble 6	1110364 IMAGE Consortium	AA674809
6137 B	9 021VAMNQ	ESTs	1110407 IMAGE Consortium	AA674839
6138 B	11 021VAMNQ	DNA segment, Chr 8, ERATO Doi 69, expressed	2102253 IMAGE Consortium	AW209351
6139 D	1 021VAMNQ	ribosomal protein L35a	1038490 IMAGE Consortium	AA624600

6140 D	3	021VAMNQ	ribosomal protein L23	1082018	IMAGE Consortium	AA674018
6141 D	5	021VAMNQ				
6142 D	7	021VAMNQ	ADP-ribosylarginine hydrolase	1110067	IMAGE Consortium	AA795220
6143 D	9	021VAMNQ	ESTs	1110423	IMAGE Consortium	AA675174
6144 D	11	021VAMNQ	PCTAIRE-motif protein kinase 1	2101725	IMAGE Consortium	AW210038
6145 F	1	021VAMNQ	ESTs, Highly similar to sorting nexin 4 [H.sapiens]	1038497	IMAGE Consortium	AA624612
6146 F	3	021VAMNQ	ESTs	1082161	IMAGE Consortium	AA815841
6147 F	5	021VAMNQ	E74-like factor 3	1109729	IMAGE Consortium	AA822889
6148 F	7	021VAMNQ	ESTs	1110113	IMAGE Consortium	AA795508
6149 F	9	021VAMNQ	ESTs, Weakly similar to similar to Human homolog of Drosophila female sterile homeotic mRNA [H.sapiens]	1110439	IMAGE Consortium	AA675191
6150 F	11	021VAMNQ	ESTs	2101903	IMAGE Consortium	AW210206
6151 H	1	021VAMNQ	heterogeneous nuclear ribonucleoprotein L	1038500	IMAGE Consortium	AA624604
6152 H	3	021VAMNQ	lysophospholipase 1	1109309	IMAGE Consortium	AA666977
6153 H	5	021VAMNQ	ESTs, Highly similar to hypothetical protein [H.sapiens]	1109734	IMAGE Consortium	AA822882
6154 H	7	021VAMNQ	ESTs	1110225	IMAGE Consortium	AA795576
6155 H	9	021VAMNQ	ESTs	1110623	IMAGE Consortium	AA684512
6156 H	11	021VAMNQ	ESTs	2101947	IMAGE Consortium	AW210242
6157 B	1	0212AMNR	epidermal growth factor-containing fibulin-like extracellular matrix protein 2	2102061	IMAGE Consortium	AW210333
6158 B	3	0212AMNR	ESTs, Weakly similar to DDX9_MOUSE ATP-DEPENDENT RNA HELICASE A [M.musculus]	2101537	IMAGE Consortium	AW209882
6159 B	5	0212AMNR	SH3-domain binding protein 1	2101142	IMAGE Consortium	AW209528
6160 B	7	0212AMNR	ESTs, Moderately similar to The ha1438 gene product is related to a C728 protein encoded in S.cerevisiae chromosome II. [H.sapiens] {IMAGE:804482}	804482	IMAGE Consortium	AA474342
6161 B	9	0212AMNR	ESTs {IMAGE:934337}	934337	IMAGE Consortium	AA544844
6162 B	11	0212AMNR	ESTs, Weakly similar to HIRA protein [M.musculus] {IMAGE:761060}	761060	IMAGE Consortium	AA387585
6163 D	1	0212AMNR	ESTs, Highly similar to QPCT_HUMAN GLUTAMINYLPOLYMERASE PRECURSOR [H.sapiens]	2102081	IMAGE Consortium	AW210354
6164 D	3	0212AMNR	ESTs, Weakly similar to GLUCOSYLTRANSFERASE ALG8 [Saccharomyces cerevisiae]	2101551	IMAGE Consortium	AW209889
6165 D	5	0212AMNR	ubiquitin-conjugating enzyme E2H	2101175	IMAGE Consortium	AW209562
6166 D	7	0212AMNR	ESTs, Moderately similar to p53BP2 [M.musculus] {IMAGE:834947}	834947	IMAGE Consortium	AA467287
6167 D	9	0212AMNR	Peroxisomal membrane protein, 70 kDa {IMAGE:805835}	805835	IMAGE Consortium	AA437485
6168 D	11	0212AMNR	ESTs {IMAGE:834225}	834225	IMAGE Consortium	AA413194
6169 F	1	0212AMNR	ESTs, Highly similar to METHIONYL-TRNA SYNTHETASE, CYTOPLASMIC [Saccharomyces cerevisiae]	2102126	IMAGE Consortium	AW210384
6170 F	3	0212AMNR	Abelson murine leukemia oncogene	2101588	IMAGE Consortium	AW209918
6171 F	5	0212AMNR	ESTs	2101189	IMAGE Consortium	AW209580

6172 F	7 0212AMNR	ESTs, Highly similar to THYMIDYLATE KINASE [Homo sapiens] {IMAGE:760921}	760921 IMAGE Consortium	AA386718
6173 F	9 0212AMNR	IMAGE EST {IMAGE:894302}	894302 IMAGE Consortium	AA517346
6174 F	11 0212AMNR	IMAGE EST {IMAGE:834617}	834617 IMAGE Consortium	AA467546
6175 H	1 0212AMNR	ankyrin 3, epithelial	2102138 IMAGE Consortium	AW210399
6176 H	3 0212AMNR	ESTs, Moderately similar to Y039_HUMAN HYPOTHETICAL PROTEIN KIAA0039 [H.sapiens]	2101664 IMAGE Consortium	AW209980
6177 H	5 0212AMNR			
6178 H	7 0212AMNR	IMAGE EST {IMAGE:834712}	834712 IMAGE Consortium	AA467053
6179 H	9 0212AMNR	ESTs, Highly similar to TRANSCRIPTION INITIATION FACTOR IIE, BETA SUBUNIT [H.sapiens] {IMAGE:833858}	833858 IMAGE Consortium	AA412879
6180 H	11 0212AMNR	Sine oculis-related homeobox 3, (Drosophila) {IMAGE:761326}	761326 IMAGE Consortium	AA388136
6181 B	1 0219AMNS	ESTs {IMAGE:805525}	805525 IMAGE Consortium	AA397161
6182 B	3 0219AMNS	ESTs {IMAGE:535718}	535718 IMAGE Consortium	AA106962
6183 B	5 0219AMNS	ESTs, Highly similar to ISOCITRATE DEHYDROGENASE [Bos taurus] {IMAGE:775189}	775189 IMAGE Consortium	AA387198
6184 B	7 0219AMNS	ESTs	619224 IMAGE Consortium	AI592555
6185 B	9 0219AMNS	ESTs	619381 IMAGE Consortium	AA175578
6186 B	11 0219AMNS	ESTs	618954 IMAGE Consortium	AI592510
6187 D	1 0219AMNS	IMAGE EST {IMAGE:805526}	805526 IMAGE Consortium	AA397153
6188 D	3 0219AMNS	ESTs {IMAGE:534015}	534015 IMAGE Consortium	AA072800
6189 D	5 0219AMNS	coagulation factor II (thrombin) receptor-like 1	619808 IMAGE Consortium	AA177852
6190 D	7 0219AMNS	fibronectin 1	619240 IMAGE Consortium	AA170781
6191 D	9 0219AMNS	ESTs	619391 IMAGE Consortium	AA172521
6192 D	11 0219AMNS	ESTs	618961 IMAGE Consortium	AI449090
6193 F	1 0219AMNS	Est2 repressor factor {IMAGE:894472}	894472 IMAGE Consortium	AA517533
6194 F	3 0219AMNS	IMAGE EST {IMAGE:536577}	536577 IMAGE Consortium	AA073904
6195 F	5 0219AMNS	EST	619004 IMAGE Consortium	AI449095
6196 F	7 0219AMNS	ESTs	619261 IMAGE Consortium	AA175763
6197 F	9 0219AMNS	ESTs	619415 IMAGE Consortium	AA172529
6198 F	11 0219AMNS	ESTs	618975 IMAGE Consortium	AA175485
6199 H	1 0219AMNS	IMAGE EST {IMAGE:932748}	932748 IMAGE Consortium	AA538419
6200 H	3 0219AMNS			
6201 H	5 0219AMNS	glycoprotein 49 B	619095 IMAGE Consortium	AA175494
6202 H	7 0219AMNS	ESTs	619289 IMAGE Consortium	AA172354
6203 H	9 0219AMNS	ESTs	619426 IMAGE Consortium	AA175623
6204 H	11 0219AMNS	ESTs	619781 IMAGE Consortium	AI604120
6205 B	1 021GAMNT	IMAGE EST {IMAGE:524177}	524177 IMAGE Consortium	AA065624
6206 B	3 021GAMNT	Mus musculus steroid receptor coactivator-1 (mSRC-1) mRNA, complete cds {IMAGE:523588}	523588 IMAGE Consortium	AA097341
6207 B	5 021GAMNT	ESTs, Weakly similar to KIAA0005 [H.sapiens]	532659 IMAGE Consortium	AA068436

		{IMAGE:532659}		
6208 B	7	021GAMNT ESTs {IMAGE:661821}	661821	IMAGE Consortium AA208426
6209 B	9	021GAMNT IMAGE EST {IMAGE:803629}	803629	IMAGE Consortium AA422805
6210 B	11	021GAMNT ESTs {IMAGE:317958}	317958	IMAGE Consortium W11432
6211 D	1	021GAMNT Mus musculus protein kinase C inhibitor (mPKCI) mRNA, complete cds {IMAGE:533117}	533117	IMAGE Consortium AA068901
6212 D	3	021GAMNT ESTs, Highly similar to HYPOTHETICAL 64.3 KD GTP-BINDING PROTEIN C02F5.3 IN CHROMOSOME III [Caenorhabditis elegans] {IMAGE:523945}	523945	IMAGE Consortium AA071987
6213 D	5	021GAMNT Lamin A {IMAGE:524119}	524119	IMAGE Consortium AA066180
6214 D	7	021GAMNT ESTs {IMAGE:662025}	662025	IMAGE Consortium AA207462
6215 D	9	021GAMNT ESTs {IMAGE:803897}	803897	IMAGE Consortium AA435252
6216 D	11	021GAMNT ESTs {IMAGE:314625}	314625	IMAGE Consortium W10903
6217 F	1	021GAMNT IMAGE EST {IMAGE:535288}	535288	IMAGE Consortium AA087768
6218 F	3	021GAMNT IMAGE EST {IMAGE:536030}	536030	IMAGE Consortium AA073247
6219 F	5	021GAMNT ESTs, Highly similar to ERYTHROCYTE ADDUCIN BETA SUBUNIT [Homo sapiens] {IMAGE:524196}	524196	IMAGE Consortium AA066202
6220 F	7	021GAMNT ESTs, Highly similar to HM-1 KILLER TOXIN RESISTANCE PROTEIN [Saccharomyces cerevisiae] {IMAGE:652131}	652131	IMAGE Consortium AA215144
6221 F	9	021GAMNT ESTs {IMAGE:944409}	944409	IMAGE Consortium AA545589
6222 F	11	021GAMNT IMAGE EST {IMAGE:314562}	314562	IMAGE Consortium W12782
6223 H	1	021GAMNT IMAGE EST {IMAGE:522544}	522544	IMAGE Consortium AA067977
6224 H	3	021GAMNT ESTs {IMAGE:534178}	534178	IMAGE Consortium AA072617
6225 H	5	021GAMNT Aquaporin 1 {IMAGE:656654}	656654	IMAGE Consortium AA241281
6226 H	7	021GAMNT ESTs, Weakly similar to ZK856.11 [C.elegans] {IMAGE:778381}	778381	IMAGE Consortium AA413555
6227 H	9	021GAMNT ESTs {IMAGE:961271}	961271	IMAGE Consortium AA547550
6228 H	11	021GAMNT ESTs {IMAGE:318817}	318817	IMAGE Consortium W34216
6229 B	1	021NAMNU		
6230 B	3	021NAMNU ESTs {IMAGE:403663}	403663	IMAGE Consortium W82199
6231 B	5	021NAMNU M.musculus mRNA for semaphorin D {IMAGE:423697}	423697	IMAGE Consortium W98303
6232 B	7	021NAMNU Wingless-related MMTV integration site 3A {IMAGE:426103}	426103	IMAGE Consortium AA000971
6233 B	9	021NAMNU ESTs {IMAGE:438463}	438463	IMAGE Consortium AA008251
6234 B	11	021NAMNU ESTs, Weakly similar to ORF YOR173w [S.cerevisiae] {IMAGE:441983}	441983	IMAGE Consortium AA015421
6235 D	1	021NAMNU Mus musculus mRNA for pyruvate dehydrogenase kinase-like protein {IMAGE:334876}	334876	IMAGE Consortium W17814
6236 D	3	021NAMNU ESTs, Highly similar to p82 [H.sapiens] {IMAGE:401305}	401305	IMAGE Consortium W98251
6237 D	5	021NAMNU ESTs {IMAGE:423276}	423276	IMAGE Consortium W98720
6238 D	7	021NAMNU ESTs {IMAGE:427248}	427248	IMAGE Consortium AA002937
6239 D	9	021NAMNU ESTs {IMAGE:437278}	437278	IMAGE Consortium AA003672

6240 D	11	021NAMNU	ESTs {IMAGE:445450}	445450 IMAGE Consortium	AA018015
6241 F	1	021NAMNU	ESTs {IMAGE:348362}	348362 IMAGE Consortium	W33467
6242 F	3	021NAMNU	ESTs {IMAGE:420131}	420131 IMAGE Consortium	W89800
6243 F	5	021NAMNU	IMAGE EST {IMAGE:421672}	421672 IMAGE Consortium	W98666
6244 F	7	021NAMNU	Vasoactive intestinal peptide receptor 2 {IMAGE:425925}	425925 IMAGE Consortium	AA000726
6245 F	9	021NAMNU	ESTs {IMAGE:439211}	439211 IMAGE Consortium	AA003898
6246 F	11	021NAMNU	Delta-like 1 homolog (Drosophila) {IMAGE:444023}	444023 IMAGE Consortium	AA015264
6247 H	1	021NAMNU	Mus musculus von Ebner minor salivary gland protein mRNA, complete cds {IMAGE:335127}	335127 IMAGE Consortium	W17563
6248 H	3	021NAMNU	IMAGE EST {IMAGE:418564}	418564 IMAGE Consortium	W99034
6249 H	5	021NAMNU	ESTs, Highly similar to unknown [H.sapiens] {IMAGE:421797}	421797 IMAGE Consortium	W99114
6250 H	7	021NAMNU	ESTs {IMAGE:426069}	426069 IMAGE Consortium	AA002913
6251 H	9	021NAMNU	ESTs {IMAGE:437755}	437755 IMAGE Consortium	AA008001
6252 H	11	021NAMNU	ESTs {IMAGE:456249}	456249 IMAGE Consortium	AA023245
6253 B	1	021UAMNV	ESTs, Highly similar to L-SERINE DEHYDRATASE [Rattus norvegicus] {IMAGE:463345}	463345 IMAGE Consortium	AA027702
6254 B	3	021UAMNV	IMAGE EST {IMAGE:472722}	472722 IMAGE Consortium	AA038172
6255 B	5	021UAMNV	ESTs	623028 IMAGE Consortium	AA185884
6256 B	7	021UAMNV	ESTs	622621 IMAGE Consortium	AI591925
6257 B	9	021UAMNV	nicotinamide N-methyltransferase	634474 IMAGE Consortium	AA185486
6258 B	11	021UAMNV	ESTs, Highly similar to POSSIBLE GUSTATORY RECEPTOR TYPE B [Bos taurus] {IMAGE:476044}	476044 IMAGE Consortium	AA050273
6259 D	1	021UAMNV	IMAGE EST {IMAGE:463157}	463157 IMAGE Consortium	AA030609
6260 D	3	021UAMNV	ESTs {IMAGE:473415}	473415 IMAGE Consortium	AA038443
6261 D	5	021UAMNV	ESTs	623054 IMAGE Consortium	AA185889
6262 D	7	021UAMNV	ESTs	621944 IMAGE Consortium	AA183713
6263 D	9	021UAMNV	ESTs	634604 IMAGE Consortium	AA189652
6264 D	11	021UAMNV	IMAGE EST {IMAGE:478788}	478788 IMAGE Consortium	AA051003
6265 F	1	021UAMNV	ESTs {IMAGE:463330}	463330 IMAGE Consortium	AA027604
6266 F	3	021UAMNV	IMAGE EST {IMAGE:475491}	475491 IMAGE Consortium	AA042014
6267 F	5	021UAMNV	ESTs, Weakly similar to KIAA0594 protein [H.sapiens]	622365 IMAGE Consortium	AI591902
6268 F	7	021UAMNV	DNA segment, Chr 6, ERATO Doi 365, expressed	622795 IMAGE Consortium	AA177625
6269 F	9	021UAMNV	ESTs {IMAGE:477796}	477796 IMAGE Consortium	AA048412
6270 F	11	021UAMNV	ESTs, Highly similar to C-1-TETRAHYDROFOLATE SYNTHASE, CYTOPLASMIC [Rattus norvegicus] {IMAGE:478383}	478383 IMAGE Consortium	AA049796
6271 H	1	021UAMNV	IMAGE EST {IMAGE:463006}	463006 IMAGE Consortium	AA030574
6272 H	3	021UAMNV	v-src suppressed transcript 1	621307 IMAGE Consortium	AA177318
6273 H	5	021UAMNV	ESTs, Weakly similar to Similarity to B.subtilis YQJC protein [C.elegans]	621671 IMAGE Consortium	AI591827
6274 H	7	021UAMNV	ESTs, Highly similar to AF121859_1 sorting nexin 9 [H.sapiens]	622825 IMAGE Consortium	AI591948

6275 H	9 021UAMNV	ESTs {IMAGE:477056}	477056 IMAGE Consortium	AA048139
6276 H	11 021UAMNV	ESTs {IMAGE:479311}	479311 IMAGE Consortium	AA048769
6277 B	1 0211AMNW	ESTs {IMAGE:483597}	483597 IMAGE Consortium	AA061662
6278 B	3 0211AMNW	ESTs {IMAGE:481362}	481362 IMAGE Consortium	AA060545
6279 B	5 0211AMNW	Mus musculus mRNA for translation initiation factor eIF2 gamma Y {IMAGE:573320}	573320 IMAGE Consortium	AA118716
6280 B	7 0211AMNW	ESTs, Moderately similar to serine/threonine protein kinase [M.musculus] {IMAGE:573966}	573966 IMAGE Consortium	AA118903
6281 B	9 0211AMNW	IMAGE EST {IMAGE:580763}	580763 IMAGE Consortium	AA145877
6282 B	11 0211AMNW			
6283 D	1 0211AMNW	IMAGE EST {IMAGE:483024}	483024 IMAGE Consortium	AA060880
6284 D	3 0211AMNW	Mouse mRNA for neuropsin, complete cds {IMAGE:483813}	483813 IMAGE Consortium	AA061507
6285 D	5 0211AMNW	IMAGE EST {IMAGE:573771}	573771 IMAGE Consortium	AA119029
6286 D	7 0211AMNW	ESTs, Highly similar to The ha3611 gene product is related to S.cerevisiae SNM1 protein. [H.sapiens] {IMAGE:576960}	576960 IMAGE Consortium	AA122530
6287 D	9 0211AMNW	ESTs {IMAGE:578345}	578345 IMAGE Consortium	AA120130
6288 D	11 0211AMNW	IMAGE EST {IMAGE:583669}	583669 IMAGE Consortium	AA153416
6289 F	1 0211AMNW	EST {IMAGE:481499}	481499 IMAGE Consortium	AA060615
6290 F	3 0211AMNW	ESTs {IMAGE:481867}	481867 IMAGE Consortium	AA059812
6291 F	5 0211AMNW	Mus musculus syntaxin 7 (Syn7) mRNA, complete cds {IMAGE:573479}	573479 IMAGE Consortium	AA111166
6292 F	7 0211AMNW	Myeloblastosis oncogene {IMAGE:575702}	575702 IMAGE Consortium	AA123009
6293 F	9 0211AMNW	ESTs {IMAGE:582936}	582936 IMAGE Consortium	AA153601
6294 F	11 0211AMNW	ESTs, Weakly similar to HYPOTHETICAL 67.0 KD PROTEIN IN PRE3-SAG1 INTERGENIC REGION [Saccharomyces cerevisiae] {IMAGE:597896}	597896 IMAGE Consortium	AA162148
6295 H	1 0211AMNW	IMAGE EST {IMAGE:483649}	483649 IMAGE Consortium	AA061366
6296 H	3 0211AMNW	IMAGE EST {IMAGE:484284}	484284 IMAGE Consortium	AA080236
6297 H	5 0211AMNW	IMAGE EST {IMAGE:575434}	575434 IMAGE Consortium	AA124928
6298 H	7 0211AMNW	IMAGE EST {IMAGE:576686}	576686 IMAGE Consortium	AA124417
6299 H	9 0211AMNW	ESTs {IMAGE:581007}	581007 IMAGE Consortium	AA139837
6300 H	11 0211AMNW	ESTs {IMAGE:583522}	583522 IMAGE Consortium	AA142685
6301 B	1 0218AMNX	Mouse mkf-1 mRNA, complete cds {IMAGE:596277}	596277 IMAGE Consortium	AA122720
6302 B	3 0218AMNX	IMAGE EST {IMAGE:618480}	618480 IMAGE Consortium	AA175279
6303 B	5 0218AMNX	ESTs {IMAGE:619687}	619687 IMAGE Consortium	AA175976
6304 B	7 0218AMNX	ESTs {IMAGE:617455}	617455 IMAGE Consortium	AA174582
6305 B	9 0218AMNX	ESTs {IMAGE:635025}	635025 IMAGE Consortium	AA163246
6306 B	11 0218AMNX	IMAGE EST {IMAGE:635565}	635565 IMAGE Consortium	AA163378
6307 D	1 0218AMNX	Histocompatibility 2, Q region locus 7 {IMAGE:596447}	596447 IMAGE Consortium	AA122791
6308 D	3 0218AMNX	Mus musculus protease-activated receptor 3 (PAR3) mRNA, complete cds {IMAGE:620379}	620379 IMAGE Consortium	AA177828

6309 D	5	0218AMNX	IMAGE EST {IMAGE:619861}	619861 IMAGE Consortium	AA177984
6310 D	7	0218AMNX	ESTs {IMAGE:619970}	619970 IMAGE Consortium	AA178026
6311 D	9	0218AMNX	IMAGE EST {IMAGE:634252}	634252 IMAGE Consortium	AA182981
6312 D	11	0218AMNX	IMAGE EST {IMAGE:635810}	635810 IMAGE Consortium	AA185313
6313 F	1	0218AMNX	Apoptosis inhibitor 2 {IMAGE:597715}	597715 IMAGE Consortium	AA144490
6314 F	3	0218AMNX	ESTs {IMAGE:617258}	617258 IMAGE Consortium	AA174859
6315 F	5	0218AMNX	Signal transducer and activator of transcription 4 {IMAGE:618949}	618949 IMAGE Consortium	AA175477
6316 F	7	0218AMNX	ESTs {IMAGE:619926}	619926 IMAGE Consortium	AA177733
6317 F	9	0218AMNX	ESTs {IMAGE:622383}	622383 IMAGE Consortium	AA184207
6318 F	11	0218AMNX	ESTs {IMAGE:634389}	634389 IMAGE Consortium	AA185231
6319 H	1	0218AMNX	IMAGE EST {IMAGE:583769}	583769 IMAGE Consortium	AA146401
6320 H	3	0218AMNX	ESTs {IMAGE:619079}	619079 IMAGE Consortium	AA170629
6321 H	5	0218AMNX	ESTs, Moderately similar to PLASMA-CELL MEMBRANE GLYCOPROTEIN PC-1 [Homo sapiens] {IMAGE:621166}	621166 IMAGE Consortium	AA177363
6322 H	7	0218AMNX	ESTs {IMAGE:619232}	619232 IMAGE Consortium	AA175750
6323 H	9	0218AMNX	ESTs {IMAGE:621820}	621820 IMAGE Consortium	AA184114
6324 H	11	0218AMNX	IMAGE EST {IMAGE:634581}	634581 IMAGE Consortium	AA189648
6325 B	1	021FAMNY	ESTs {IMAGE:622792}	622792 IMAGE Consortium	AA177621
6326 B	3	021FAMNY	ESTs {IMAGE:638302}	638302 IMAGE Consortium	AA200877
6327 B	5	021FAMNY	IMAGE EST {IMAGE:638244}	638244 IMAGE Consortium	AA200336
6328 B	7	021FAMNY	Hormone receptor {IMAGE:641865}	641865 IMAGE Consortium	AA209882
6329 B	9	021FAMNY	IMAGE EST {IMAGE:652735}	652735 IMAGE Consortium	AA217196
6330 B	11	021FAMNY	RNA binding motif protein, X chromosome	635076 IMAGE Consortium	AA163256
6331 D	1	021FAMNY	ESTs, Weakly similar to HIGH AFFINITY IMMUNOGLOBULIN GAMMA FC RECEPTOR I PRECURSOR [M.musculus] {IMAGE:622960}	622960 IMAGE Consortium	AA177761
6332 D	3	021FAMNY	ESTs {IMAGE:639990}	639990 IMAGE Consortium	AA197474
6333 D	5	021FAMNY	ESTs, Highly similar to The KIAA0136 gene product is novel. [H.sapiens] {IMAGE:638129}	638129 IMAGE Consortium	AA200976
6334 D	7	021FAMNY	ESTs, Highly similar to similar to human ZFY protein. [H.sapiens] {IMAGE:641892}	641892 IMAGE Consortium	AA209887
6335 D	9	021FAMNY	ESTs {IMAGE:642409}	642409 IMAGE Consortium	AA184882
6336 D	11	021FAMNY	ESTs	635170 IMAGE Consortium	AI592809
6337 F	1	021FAMNY	IMAGE EST {IMAGE:636318}	636318 IMAGE Consortium	AA189229
6338 F	3	021FAMNY	ESTs {IMAGE:637444}	637444 IMAGE Consortium	AA185052
6339 F	5	021FAMNY	IMAGE EST {IMAGE:639768}	639768 IMAGE Consortium	AA197519
6340 F	7	021FAMNY	ESTs {IMAGE:644857}	644857 IMAGE Consortium	AA210597
6341 F	9	021FAMNY	ESTs {IMAGE:643902}	643902 IMAGE Consortium	AA204090
6342 F	11	021FAMNY	ESTs	635227 IMAGE Consortium	AA189760
6343 H	1	021FAMNY	IMAGE EST {IMAGE:622182}	622182 IMAGE Consortium	AA183431
6344 H	3	021FAMNY	ESTs {IMAGE:639215}	639215 IMAGE Consortium	AA200989
6345 H	5	021FAMNY	ESTs {IMAGE:638593}	638593 IMAGE Consortium	AA200890

6346 H	7 021FAMNY	ESTs, Highly similar to zinc finger protein C2H2-171 [H.sapiens] {IMAGE:642975}	642975 IMAGE Consortium	AA185217
6347 H	9 021FAMNY	ESTs {IMAGE:644358}	644358 IMAGE Consortium	AA265226
6348 H	11 021FAMNY	sushi-repeat-containing protein, X chromosome	635234 IMAGE Consortium	AA189487
6349 B	1 021MAMNZ	ESTs	635399 IMAGE Consortium	AA216979
6350 B	3 021MAMNZ	EST	635648 IMAGE Consortium	AI451255
6351 B	5 021MAMNZ	ESTs, Highly similar to IMA3_MOUSE IMPORTIN ALPHA-3 SUBUNIT [M.musculus]	635985 IMAGE Consortium	AI645123
6352 B	7 021MAMNZ	tubulin, beta 4	318392 IMAGE Consortium	W13768
6353 B	9 021MAMNZ	small inducible cytokine subfamily D, 1	333711 IMAGE Consortium	W16003
6354 B	11 021MAMNZ	channel-interacting PDZ domain protein	356188 IMAGE Consortium	AI427358
6355 D	1 021MAMNZ	calcitonin receptor-like	634704 IMAGE Consortium	AA154791
6356 D	3 021MAMNZ	ESTs	634925 IMAGE Consortium	AI592775
6357 D	5 021MAMNZ	CCAAT/enhancer binding protein (C/EBP), delta	303612 IMAGE Consortium	W09210
6358 D	7 021MAMNZ	Sfpi1/PU.1 interaction partner	330172 IMAGE Consortium	AI325336
6359 D	9 021MAMNZ	exocyst component protein 70 kDa homolog (S. cerevisiae)	334130 IMAGE Consortium	W16220
6360 D	11 021MAMNZ	transcription factor 7-like 1	368160 IMAGE Consortium	W54087
6361 F	1 021MAMNZ	ESTs	634711 IMAGE Consortium	AI592746
6362 F	3 021MAMNZ	ESTs, Highly similar to EVI2B PROTEIN PRECURSOR TROPIC VIRAL INTEGRATION SITE 2B PROTEIN) [Homo sapiens]	634957 IMAGE Consortium	AI893742
6363 F	5 021MAMNZ	chitinase 3-like 1	313884 IMAGE Consortium	W10705
6364 F	7 021MAMNZ	early B-cell factor 3	331106 IMAGE Consortium	W14732
6365 F	9 021MAMNZ	ADP-ribosyltransferase 1	334682 IMAGE Consortium	W19016
6366 F	11 021MAMNZ	transforming growth factor, beta 3	368545 IMAGE Consortium	W53865
6367 H	1 021MAMNZ	ESTs	634781 IMAGE Consortium	AA163353
6368 H	3 021MAMNZ	ESTs	634988 IMAGE Consortium	AI592785
6369 H	5 021MAMNZ	small inducible cytokine A21a (leucine)	316441 IMAGE Consortium	W11170
6370 H	7 021MAMNZ	leucine rich protein, B7 gene	331513 IMAGE Consortium	W13869
6371 H	9 021MAMNZ	Moloney leukemia virus 10	334747 IMAGE Consortium	W36421
6372 H	11 021MAMNZ	phospholipase D2	374955 IMAGE Consortium	W61990
6373 B	1 021MAMOO	zinc finger protein 57	404508 IMAGE Consortium	AI893971
6374 B	3 021MAMOO	low density lipoprotein receptor	426185 IMAGE Consortium	AA003138
6375 B	5 021MAMOO	metallothionein 3	478292 IMAGE Consortium	AA049749
6376 B	7 021MAMOO	RAS, guanyl releasing protein 2	478922 IMAGE Consortium	AA048933
6377 B	9 021MAMOO	eukaryotic translation initiation factor 2 alpha kinase 2	597390 IMAGE Consortium	AA153156
6378 B	11 021MAMOO	heat shock protein, 70 kDa 3	582093 IMAGE Consortium	AA145895
6379 D	1 021MAMOO	elastin	372899 IMAGE Consortium	W59510
6380 D	3 021MAMOO	glycoprotein 49 B	440614 IMAGE Consortium	AI894044
6381 D	5 021MAMOO	forkhead box D3	479327 IMAGE Consortium	AI323053
6382 D	7 021MAMOO	factor 8-associated gene A	572416 IMAGE Consortium	AA105827
6383 D	9 021MAMOO	2'-5' oligoadenylate synthetase 1A	598851 IMAGE Consortium	AI322733



6384 D	11 021MAM00	tumor necrosis factor (ligand) superfamily, member 11	618692 IMAGE Consortium	AA170348
6385 F	1 021MAM00	sema domain, transmembrane domain (TM), and cytoplasmic domain, (semaphorin) 6B	387447 IMAGE Consortium	W65836
6386 F	3 021MAM00	cofilin 2, muscle	443000 IMAGE Consortium	AA014178
6387 F	5 021MAM00	fat specific gene 27	482917 IMAGE Consortium	AA062151
6388 F	7 021MAM00	Fc receptor, IgG, high affinity I	576062 IMAGE Consortium	AA120333
6389 F	9 021MAM00	apolipoprotein B editing complex 1	618120 IMAGE Consortium	AA175231
6390 F	11 021MAM00	LIM-domain containing, protein kinase	635736 IMAGE Consortium	AA165899
6391 H	1 021MAM00	DNA segment, Chr 17, Wayne State University 76, expressed	391583 IMAGE Consortium	AI430217
6392 H	3 021MAM00			
6393 H	5 021MAM00	0-6-methylguanine-DNA methyltransferase	493108 IMAGE Consortium	AA097896
6394 H	7 021MAM00	klotho	578480 IMAGE Consortium	AA162364
6395 H	9 021MAM00	a disintegrin and metalloproteinase domain 1a (fertilin alpha)	619528 IMAGE Consortium	AA172412
6396 H	11 021MAM00	coagulation factor VIII	637486 IMAGE Consortium	AA184901
6397 B	1 021TAM01	purinergic receptor P2Y, G-protein coupled 1	671513 IMAGE Consortium	AA241971
6398 B	3 021TAM01	cadherin 4	694334 IMAGE Consortium	AA221467
6399 B	5 021TAM01	kit ligand	657946 IMAGE Consortium	AA198236
6400 B	7 021TAM01	nuclear receptor subfamily 1, group H, member 4	738199 IMAGE Consortium	AA270904
6401 B	9 021TAM01	small inducible cytokine A5	832342 IMAGE Consortium	AA426997
6402 B	11 021TAM01	solute carrier family 22 (organic cation transporter)-like 2	571096 IMAGE Consortium	AA109757
6403 D	1 021TAM01	flavin containing monooxygenase 3	676386 IMAGE Consortium	AA209044
6404 D	3 021TAM01	myelin transcription factor 1	716942 IMAGE Consortium	AA266095
6405 D	5 021TAM01	SFFV proviral integration 1	671036 IMAGE Consortium	AA239990
6406 D	7 021TAM01	deiodinase, iodothyronine, type I	748113 IMAGE Consortium	AA260525
6407 D	9 021TAM01	ryanodine receptor 2, cardiac	871232 IMAGE Consortium	AA462877
6408 D	11 021TAM01	UDP-glucose ceramide glucosyltransferase	577841 IMAGE Consortium	AA116263
6409 F	1 021TAM01	P glycoprotein 2	678733 IMAGE Consortium	AA239419
6410 F	3 021TAM01	Fc receptor, IgE, low affinity II, alpha polypeptide	717625 IMAGE Consortium	AA250331
6411 F	5 021TAM01	carboxylesterase 1	680175 IMAGE Consortium	AI451839
6412 F	7 021TAM01	sodium channel, voltage-gated, type VIII, alpha polypeptide	748459 IMAGE Consortium	AA272867
6413 F	9 021TAM01	protein kinase C, alpha	920570 IMAGE Consortium	AA541955
6414 F	11 021TAM01	Tnf receptor-associated factor 1	636225 IMAGE Consortium	AA183167
6415 H	1 021TAM01	solute carrier family 2 (facilitated glucose transporter), member 2	680470 IMAGE Consortium	AA237317
6416 H	3 021TAM01	erythropoietin receptor	722259 IMAGE Consortium	AA254203
6417 H	5 021TAM01	bile acid-Coenzyme A dehydrogenase: amino acid n-acyltransferase	681783 IMAGE Consortium	AA244588
6418 H	7 021TAM01	interleukin 2	750652 IMAGE Consortium	AA290106
6419 H	9 021TAM01	cadherin 5	963176 IMAGE Consortium	AA553029
6420 H	11 021TAM01	ATPase, H+/K+ transporting, alpha polypeptide	670834 IMAGE Consortium	AA221980

6421 B	1	0210AMO2	ESTs	638240	IMAGE Consortium	AA200334
6422 B	3	0210AMO2	ESTs	639438	IMAGE Consortium	AA198595
6423 B	5	0210AMO2	ESTs, Highly similar to unnamed protein product [H.sapiens]	640166	IMAGE Consortium	AA197627
6424 B	7	0210AMO2	ESTs	636923	IMAGE Consortium	AA189446
6425 B	9	0210AMO2	ESTs	636368	IMAGE Consortium	AI645171
6426 B	11	0210AMO2	ESTs	637257	IMAGE Consortium	AA189907
6427 D	1	0210AMO2	ESTs	639083	IMAGE Consortium	AA199568
6428 D	3	0210AMO2	ESTs	638751	IMAGE Consortium	AI646187
6429 D	5	0210AMO2	ESTs	640197	IMAGE Consortium	AA197709
6430 D	7	0210AMO2	ESTs	636947	IMAGE Consortium	AA185123
6431 D	9	0210AMO2	POU domain, class 2, associating factor 1	636390	IMAGE Consortium	AA183024
6432 D	11	0210AMO2	histocompatibility 2, O region alpha locus	636533	IMAGE Consortium	AA182323
6433 F	1	0210AMO2	desert hedgehog homolog, (Drosophila)	639130	IMAGE Consortium	AI645932
6434 F	3	0210AMO2	ESTs	639604	IMAGE Consortium	AA198887
6435 F	5	0210AMO2	ESTs	640460	IMAGE Consortium	AA204188
6436 F	7	0210AMO2	ESTs	636958	IMAGE Consortium	AI645218
6437 F	9	0210AMO2	ESTs	637112	IMAGE Consortium	AI645244
6438 F	11	0210AMO2	ESTs	636564	IMAGE Consortium	AA183014
6439 H	1	0210AMO2	ESTs	639180	IMAGE Consortium	AI645939
6440 H	3	0210AMO2	ESTs	638924	IMAGE Consortium	AI645907
6441 H	5	0210AMO2	ESTs, Highly similar to embryonic lung protein [H.sapiens]	640532	IMAGE Consortium	AA204221
6442 H	7	0210AMO2	heterogeneous nuclear ribonucleoprotein A/B	636188	IMAGE Consortium	AA183307
6443 H	9	0210AMO2	ESTs	636435	IMAGE Consortium	AA183258
6444 H	11	0210AMO2	ESTs	636607	IMAGE Consortium	AA183295
6445 B	1	0217AMO3	spleen tyrosine kinase	642318	IMAGE Consortium	AA185105
6446 B	3	0217AMO3	ESTs	642537	IMAGE Consortium	AI606029
6447 B	5	0217AMO3	ESTs	336806	IMAGE Consortium	AI390970
6448 B	7	0217AMO3	ESTs	336460	IMAGE Consortium	AI390949
6449 B	9	0217AMO3	ESTs	643317	IMAGE Consortium	AA189897
6450 B	11	0217AMO3	ESTs	642805	IMAGE Consortium	AA189214
6451 D	1	0217AMO3	CD44 antigen	642342	IMAGE Consortium	AA185112
6452 D	3	0217AMO3	B-cell leukemia/lymphoma 2 related protein A1d	642577	IMAGE Consortium	AA185513
6453 D	5	0217AMO3	ESTs, Moderately similar to unnamed protein product [H.sapiens]	337790	IMAGE Consortium	AI391134
6454 D	7	0217AMO3	ESTs	335894	IMAGE Consortium	W17965
6455 D	9	0217AMO3	ESTs	643360	IMAGE Consortium	AA189744
6456 D	11	0217AMO3	ESTs	642831	IMAGE Consortium	AA189220
6457 F	1	0217AMO3	sema domain, transmembrane domain (TM), and cytoplasmic domain, (semaphorin) 6C	642361	IMAGE Consortium	AI606008
6458 F	3	0217AMO3	ESTs	642603	IMAGE Consortium	AI606034
6459 F	5	0217AMO3	ESTs	337921	IMAGE Consortium	AI391001
6460 F	7	0217AMO3	ESTs	644983	IMAGE Consortium	AI450476

6461 F	9 0217AMO3	ESTs	644118 IMAGE Consortium	AI450439
6462 F	11 0217AMO3	dishevelled, dsh homolog (Drosophila)	642837 IMAGE Consortium	AA185273
6463 H	1 0217AMO3	Sjogren syndrome antigen B	642393 IMAGE Consortium	AA184809
6464 H	3 0217AMO3	ESTs	642610 IMAGE Consortium	AA184889
6465 H	5 0217AMO3	ESTs	347817 IMAGE Consortium	W29750
6466 H	7 0217AMO3	lymphocyte cytosolic protein 2	643111 IMAGE Consortium	AA189872
6467 H	9 0217AMO3	ESTs	643434 IMAGE Consortium	AI451661
6468 H	11 0217AMO3	REV1 protein	642872 IMAGE Consortium	AA189247
6469 B	1 021EAMO4	ESTs	642950 IMAGE Consortium	AA185188
6470 B	3 021EAMO4	protein tyrosine phosphatase, non-receptor type 5	670128 IMAGE Consortium	AA221322
6471 B	5 021EAMO4	ESTs, Moderately similar to NEUROMEDIN B PRECURSOR [Homo sapiens]	658885 IMAGE Consortium	AA217878
6472 B	7 021EAMO4	ESTs	660269 IMAGE Consortium	AA220774
6473 B	9 021EAMO4	ESTs, Highly similar to AF171102_1 retinal degeneration B beta [H.sapiens]	660688 IMAGE Consortium	AA230387
6474 B	11 021EAMO4	ESTs	653605 IMAGE Consortium	AI390751
6475 D	1 021EAMO4	ESTs	643687 IMAGE Consortium	AA204045
6476 D	3 021EAMO4	EST	660971 IMAGE Consortium	AI464420
6477 D	5 021EAMO4	ESTs	658913 IMAGE Consortium	AI390431
6478 D	7 021EAMO4	ESTs, Highly similar to otoraplin [M.musculus]	661055 IMAGE Consortium	AA222797
6479 D	9 021EAMO4	ESTs, Weakly similar to U83194_1 TRAF4-associated factor 2 [H.sapiens]	670148 IMAGE Consortium	AA231064
6480 D	11 021EAMO4	cytochrome c oxidase, subunit VIIC	652935 IMAGE Consortium	AA217122
6481 F	1 021EAMO4	ESTs	642987 IMAGE Consortium	AA185587
6482 F	3 021EAMO4	ESTs	670568 IMAGE Consortium	AA222016
6483 F	5 021EAMO4	ESTs	659653 IMAGE Consortium	AA219919
6484 F	7 021EAMO4	nuclear receptor subfamily 2, group F, member 1	669736 IMAGE Consortium	AA230392
6485 F	9 021EAMO4	ESTs, Highly similar to thyroid hormone receptor-associated protein complex component TRAP240 [H.sapiens]	660729 IMAGE Consortium	AI644845
6486 F	11 021EAMO4	ATP-binding cassette, sub-family G (WHITE), member 2	652993 IMAGE Consortium	AA217253
6487 H	1 021EAMO4	ESTs	643707 IMAGE Consortium	AI390472
6488 H	3 021EAMO4	G1 to phase transition 2	659212 IMAGE Consortium	AA218194
6489 H	5 021EAMO4	ESTs, Highly similar to AA2A_MOUSE ADENOSINE A2A RECEPTOR [M.musculus]	658987 IMAGE Consortium	AI390437
6490 H	7 021EAMO4	ESTs	661114 IMAGE Consortium	AA222660
6491 H	9 021EAMO4	ESTs	652713 IMAGE Consortium	AA217243
6492 H	11 021EAMO4	ATPase, aminophospholipid transporter-like, class I, type 8A, member 2	658095 IMAGE Consortium	AI390382
6493 B	1 021LAMO5	ESTs	653188 IMAGE Consortium	AA217118
6494 B	3 021LAMO5	ESTs	652704 IMAGE Consortium	AI390603
6495 B	5 021LAMO5	pyruvate kinase liver and red blood cell	680323 IMAGE Consortium	AA237607
6496 B	7 021LAMO5	ESTs	677565 IMAGE Consortium	AI390275

6497 B	9 021LAMO5	ESTs, Highly similar to GTPASE-ACTIVATING PROTEIN [Homo sapiens]	697237 IMAGE Consortium	AI425461
6498 B	11 021LAMO5	ESTs	680754 IMAGE Consortium	AA242553
6499 D	1 021LAMO5	ESTs	653231 IMAGE Consortium	AA217433
6500 D	3 021LAMO5	ESTs	678856 IMAGE Consortium	AI390305
6501 D	5 021LAMO5	ESTs, Weakly similar to 3-MERCAPTOPYRUVATE SULFURTRANSFERASE [H.sapiens]	680334 IMAGE Consortium	AA250153
6502 D	7 021LAMO5	ESTs	677623 IMAGE Consortium	AA212925
6503 D	9 021LAMO5	ESTs	697256 IMAGE Consortium	AI425463
6504 D	11 021LAMO5	mannan-binding lectin serine protease 1	681501 IMAGE Consortium	AA237632
6505 F	1 021LAMO5	ESTs	652553 IMAGE Consortium	AA217488
6506 F	3 021LAMO5	ESTs	679615 IMAGE Consortium	AA242681
6507 F	5 021LAMO5	ESTs, Weakly similar to KIAA0982 protein [H.sapiens]	671688 IMAGE Consortium	AI391340
6508 F	7 021LAMO5	ESTs	679213 IMAGE Consortium	AA245987
6509 F	9 021LAMO5	solute carrier family 6 (neurotransmitter transporter), member 14	697546 IMAGE Consortium	AA230972
6510 F	11 021LAMO5	ESTs	681582 IMAGE Consortium	AA237735
6511 H	1 021LAMO5	purine-nucleoside phosphorylase	652603 IMAGE Consortium	AI390585
6512 H	3 021LAMO5	procollagen, type IX, alpha 3	671056 IMAGE Consortium	AA239995
6513 H	5 021LAMO5	cytochrome P450, 39a1 (oxysterol 7alpha-hydroxylase)	680424 IMAGE Consortium	AI451832
6514 H	7 021LAMO5	ESTs	679346 IMAGE Consortium	AI390325
6515 H	9 021LAMO5	CEA-related cell adhesion molecule 1	681026 IMAGE Consortium	AA242637
6516 H	11 021LAMO5	ESTs	681598 IMAGE Consortium	AA237974
6517 B	1 021SAMO6	ESTs, Moderately similar to AF163315_1 putative N-acetyltransferase Camello 2 [M.musculus]	693021 IMAGE Consortium	AI466969
6518 B	3 021SAMO6	ESTs	693853 IMAGE Consortium	AI425378
6519 B	5 021SAMO6	ESTs	719317 IMAGE Consortium	AI464752
6520 B	7 021SAMO6	ESTs	718936 IMAGE Consortium	AI464682
6521 B	9 021SAMO6	hematopoietic cell specific Lyn substrate 1	720655 IMAGE Consortium	AA267468
6522 B	11 021SAMO6	EST	717726 IMAGE Consortium	AI661098
6523 D	1 021SAMO6	interferon activated gene 203	692306 IMAGE Consortium	AA238257
6524 D	3 021SAMO6	ESTs	719060 IMAGE Consortium	AI464705
6525 D	5 021SAMO6	ESTs	719337 IMAGE Consortium	AA255071
6526 D	7 021SAMO6	ESTs	719887 IMAGE Consortium	AI464795
6527 D	9 021SAMO6	ESTs	720676 IMAGE Consortium	AI464833
6528 D	11 021SAMO6	ESTs	718526 IMAGE Consortium	AA266331
6529 F	1 021SAMO6	ESTs, Moderately similar to hypothetical protein [H.sapiens]	692582 IMAGE Consortium	AI390633
6530 F	3 021SAMO6	ESTs	719105 IMAGE Consortium	AI464714
6531 F	5 021SAMO6	ESTs	719360 IMAGE Consortium	AI464758
6532 F	7 021SAMO6	ESTs	719964 IMAGE Consortium	AI464799
6533 F	9 021SAMO6	ESTs	717603 IMAGE Consortium	AI466892
6534 F	11 021SAMO6	ESTs	718554 IMAGE Consortium	AI464616

6535 H	1 021SAMO6	cytochrome P450, 26, retinoic acid	692620 IMAGE Consortium	AA239785
6536 H	3 021SAMO6	ESTs	719120 IMAGE Consortium	AA267295
6537 H	5 021SAMO6	ESTs	719422 IMAGE Consortium	AA254700
6538 H	7 021SAMO6	ESTs	720135 IMAGE Consortium	AA254866
6539 H	9 021SAMO6	ESTs	718357 IMAGE Consortium	AI464593
6540 H	11 021SAMO6	ESTs	717839 IMAGE Consortium	AI466905
6541 B	1 021ZAMO7	ESTs	718718 IMAGE Consortium	AI464648
6542 B	3 021ZAMO7	ESTs	718087 IMAGE Consortium	AA266447
6543 B	5 021ZAMO7	ESTs, Weakly similar to putative phosphoinositide 5-phosphatase type II [M.musculus]	721258 IMAGE Consortium	AA267017
6544 B	7 021ZAMO7	ESTs	722179 IMAGE Consortium	AI608059
6545 B	9 021ZAMO7	ESTs	720890 IMAGE Consortium	AI464866
6546 B	11 021ZAMO7	ESTs	721868 IMAGE Consortium	AI595315
6547 D	1 021ZAMO7	ESTs	718723 IMAGE Consortium	AI464655
6548 D	3 021ZAMO7	ESTs	718142 IMAGE Consortium	AI466953
6549 D	5 021ZAMO7	ESTs	722021 IMAGE Consortium	AI608045
6550 D	7 021ZAMO7	ESTs, Highly similar to Rab8-interacting protein [M.musculus]	721466 IMAGE Consortium	AA267626
6551 D	9 021ZAMO7	ESTs	720901 IMAGE Consortium	AA267210
6552 D	11 021ZAMO7	ESTs, Moderately similar to estradiol 17beta-dehydrogenase [M.musculus]	721959 IMAGE Consortium	AI595330
6553 F	1 021ZAMO7	exostoses (multiple) 2	699160 IMAGE Consortium	AI466760
6554 F	3 021ZAMO7	ESTs	718313 IMAGE Consortium	AA261105
6555 F	5 021ZAMO7	ESTs	722047 IMAGE Consortium	AA268472
6556 F	7 021ZAMO7	ESTs	720763 IMAGE Consortium	AI464847
6557 F	9 021ZAMO7	ESTs	721680 IMAGE Consortium	AI608034
6558 F	11 021ZAMO7	glutamate receptor, ionotropic, kainate 5 (gamma 2)	733368 IMAGE Consortium	AA261334
6559 H	1 021ZAMO7	ESTs, Moderately similar to hypothetical protein, similar to AF033339 [H.sapiens]	699291 IMAGE Consortium	AA245397
6560 H	3 021ZAMO7	ESTs	721006 IMAGE Consortium	AI464880
6561 H	5 021ZAMO7	ESTs, Moderately similar to NICE-5 protein [H.sapiens]	722076 IMAGE Consortium	AI608050
6562 H	7 021ZAMO7	ESTs	721508 IMAGE Consortium	AA266890
6563 H	9 021ZAMO7	transforming growth factor beta regulated gene 1	720975 IMAGE Consortium	AI464879
6564 H	11 021ZAMO7	ESTs, Moderately similar to F26L_MOUSE 6PF-2-K/FRU-2,6-P2ASE LIVER ISOZYME [M.musculus]	735122 IMAGE Consortium	AI529513
6565 B	1 0216AMO8	ESTs	734904 IMAGE Consortium	AI529488
6566 B	3 0216AMO8	ESTs	723184 IMAGE Consortium	AI463750
6567 B	5 0216AMO8	ESTs	722759 IMAGE Consortium	AI608126
6568 B	7 0216AMO8	transcription factor EB	733533 IMAGE Consortium	AA261233
6569 B	9 0216AMO8	ESTs	749292 IMAGE Consortium	AA288545
6570 B	11 0216AMO8	ESTs, Highly similar to GFG_RAT PROTEIN GFG [R.norvegicus]	748869 IMAGE Consortium	AA288191
6571 D	1 0216AMO8	formin binding protein 3	734948 IMAGE Consortium	AA261241

6572 D	3 0216AMO8	ESTs	722478 IMAGE Consortium	AA253984
6573 D	5 0216AMO8	ESTs	722797 IMAGE Consortium	AI608132
6574 D	7 0216AMO8	procollagen, type XI, alpha 1	734278 IMAGE Consortium	AA268923
6575 D	9 0216AMO8	ESTs	749367 IMAGE Consortium	AI644813
6576 D	11 0216AMO8	ESTs	749616 IMAGE Consortium	AA388702
6577 F	1 0216AMO8	ESTs	722263 IMAGE Consortium	AI608071
6578 F	3 0216AMO8	hypothetical protein	723209 IMAGE Consortium	AI463754
6579 F	5 0216AMO8	ESTs	722815 IMAGE Consortium	AI608136
6580 F	7 0216AMO8	ESTs	734341 IMAGE Consortium	AA268699
6581 F	9 0216AMO8	peroxisomal biogenesis factor 11a	747923 IMAGE Consortium	AA260931
6582 F	11 0216AMO8	ESTs	748989 IMAGE Consortium	AA288043
6583 H	1 0216AMO8	ESTs	723030 IMAGE Consortium	AI463724
6584 H	3 0216AMO8	ESTs	723226 IMAGE Consortium	AI463757
6585 H	5 0216AMO8	ESTs, Moderately similar to KIAA0986 protein [H.sapiens]	723564 IMAGE Consortium	AA268558
6586 H	7 0216AMO8	ESTs, Weakly similar to 1010301A contrapsin [M.musculus]	748140 IMAGE Consortium	AI644723
6587 H	9 0216AMO8	S100 calcium-binding protein A13	747966 IMAGE Consortium	AA260961
6588 H	11 0216AMO8	ESTs	749718 IMAGE Consortium	AA388770
6589 B	1 021DAMO9	ESTs	750159 IMAGE Consortium	AI606525
6590 B	3 021DAMO9	ESTs, Highly similar to schlafen2 [M.musculus]	334371 IMAGE Consortium	W36524
6591 B	5 021DAMO9	ameloblastin	334870 IMAGE Consortium	W36463
6592 B	7 021DAMO9	ESTs	738309 IMAGE Consortium	AA271129
6593 B	9 021DAMO9	ESTs	750443 IMAGE Consortium	AA390015
6594 B	11 021DAMO9	ESTs	751388 IMAGE Consortium	AA396310
6595 D	1 021DAMO9	carbonic anhydrase 1	750203 IMAGE Consortium	AA396359
6596 D	3 021DAMO9	ESTs, Highly similar to Y176_HUMAN HYPOTHETICAL PROTEIN KIAA0176 [H.sapiens]	334413 IMAGE Consortium	W18281
6597 D	5 021DAMO9	ESTs, Highly similar to GLANDULAR KALLIKREIN 2 PRECURSOR [Homo sapiens]	334909 IMAGE Consortium	AI391329
6598 D	7 021DAMO9	DNA segment, Human S2298E	737614 IMAGE Consortium	AI509041
6599 D	9 021DAMO9	ESTs, Weakly similar to F29B9.1 [C.elegans]	751178 IMAGE Consortium	AI508691
6600 D	11 021DAMO9	ESTs, Weakly similar to MRP5_MOUSE MULTIDRUG RESISTANCE-ASSOCIATED PROTEIN 5 [M.musculus]	750693 IMAGE Consortium	AA288247
6601 F	1 021DAMO9	cholinergic receptor, nicotinic, beta polypeptide 1 (muscle)	750210 IMAGE Consortium	AA288642
6602 F	3 021DAMO9	ESTs, Weakly similar to TRLC_DROME TRANSIENT RECEPTOR POTENTIAL LOCUS C PROTEIN PRECURSOR [D.melanogaster]	334445 IMAGE Consortium	W36621
6603 F	5 021DAMO9	ESTs	736356 IMAGE Consortium	AA270575
6604 F	7 021DAMO9	ATP-binding cassette, sub-family A (ABC1), member 8	737699 IMAGE Consortium	AI509048
6605 F	9 021DAMO9	ESTs	751190 IMAGE Consortium	AI508693
6606 F	11 021DAMO9	ESTs, Highly similar to T02672 hypothetical protein R31449_3 - human [H.sapiens]	751403 IMAGE Consortium	AI508714
6607 H	1 021DAMO9	interferon regulatory factor 2	334512 IMAGE Consortium	AI893892

6608 H	3 021DAMO9	ESTs	334477 IMAGE Consortium	AI391310
6609 H	5 021DAMO9	ESTs	735655 IMAGE Consortium	AA269685
6610 H	7 021DAMO9	DNA segment, Chr 4, ERATO Doi 321, expressed	746496 IMAGE Consortium	AA260117
6611 H	9 021DAMO9	ESTs, Moderately similar to Y050_HUMAN HYPOTHETICAL PROTEIN KIAA0050 [H.sapiens]	750531 IMAGE Consortium	AI508471
6612 H	11 021DAMO9	ESTs	751460 IMAGE Consortium	AA396369
6613 B	1 021XAMOA	ESTs	751620 IMAGE Consortium	AI508742
6614 B	3 021XAMOA	ESTs	751972 IMAGE Consortium	AA396245
6615 B	5 021XAMOA	ESTs, Weakly similar to BIT [M.musculus]	750344 IMAGE Consortium	AI606547
6616 B	7 021XAMOA	ESTs	752473 IMAGE Consortium	AI508596
6617 B	9 021XAMOA	ESTs	761713 IMAGE Consortium	AI508993
6618 B	11 021XAMOA	ESTs	764195 IMAGE Consortium	AA285799
6619 D	1 021XAMOA	ESTs	750901 IMAGE Consortium	AA290154
6620 D	3 021XAMOA	ESTs	751982 IMAGE Consortium	AI508547
6621 D	5 021XAMOA	killer cell lectin-like receptor, subfamily A, member 1	752181 IMAGE Consortium	AA289069
6622 D	7 021XAMOA	ESTs	752508 IMAGE Consortium	AA290361
6623 D	9 021XAMOA	ESTs	762523 IMAGE Consortium	AI595041
6624 D	11 021XAMOA	acetylcholinesterase	762761 IMAGE Consortium	AA271549
6625 F	1 021XAMOA	ESTs	751639 IMAGE Consortium	AI508747
6626 F	3 021XAMOA	regulator of G-protein signaling 16	750226 IMAGE Consortium	AA390086
6627 F	5 021XAMOA	ESTs, Weakly similar to alternatively spliced form [H.sapiens]	752166 IMAGE Consortium	AI508554
6628 F	7 021XAMOA	ESTs	752513 IMAGE Consortium	AI508601
6629 F	9 021XAMOA	ESTs	762621 IMAGE Consortium	AI595047
6630 F	11 021XAMOA	ESTs	764276 IMAGE Consortium	AA285420
6631 H	1 021XAMOA	ESTs, Weakly similar to ORF2 [M.musculus]	751658 IMAGE Consortium	AA396084
6632 H	3 021XAMOA	EST AI425994	750260 IMAGE Consortium	AA396777
6633 H	5 021XAMOA	ESTs	752191 IMAGE Consortium	AI508560
6634 H	7 021XAMOA	ESTs	762123 IMAGE Consortium	AA277192
6635 H	9 021XAMOA	ESTs	762633 IMAGE Consortium	AI595049
6636 H	11 021XAMOA	cathepsin F	764342 IMAGE Consortium	AA277533
6637 B	1 0214AMOB	spermidine/spermine N1-acetyl transferase	778311 IMAGE Consortium	AA413899
6638 B	3 0214AMOB	ESTs, Highly similar to KIAA1468 protein [H.sapiens]	778486 IMAGE Consortium	AA413514
6639 B	5 0214AMOB	serum albumin variant	778707 IMAGE Consortium	AA413684
6640 B	7 0214AMOB	ESTs	777258 IMAGE Consortium	AA276944
6641 B	9 0214AMOB	TATA box binding protein (TBP)-associated factor, RNA polymerase II, H, 30 kDa	777561 IMAGE Consortium	AA276759
6642 B	11 0214AMOB	ESTs	765355 IMAGE Consortium	AA275254
6643 D	1 0214AMOB	ESTs, Weakly similar to unnamed protein product [M.musculus]	778315 IMAGE Consortium	AA413868
6644 D	3 0214AMOB	enhancer of zeste homolog 2 (Drosophila)	779236 IMAGE Consortium	AA445767
6645 D	5 0214AMOB	ESTs	778752 IMAGE Consortium	AA413909
6646 D	7 0214AMOB			

6647 D	9 0214AMOB	natural killer cell BY55 precursor	777636 IMAGE Consortium	AA276726
6648 D	11 0214AMOB	ESTs	764638 IMAGE Consortium	AA273374
6649 F	1 0214AMOB	ESTs, Weakly similar to HYPOTHETICAL 97.1 KD PROTEIN R05D3.4 IN CHROMOSOME III [Caenorhabditis elegans]	779069 IMAGE Consortium	AA413953
6650 F	3 0214AMOB	ESTs, Highly similar to BETA-ADAPTIN [Homo sapiens; Rattus norvegicus; Bos taurus]	778523 IMAGE Consortium	AA413703
6651 F	5 0214AMOB	Mus musculus antigen containing epitope to monoclonal antibody MMS-85/12 mRNA, partial cds	778780 IMAGE Consortium	AA413916
6652 F	7 0214AMOB	ESTs, Weakly similar to unknown protein [R.norvegicus]	777325 IMAGE Consortium	AI464239
6653 F	9 0214AMOB	bromodomain adjacent to zinc finger domain, 1B	777800 IMAGE Consortium	AA286377
6654 F	11 0214AMOB	ESTs	765454 IMAGE Consortium	AA275308
6655 H	1 0214AMOB	ESTs	778365 IMAGE Consortium	AA413472
6656 H	3 0214AMOB	ESTs	778525 IMAGE Consortium	AA413704
6657 H	5 0214AMOB	ESTs	778808 IMAGE Consortium	AA414064
6658 H	7 0214AMOB	ESTs, Weakly similar to nuclear dual-specificity phosphatase [H.sapiens]	776613 IMAGE Consortium	AA275909
6659 H	9 0214AMOB	ESTs	777834 IMAGE Consortium	AI508931
6660 H	11 0214AMOB	ESTs	765533 IMAGE Consortium	AA275116
6661 B	1 021BAMOC	ESTs	776163 IMAGE Consortium	AA276095
6662 B	3 021BAMOC	UDP-Gal:betaGlcNAc beta 1,4-galactosyltransferase, polypeptide 6	779077 IMAGE Consortium	AA414080
6663 B	5 021BAMOC	embryonic ectoderm development	779364 IMAGE Consortium	AA414109
6664 B	7 021BAMOC	ESTs, Highly similar to REPLICATION PROTEIN A 14 KD SUBUNIT [Homo sapiens]	779479 IMAGE Consortium	AA414248
6665 B	9 021BAMOC	ESTs	779734 IMAGE Consortium	AA414636
6666 B	11 021BAMOC	ATPase, Cu <sup>++</sup> transporting, alpha polypeptide	791019 IMAGE Consortium	AA415192
6667 D	1 021BAMOC	serine protease inhibitor 17	777096 IMAGE Consortium	AA276326
6668 D	3 021BAMOC	ESTs, Highly similar to PUTATIVE ATP-DEPENDENT RNA HELICASE C22F3.08C [Schizosaccharomyces pombe]	779111 IMAGE Consortium	AA413830
6669 D	5 021BAMOC	phosphatidylinositol glycan, class A	779367 IMAGE Consortium	AA414112
6670 D	7 021BAMOC	ESTs	779593 IMAGE Consortium	AA413696
6671 D	9 021BAMOC	ESTs, Weakly similar to RO52_MOUSE 52 KD RO PROTEIN [M.musculus]	779821 IMAGE Consortium	AA414713
6672 D	11 021BAMOC	spermine synthase	791067 IMAGE Consortium	AA415202
6673 F	1 021BAMOC	ESTs, Moderately similar to SERINE/THREONINE PROTEIN PHOSPHATASE 2B CATALYTIC SUBUNIT, BETA ISOFORM [Mus musculus]	777167 IMAGE Consortium	AA276383
6674 F	3 021BAMOC	ESTs, Highly similar to p300/CBP/Co-Integrator protein [M.musculus]	779126 IMAGE Consortium	AA445710
6675 F	5 021BAMOC	signal recognition particle 9 kDa	779385 IMAGE Consortium	AA414116
6676 F	7 021BAMOC	ESTs	779608 IMAGE Consortium	AA466922
6677 F	9 021BAMOC	ESTs, Highly similar to HEXIM1 protein [H.sapiens]	779895 IMAGE Consortium	AA414822



6678 F	11 021BAMOC	ESTs	791078 IMAGE Consortium	AA415181
6679 H	1 021BAMOC	solute carrier family 13 (sodium/sulphate symporters), member 1	775737 IMAGE Consortium	AI508944
6680 H	3 021BAMOC	heat shock protein cognate 70, testis	779196 IMAGE Consortium	AA445731
6681 H	5 021BAMOC	ESTs, Highly similar to nucleoporin p54 [R.norvegicus]	779395 IMAGE Consortium	AA414836
6682 H	7 021BAMOC	poly (A) polymerase	779628 IMAGE Consortium	AA413822
6683 H	9 021BAMOC	cDNA sequence AF244542	779918 IMAGE Consortium	AA414828
6684 H	11 021BAMOC	muscle, skeletal, receptor tyrosine kinase	791095 IMAGE Consortium	AA415211
6685 B	1 021IAMOD	ESTs	791164 IMAGE Consortium	AA415273
6686 B	3 021IAMOD	nucleotide binding protein 2	790823 IMAGE Consortium	AA388597
6687 B	5 021IAMOD	ESTs	791849 IMAGE Consortium	AA414388
6688 B	7 021IAMOD	ESTs, Moderately similar to NOF1 [H.sapiens]	791944 IMAGE Consortium	AA414434
6689 B	9 021IAMOD	dystrophia myotonica linked gene, N9	792964 IMAGE Consortium	AA415346
6690 B	11 021IAMOD			
6691 D	1 021IAMOD	EST AI181014	791181 IMAGE Consortium	AA415500
6692 D	3 021IAMOD	hypothetical protein from clone MNCb-1932, similar to Homo sapiens FLJ20644	790920 IMAGE Consortium	AA415328
6693 D	5 021IAMOD	I3 protein	792597 IMAGE Consortium	AA415225
6694 D	7 021IAMOD	plexin 1	791967 IMAGE Consortium	AA414276
6695 D	9 021IAMOD	ESTs, Moderately similar to HYPOTHETICAL 104.7 KD PROTEIN IN PKC1-RTG3 INTERGENIC REGION [Saccharomyces cerevisiae]	792990 IMAGE Consortium	AA415473
6696 D	11 021IAMOD	karyopherin (importin) alpha 6	793177 IMAGE Consortium	AA438018
6697 F	1 021IAMOD	ESTs	791217 IMAGE Consortium	AA416264
6698 F	3 021IAMOD	cytidine monophospho-N-acetylneuraminic acid synthetase	790926 IMAGE Consortium	AA415331
6699 F	5 021IAMOD	ESTs, Highly similar to p68 RNA helicase [M.musculus]	791872 IMAGE Consortium	AA414411
6700 F	7 021IAMOD	ESTs, Highly similar to 40S RIBOSOMAL PROTEIN S25 [Homo sapiens; Rattus norvegicus]	792739 IMAGE Consortium	AA416439
6701 F	9 021IAMOD	ATPase, Na <sup>+</sup> /K <sup>+</sup> beta 3 polypeptide	792041 IMAGE Consortium	AA414588
6702 F	11 021IAMOD	ESTs, Moderately similar to hypothetical protein [H.sapiens]	791751 IMAGE Consortium	AA414681
6703 H	1 021IAMOD	nuclear RNA export factor 1 homolog (S. cerevisiae)	791266 IMAGE Consortium	AA416326
6704 H	3 021IAMOD	ESTs	790930 IMAGE Consortium	AA415333
6705 H	5 021IAMOD	ESTs, Highly similar to PHOSPHORYLASE B KINASE BETA REGULATORY CHAIN [Oryctolagus cuniculus]	792615 IMAGE Consortium	AA415285
6706 H	7 021IAMOD	ESTs	792846 IMAGE Consortium	AA416380
6707 H	9 021IAMOD	squalene epoxidase	793035 IMAGE Consortium	AA444230
6708 H	11 021IAMOD	ESTs	791776 IMAGE Consortium	AA414536
6709 B	1 021PAMOE	ESTs, Highly similar to DTDP-GLUCOSE 4,6-DEHYDRATASE [Escherichia coli]	791361 IMAGE Consortium	AA415408
6710 B	3 021PAMOE	ESTs	791558 IMAGE Consortium	AA437451
6711 B	5 021PAMOE	ESTs	793266 IMAGE Consortium	AA438095

6712 B	7 021PAMOE	neighbor of Cox4	792951 IMAGE Consortium	AA437616
6713 B	9 021PAMOE	stromal antigen 3	793979 IMAGE Consortium	AA422865
6714 B	11 021PAMOE	seven in absentia 2	793125 IMAGE Consortium	AA437881
6715 D	1 021PAMOE	open reading frame 60	791375 IMAGE Consortium	AA473975
6716 D	3 021PAMOE	ESTs, Highly similar to AF118890_1 s-tomosyn isoform [R.norvegicus]	791579 IMAGE Consortium	AA437465
6717 D	5 021PAMOE	ESTs	793367 IMAGE Consortium	AA438249
6718 D	7 021PAMOE	DNA segment, Chr 13, ERATO Doi 372, expressed	792974 IMAGE Consortium	AA437620
6719 D	9 021PAMOE	ESTs, Highly similar to dJ50O24.2.1 [H.sapiens]	793016 IMAGE Consortium	AA437631
6720 D	11 021PAMOE	ESTs, Weakly similar to ankyrin 3 [M.musculus]	793136 IMAGE Consortium	AA437976
6721 F	1 021PAMOE	Wolf-Hirschhorn syndrome candidate 2 homolog (human)	791395 IMAGE Consortium	AA473980
6722 F	3 021PAMOE	ESTs	791605 IMAGE Consortium	AA437577
6723 F	5 021PAMOE	pyruvate dehydrogenase complex, component X	793529 IMAGE Consortium	AA415447
6724 F	7 021PAMOE	ESTs, Moderately similar to T34531 hypothetical protein DKFZp434P1215.1 - human [H.sapiens]	792979 IMAGE Consortium	AA437717
6725 F	9 021PAMOE	DNA segment, Chr 7, ERATO Doi 143, expressed	793027 IMAGE Consortium	AA437728
6726 F	11 021PAMOE	mitogen-activated protein kinase 6	793145 IMAGE Consortium	AA437990
6727 H	1 021PAMOE	phenylalanine-tRNA synthetase-like	791413 IMAGE Consortium	AA416399
6728 H	3 021PAMOE	secretory carrier membrane protein 2	791611 IMAGE Consortium	AA474004
6729 H	5 021PAMOE	ESTs	792850 IMAGE Consortium	AA415534
6730 H	7 021PAMOE	geminin	792996 IMAGE Consortium	AA437625
6731 H	9 021PAMOE	ESTs, Highly similar to serine/threonine protein kinase TAO1 [R.norvegicus]	793043 IMAGE Consortium	AA444234
6732 H	11 021PAMOE	ESTs, Highly similar to protein inhibitor of nitric oxide synthase [M.musculus]	793166 IMAGE Consortium	AA438000
6733 B	1 021WAMOF	ESTs	352820 IMAGE Consortium	AI430693
6734 B	3 021WAMOF	ectonucleoside triphosphate diphosphohydrolase 3	354717 IMAGE Consortium	W46136
6735 B	5 021WAMOF	ESTs	351000 IMAGE Consortium	AI450521
6736 B	7 021WAMOF	ESTs	353005 IMAGE Consortium	AI430709
6737 B	9 021WAMOF	ESTs	793870 IMAGE Consortium	AA422606
6738 B	11 021WAMOF	ESTs	353609 IMAGE Consortium	W41732
6739 D	1 021WAMOF	ESTs, Weakly similar to CR16 [R.norvegicus]	354769 IMAGE Consortium	W45973
6740 D	3 021WAMOF	ESTs, Weakly similar to S12207 hypothetical protein [M.musculus]	350323 IMAGE Consortium	AI391176
6741 D	5 021WAMOF	ESTs, Weakly similar to AF191309_1 zinc finger protein [M.musculus]	350927 IMAGE Consortium	AI450518
6742 D	7 021WAMOF	ESTs, Weakly similar to sushi-repeat-containing protein [M.musculus]	806802 IMAGE Consortium	AA403815
6743 D	9 021WAMOF	desmoglein 2	808692 IMAGE Consortium	AA467296
6744 D	11 021WAMOF	ESTs	355113 IMAGE Consortium	W48255
6745 F	1 021WAMOF	ESTs	349757 IMAGE Consortium	W30094
6746 F	3 021WAMOF	ESTs	350349 IMAGE Consortium	W40747
6747 F	5 021WAMOF	ESTs		

6748 F	7 021WAMOF	ESTs	353167 IMAGE Consortium	AI430726
6749 F	9 021WAMOF	ESTs	807539 IMAGE Consortium	AI597424
6750 F	11 021WAMOF	eukaryotic translation initiation factor 1A	793988 IMAGE Consortium	AA422861
6751 H	1 021WAMOF	ring finger protein 22	353706 IMAGE Consortium	W41965
6752 H	3 021WAMOF	ESTs, Highly similar to dJ979N1.1 [H.sapiens]	349833 IMAGE Consortium	W30583
6753 H	5 021WAMOF	ESTs, Highly similar to COLLAGEN 1(X) CHAIN PRECURSOR [Bos taurus]	350489 IMAGE Consortium	W41045
6754 H	7 021WAMOF	SH3-domain GRB2-like B1 (endophilin)	351776 IMAGE Consortium	AI430540
6755 H	9 021WAMOF	ESTs	806894 IMAGE Consortium	AA403916
6756 H	11 021WAMOF	double C2, gamma	808846 IMAGE Consortium	AA467531
6757 B	1 0213AMOG	ESTs	804310 IMAGE Consortium	AA474841
6758 B	3 0213AMOG	Mus musculus PES1 protein (Pes1) mRNA, complete cds	803839 IMAGE Consortium	AA422979
6759 B	5 0213AMOG	ESTs	833286 IMAGE Consortium	AA465847
6760 B	7 0213AMOG	membrane interacting protein of RGS16	836062 IMAGE Consortium	AA426737
6761 B	9 0213AMOG	ESTs	835890 IMAGE Consortium	AA422198
6762 B	11 0213AMOG	ESTs	831943 IMAGE Consortium	AI549742
6763 D	1 0213AMOG	hypothetical protein MNCb-4193	803610 IMAGE Consortium	AA422756
6764 D	3 0213AMOG	ESTs	803870 IMAGE Consortium	AA427312
6765 D	5 0213AMOG	ESTs	832578 IMAGE Consortium	AI549815
6766 D	7 0213AMOG	ESTs	836085 IMAGE Consortium	AA426779
6767 D	9 0213AMOG	ESTs, Weakly similar to trg [R.norvegicus]	835931 IMAGE Consortium	AA422363
6768 D	11 0213AMOG	inositol 1,4,5-triphosphate receptor 1	831995 IMAGE Consortium	AA444527
6769 F	1 0213AMOG	ESTs, Moderately similar to unnamed protein product [H.sapiens]	803615 IMAGE Consortium	AA422788
6770 F	3 0213AMOG	ESTs	803887 IMAGE Consortium	AA435248
6771 F	5 0213AMOG	ESTs	832638 IMAGE Consortium	AI549822
6772 F	7 0213AMOG	ESTs	836100 IMAGE Consortium	AA426806
6773 F	9 0213AMOG	ESTs	832057 IMAGE Consortium	AA444862
6774 F	11 0213AMOG	ESTs	818271 IMAGE Consortium	AI596286
6775 H	1 0213AMOG	Mus musculus mRNA, zygotic activation gene inactivated by frameshift	803639 IMAGE Consortium	AA422810
6776 H	3 0213AMOG	ESTs	803936 IMAGE Consortium	AA435317
6777 H	5 0213AMOG	zinc finger protein 42	833416 IMAGE Consortium	AA432569
6778 H	7 0213AMOG	ESTs, Weakly similar to almondex [D.melanogaster]	835496 IMAGE Consortium	AA422438
6779 H	9 0213AMOG	ESTs	831646 IMAGE Consortium	AI549699
6780 H	11 0213AMOG	ESTs	818384 IMAGE Consortium	AI596291
6781 B	1 021AAMOH	AU RNA-binding protein/enoyl-coenzyme A hydratase	818743 IMAGE Consortium	AA466669
6782 B	3 021AAMOH	ESTs	820899 IMAGE Consortium	AA646765
6783 B	5 021AAMOH	Mus musculus brain cDNA, clone MNCb-3307	836442 IMAGE Consortium	AA426904
6784 B	7 021AAMOH	ESTs, Highly similar to TB1 [H.sapiens]	846452 IMAGE Consortium	AI553049
6785 B	9 021AAMOH			
6786 B	11 021AAMOH			
6787 D	1 021AAMOH	ATP-binding cassette, sub-family B (MDR/TAP), member	818939 IMAGE Consortium	AA434922

6788 D	3 021AAMOH	ESTs	849415 IMAGE Consortium	AI552264
6789 D	5 021AAMOH	ESTs	836482 IMAGE Consortium	AA426941
6790 D	7 021AAMOH	ESTs	847196 IMAGE Consortium	AA420047
6791 D	9 021AAMOH	ESTs, Weakly similar to LCFB_MOUSE LONG-CHAIN-FATTY-ACID--COA LIGASE 2 [M.musculus]	846790 IMAGE Consortium	AA422420
6792 D	11 021AAMOH	ESTs	847819 IMAGE Consortium	AA433804
6793 F	1 021AAMOH	ESTs	820180 IMAGE Consortium	AA423190
6794 F	3 021AAMOH	ESTs, Weakly similar to F31D4.2 [C.elegans]	849432 IMAGE Consortium	AA457882
6795 F	5 021AAMOH	ESTs	846033 IMAGE Consortium	AI553027
6796 F	7 021AAMOH	ESTs, Highly similar to lymphocyte specific adaptor protein Lnk [M.musculus]	847231 IMAGE Consortium	AI553101
6797 F	9 021AAMOH	adrenergic receptor, alpha 1b	846842 IMAGE Consortium	AA422482
6798 F	11 021AAMOH	ESTs	848654 IMAGE Consortium	AI552209
6799 H	1 021AAMOH	ESTs	820236 IMAGE Consortium	AA423177
6800 H	3 021AAMOH	FK506 binding protein 7 (23 kDa)	849512 IMAGE Consortium	AA457921
6801 H	5 021AAMOH	ESTs	846290 IMAGE Consortium	AI553038
6802 H	7 021AAMOH			
6803 H	9 021AAMOH	EST AA450473	849038 IMAGE Consortium	AA470188
6804 H	11 021AAMOH	ESTs	847936 IMAGE Consortium	AA469493
6805 B	1 021HAMOI	ESTs	849519 IMAGE Consortium	AA457932
6806 B	3 021HAMOI	casein kappa	850292 IMAGE Consortium	AA451500
6807 B	5 021HAMOI	ESTs, Weakly similar to REQU_MOUSE ZINC-FINGER PROTEIN UBI-D4 [M.musculus]	851370 IMAGE Consortium	AA461681
6808 B	7 021HAMOI	ESTs	860444 IMAGE Consortium	AI613612
6809 B	9 021HAMOI	ESTs	875051 IMAGE Consortium	AA472125
6810 B	11 021HAMOI			
6811 D	1 021HAMOI	solute carrier family 7 (cationic amino acid, transporter, y+ system), member 10	849597 IMAGE Consortium	AI552288
6812 D	3 021HAMOI	ESTs, Weakly similar to Z151_MOUSE ZINC FINGER PROTEIN 151 [M.musculus]	850329 IMAGE Consortium	AA451536
6813 D	5 021HAMOI	ESTs	851390 IMAGE Consortium	AA461870
6814 D	7 021HAMOI	ESTs	860467 IMAGE Consortium	AA499747
6815 D	9 021HAMOI	ESTs	873662 IMAGE Consortium	AA471907
6816 D	11 021HAMOI	ESTs, Weakly similar to I49278 t complex testis-specific protein - mouse [M.musculus]	875386 IMAGE Consortium	AA475555
6817 F	1 021HAMOI	ESTs, Weakly similar to CG13189 gene product [D.melanogaster]	849799 IMAGE Consortium	AI608331
6818 F	3 021HAMOI	ESTs	850359 IMAGE Consortium	AA451219
6819 F	5 021HAMOI	ESTs	860168 IMAGE Consortium	AA500847
6820 F	7 021HAMOI	phospholipase C-like protein	851855 IMAGE Consortium	AA522203
6821 F	9 021HAMOI	ESTs	875216 IMAGE Consortium	AA472692
6822 F	11 021HAMOI	transient receptor potential-related protein	873970 IMAGE Consortium	AA472153

6823 H	1 021HAMOI	ESTs	849840 IMAGE Consortium	AA450488
6824 H	3 021HAMOI	ESTs	850451 IMAGE Consortium	AI480957
6825 H	5 021HAMOI	ESTs	860223 IMAGE Consortium	AA500978
6826 H	7 021HAMOI	muscle-specific serine kinase 1	860600 IMAGE Consortium	AA500064
6827 H	9 021HAMOI	CCR4-NOT transcription complex, subunit 4	873807 IMAGE Consortium	AA471811
6828 H	11 021HAMOI	period homolog 2 (Drosophila)	876194 IMAGE Consortium	AA475629
6829 B	1 021OAMOJ	ESTs	875859 IMAGE Consortium	AA575209
6830 B	3 021OAMOJ	ectonucleoside triphosphate diphosphohydrolase 2	864751 IMAGE Consortium	AA444620
6831 B	5 021OAMOJ	mannose receptor, C type 1	870901 IMAGE Consortium	AA462138
6832 B	7 021OAMOJ	ESTs	890703 IMAGE Consortium	AA510002
6833 B	9 021OAMOJ	ESTs	876471 IMAGE Consortium	AA475934
6834 B	11 021OAMOJ	ESTs	889538 IMAGE Consortium	AA498703
6835 D	1 021OAMOJ	M.musculus mRNA (1F6) for IgA V-D-J-heavy chain	875880 IMAGE Consortium	AA575218
6836 D	3 021OAMOJ	flotillin 2	864783 IMAGE Consortium	AA444808
6837 D	5 021OAMOJ	EST	864432 IMAGE Consortium	AI482499
6838 D	7 021OAMOJ	ESTs	876180 IMAGE Consortium	AA475611
6839 D	9 021OAMOJ	ESTs	876575 IMAGE Consortium	AA476020
6840 D	11 021OAMOJ	sphingosine phosphate lyase 1	902269 IMAGE Consortium	AA516927
6841 F	1 021OAMOJ	EST, Moderately similar to alias DLC1 [H.sapiens]	872087 IMAGE Consortium	AI481126
6842 F	3 021OAMOJ	ESTs, Weakly similar to Z177_HUMAN ZINC FINGER PROTEIN 177 [H.sapiens]	864827 IMAGE Consortium	AA445023
6843 F	5 021OAMOJ	S100 calcium binding protein A8 (calgranulin A)	864657 IMAGE Consortium	AA458347
6844 F	7 021OAMOJ	ESTs	890791 IMAGE Consortium	AA510512
6845 F	9 021OAMOJ	ESTs, Moderately similar to KIAA0697 protein [H.sapiens]	876720 IMAGE Consortium	AA497520
6846 F	11 021OAMOJ	ESTs	889875 IMAGE Consortium	AA498588
6847 H	1 021OAMOJ	ESTs	871508 IMAGE Consortium	AA462085
6848 H	3 021OAMOJ	ESTs	871034 IMAGE Consortium	AI552895
6849 H	5 021OAMOJ	ESTs, Weakly similar to A27353 collagen alpha 1(III) chain precursor - mouse [M.musculus]	891116 IMAGE Consortium	AI614496
6850 H	7 021OAMOJ	ESTs, Weakly similar to BAF53a [M.musculus]	890802 IMAGE Consortium	AA510507
6851 H	9 021OAMOJ	ESTs	888463 IMAGE Consortium	AI481700
6852 H	11 021OAMOJ	ESTs	904229 IMAGE Consortium	AI553220
6853 B	1 021VAMOK	ESTs	905129 IMAGE Consortium	AI553293
6854 B	3 021VAMOK	ESTs	905435 IMAGE Consortium	AA529139
6855 B	5 021VAMOK	ESTs	902011 IMAGE Consortium	AA517891
6856 B	7 021VAMOK	ESTs	902701 IMAGE Consortium	AI553118
6857 B	9 021VAMOK	ESTs	948924 IMAGE Consortium	AI596699
6858 B	11 021VAMOK	ESTs, Weakly similar to brain specific cortactin-binding protein CBP90 [R.norvegicus]	959695 IMAGE Consortium	AA546842
6859 D	1 021VAMOK	ESTs	905189 IMAGE Consortium	AI553296
6860 D	3 021VAMOK	ESTs	905493 IMAGE Consortium	AI553308
6861 D	5 021VAMOK	steroidogenic acute regulatory protein	903163 IMAGE Consortium	AA521869
6862 D	7 021VAMOK	ESTs	904172 IMAGE Consortium	AI553217

6863 D	9 021VAMOK	ESTs	949671 IMAGE Consortium	AA537359
6864 D	11 021VAMOK	ESTs	959728 IMAGE Consortium	AA546857
6865 F	1 021VAMOK	ESTs	903738 IMAGE Consortium	AI553193
6866 F	3 021VAMOK	ESTs	904858 IMAGE Consortium	AI553263
6867 F	5 021VAMOK	SH3 domain protein D19	903241 IMAGE Consortium	AA518977
6868 F	7 021VAMOK	ESTs	903455 IMAGE Consortium	AA575288
6869 F	9 021VAMOK	ESTs	959501 IMAGE Consortium	AA546474
6870 F	11 021VAMOK	ESTs, Highly similar to KIAA0160 gene product is novel. [H.sapiens]	959755 IMAGE Consortium	AA546919
6871 H	1 021VAMOK	ESTs	904496 IMAGE Consortium	AA617112
6872 H	3 021VAMOK	ESTs	904883 IMAGE Consortium	AA519023
6873 H	5 021VAMOK	a disintegrin-like and metalloprotease (reprolysin type) with thrombospondin type 1 motif, 5 (aggrecanase-2)	902598 IMAGE Consortium	AI553108
6874 H	7 021VAMOK	ESTs, Highly similar to AF132966_1 CGI-32 protein [H.sapiens]	948575 IMAGE Consortium	AA544717
6875 H	9 021VAMOK	ESTs	959543 IMAGE Consortium	AA546500
6876 H	11 021VAMOK	ESTs	959771 IMAGE Consortium	AA546934
6877 B	1 0212AMOL	Mus musculus nuclear body associated kinase 1a (Nbak1) mRNA, alternatively spliced, complete cds	959952 IMAGE Consortium	AA547452
6878 B	3 0212AMOL	mitogen activated protein kinase 13	368273 IMAGE Consortium	W53837
6879 B	5 0212AMOL	frizzled homolog 8 (Drosophila)	368781 IMAGE Consortium	W56988
6880 B	7 0212AMOL	ESTs, Highly similar to S42924 ZG-16p protein - rat [R.norvegicus]	921936 IMAGE Consortium	AI550886
6881 B	9 0212AMOL	ESTs	944404 IMAGE Consortium	AA545578
6882 B	11 0212AMOL	proteasome (prosome, macropain) 26S subunit, ATPase 3, interacting protein	944886 IMAGE Consortium	AA545416
6883 D	1 0212AMOL	ESTs, Highly similar to KIAA0368 [H.sapiens]	960032 IMAGE Consortium	AA547260
6884 D	3 0212AMOL	trinucleotide repeat containing 11 (THR-associated protein, 230 kDa subunit)	368355 IMAGE Consortium	AI893925
6885 D	5 0212AMOL	ESTs	368882 IMAGE Consortium	W53241
6886 D	7 0212AMOL	early B-cell factor	935477 IMAGE Consortium	AA543172
6887 D	9 0212AMOL	kit oncogene	944489 IMAGE Consortium	AA545105
6888 D	11 0212AMOL	ESTs	960947 IMAGE Consortium	AA549547
6889 F	1 0212AMOL	ESTs	960124 IMAGE Consortium	AA547371
6890 F	3 0212AMOL	interleukin 11 receptor, alpha chain 1	366999 IMAGE Consortium	W54301
6891 F	5 0212AMOL	ESTs	921488 IMAGE Consortium	AA529770
6892 F	7 0212AMOL	ESTs	935539 IMAGE Consortium	AA543485
6893 F	9 0212AMOL	Mouse mRNA for cytotoxic T-cell membrane glycoprotein Ly-3 3'flank	944532 IMAGE Consortium	AA545114
6894 F	11 0212AMOL	ESTs	960956 IMAGE Consortium	AA549540
6895 H	1 0212AMOL	chromogranin B	367410 IMAGE Consortium	W54256
6896 H	3 0212AMOL	nuclear factor of activated T-cells, cytoplasmic 3	368464 IMAGE Consortium	W53705
6897 H	5 0212AMOL	UBIQUITIN-LIKE PROTEIN GDX	922316 IMAGE Consortium	AI550986

6898 H	7 0212AMOL			
6899 H	9 0212AMOL	selenocysteine tRNA gene transcription activating factor	944729 IMAGE Consortium	AA545291
6900 H	11 0212AMOL	ESTs	959903 IMAGE Consortium	AA547122
6901 B	1 0219AMOM	chromobox homolog 5 (Drosophila HP1a)	960068 IMAGE Consortium	AA547482
6902 B	3 0219AMOM	Purkinje cell protein 1	961035 IMAGE Consortium	AA547170
6903 B	5 0219AMOM	Mus musculus sentrin/SUMO-specific protease (SEN3) mRNA, complete cds	960600 IMAGE Consortium	AA549588
6904 B	7 0219AMOM	mitogen activated protein kinase 14	961617 IMAGE Consortium	AA549577
6905 B	9 0219AMOM	ESTs	961990 IMAGE Consortium	AA672330
6906 B	11 0219AMOM	Public domain EST	961494 IMAGE Consortium	
6907 D	1 0219AMOM	ESTs	960106 IMAGE Consortium	AA547353
6908 D	3 0219AMOM	ESTs	961136 IMAGE Consortium	AA547393
6909 D	5 0219AMOM	ESTs, Highly similar to CRM1 [H.sapiens]	960606 IMAGE Consortium	AA549591
6910 D	7 0219AMOM	ESTs	961705 IMAGE Consortium	AA672128
6911 D	9 0219AMOM	ESTs	962963 IMAGE Consortium	AA671824
6912 D	11 0219AMOM	polypyrimidine tract binding protein	961535 IMAGE Consortium	AA549391
6913 F	1 0219AMOM	DNA segment, Chr 1, ERATO Doi 654, expressed	960178 IMAGE Consortium	AA547574
6914 F	3 0219AMOM	ESTs, Weakly similar to ubiquitin-conjugating enzyme UbcM2 [M.musculus]	960434 IMAGE Consortium	AA549225
6915 F	5 0219AMOM	surfeit gene 5	960640 IMAGE Consortium	AA549617
6916 F	7 0219AMOM	chemokine (C-C) receptor 5	963216 IMAGE Consortium	AA671573
6917 F	9 0219AMOM	interferon dependent positive acting transcription factor 3 gamma	961095 IMAGE Consortium	AA547214
6918 F	11 0219AMOM	ESTs, Weakly similar to Phosphatidic acid phosphatase [M.musculus]	961560 IMAGE Consortium	AA549462
6919 H	1 0219AMOM	ESTs, Weakly similar to I49636 DNA-binding protein - mouse [M.musculus]	960202 IMAGE Consortium	AA547593
6920 H	3 0219AMOM	ESTs	961184 IMAGE Consortium	AA547433
6921 H	5 0219AMOM	ESTs	960715 IMAGE Consortium	AA549037
6922 H	7 0219AMOM	ESTs	961856 IMAGE Consortium	AA553240
6923 H	9 0219AMOM	ESTs	961142 IMAGE Consortium	AA547395
6924 H	11 0219AMOM	ESTs, Highly similar to EUKARYOTIC INITIATION FACTOR 4 GAMMA [Homo sapiens]	1003011 IMAGE Consortium	AA684461
6925 B	1 021GAMON	Cd27 binding protein (Hindu God of destruction)	1003064 IMAGE Consortium	AA684490
6926 B	3 021GAMON	ESTs, Moderately similar to KIAA0423 [H.sapiens]	1003381 IMAGE Consortium	AA684172
6927 B	5 021GAMON	ESTs, Weakly similar to The KIAA0191 gene is expressed ubiquitously. [H.sapiens]	1003498 IMAGE Consortium	AA684341
6928 B	7 021GAMON	TATA box binding protein (TBP) associated factor, RNA polymerase II, F, 55 kDa	976417 IMAGE Consortium	AA619797
6929 B	9 021GAMON	ESTs, Weakly similar to Similarity to Human ADP/ATP carrier protein [C.elegans]	976495 IMAGE Consortium	AA619953
6930 B	11 021GAMON	proteasome (prosome, macropain) 26S subunit, ATPase, 4	976640 IMAGE Consortium	AA619748

6931 D	1	021GAMON	solute carrier family 4 (anion exchanger), member 8	1003097	IMAGE Consortium	AA684011
6932 D	3	021GAMON	CDC-like kinase	1003405	IMAGE Consortium	AA684191
6933 D	5	021GAMON	ESTs	1003520	IMAGE Consortium	AA684403
6934 D	7	021GAMON	TATA box binding protein (TBP)-associated factor, RNA polymerase II, E	976425	IMAGE Consortium	AA619800
6935 D	9	021GAMON	ESTs, Moderately similar to GPDA_MOUSE GLYCEROL-3-PHOSPHATE DEHYDROGENASE [M.musculus]	976513	IMAGE Consortium	AA619970
6936 D	11	021GAMON	B lymphoma Mo-MLV insertion region 1	976651	IMAGE Consortium	AA619905
6937 F	1	021GAMON	ESTs, Highly similar to C214_HUMAN 17.9 KDA MEMBRANE PROTEIN C21ORF4 [H.sapiens]	1003203	IMAGE Consortium	AA684097
6938 F	3	021GAMON	TRAF-interacting protein	1003411	IMAGE Consortium	AA684194
6939 F	5	021GAMON	ESTs	1003537	IMAGE Consortium	AA684428
6940 F	7	021GAMON	DNA segment, Chr 3, ERATO Doi 265, expressed	976436	IMAGE Consortium	AA619795
6941 F	9	021GAMON	ESTs, Highly similar to A56465 TFIIIC2 subunit - human [H.sapiens]	976521	IMAGE Consortium	AA619974
6942 F	11	021GAMON	chaperonin subunit 3 (gamma)	976658	IMAGE Consortium	AA619757
6943 H	1	021GAMON	ESTs	1003214	IMAGE Consortium	AA684092
6944 H	3	021GAMON	spermatogenesis associated factor	1003435	IMAGE Consortium	AA684252
6945 H	5	021GAMON	ESTs	1003556	IMAGE Consortium	AA684426
6946 H	7	021GAMON	septin 7	976450	IMAGE Consortium	AA619856
6947 H	9	021GAMON	glutathione transferase zeta 1 (maleylacetoacetate isomerase)	976523	IMAGE Consortium	AA619975
6948 H	11	021GAMON	ornithine decarboxylase antizyme	976667	IMAGE Consortium	AA619767
6949 B	1	021NAMOO	RAB10, member RAS oncogene family	976707	IMAGE Consortium	AA619822
6950 B	3	021NAMOO	ESTs	1025528	IMAGE Consortium	AA590599
6951 B	5	021NAMOO	ESTs, Highly similar to AF181856_1 tRNA selenocysteine associated protein [R.norvegicus]	1003856	IMAGE Consortium	AA674491
6952 B	7	021NAMOO	ESTs	1003996	IMAGE Consortium	AA674771
6953 B	9	021NAMOO	ESTs	1051101	IMAGE Consortium	AA592360
6954 B	11	021NAMOO	ESTs	1052108	IMAGE Consortium	AA592861
6955 D	1	021NAMOO	T-cell lymphoma invasion and metastasis 1	976719	IMAGE Consortium	AA619827
6956 D	3	021NAMOO	solute carrier family 6 (neurotransmitter transporter, serotonin), member 4	1025582	IMAGE Consortium	AA590845
6957 D	5	021NAMOO	ESTs, Weakly similar to S1-1 protein [R.norvegicus]	1003872	IMAGE Consortium	AA674506
6958 D	7	021NAMOO	ESTs	1049539	IMAGE Consortium	AI594352
6959 D	9	021NAMOO	ESTs	1051167	IMAGE Consortium	AA597125
6960 D	11	021NAMOO	ESTs, Weakly similar to LCFB_MOUSE LONG-CHAIN-FATTY-ACID--COA LIGASE 2 [M.musculus]	1052193	IMAGE Consortium	AA596254
6961 F	1	021NAMOO	ESTs, Highly similar to similar to human DNA-binding protein 5. [H.sapiens]	976728	IMAGE Consortium	AA619828
6962 F	3	021NAMOO	dynamamin	1025651	IMAGE Consortium	AI647249
6963 F	5	021NAMOO	GA repeat binding protein, beta 1	1003892	IMAGE Consortium	AA674510
6964 F	7	021NAMOO	cytochrome P450, 51	1025252	IMAGE Consortium	AA590222



6965 F	9 021NAMOO	ESTs	1051213 IMAGE Consortium	AI429015
6966 F	11 021NAMOO	ESTs	1052214 IMAGE Consortium	AI594457
6967 H	1 021NAMOO	ryanodine receptor 1, skeletal muscle	976734 IMAGE Consortium	AA619831
6968 H	3 021NAMOO	ESTs, Highly similar to GC-RICH SEQUENCE DNA-BINDING FACTOR [Homo sapiens]	1025685 IMAGE Consortium	AA608265
6969 H	5 021NAMOO	Mus musculus CNP-inducible osteoblast mRNA sequence	1003902 IMAGE Consortium	AA674566
6970 H	7 021NAMOO	ESTs	1026037 IMAGE Consortium	AA616343
6971 H	9 021NAMOO	ESTs	1051242 IMAGE Consortium	AI429017
6972 H	11 021NAMOO	ESTs	1052344 IMAGE Consortium	AA839289
6973 B	1 021UAMOP	ESTs, Weakly similar to EL2_MOUSE ELASTASE 2 PRECURSOR [M.musculus]	1053193 IMAGE Consortium	AA607713
6974 B	3 021UAMOP	ESTs	1053526 IMAGE Consortium	AA608466
6975 B	5 021UAMOP	ESTs	1068534 IMAGE Consortium	AI585465
6976 B	7 021UAMOP	ESTs	1079007 IMAGE Consortium	AA734892
6977 B	9 021UAMOP	ESTs	1068011 IMAGE Consortium	AA764383
6978 B	11 021UAMOP	ESTs	1054943 IMAGE Consortium	AI603969
6979 D	1 021UAMOP	ESTs	1053206 IMAGE Consortium	AI594015
6980 D	3 021UAMOP	ESTs	1052904 IMAGE Consortium	AI593995
6981 D	5 021UAMOP	ADP-ribosyltransferase 2a	1067825 IMAGE Consortium	AA763726
6982 D	7 021UAMOP	ESTs, Weakly similar to testis specific DNAj-homolog [M.musculus]	1079076 IMAGE Consortium	AI593610
6983 D	9 021UAMOP	SRY-box containing gene 6	1068053 IMAGE Consortium	AI536532
6984 D	11 021UAMOP	glypican 1	1065141 IMAGE Consortium	AA611085
6985 F	1 021UAMOP	ESTs	1051856 IMAGE Consortium	AI594426
6986 F	3 021UAMOP	ESTs	1053688 IMAGE Consortium	AI594090
6987 F	5 021UAMOP	ESTs	1078030 IMAGE Consortium	AA792956
6988 F	7 021UAMOP	cannabinoid receptor 2 (macrophage)	1068955 IMAGE Consortium	AA796154
6989 F	9 021UAMOP	ESTs	1068061 IMAGE Consortium	AA764465
6990 F	11 021UAMOP	ESTs	1065166 IMAGE Consortium	AI594067
6991 H	1 021UAMOP	ESTs	1051908 IMAGE Consortium	AA597135
6992 H	3 021UAMOP	ESTs, Moderately similar to protocadherin-3 [R.norvegicus]	1052992 IMAGE Consortium	AA606540
6993 H	5 021UAMOP	ESTs	1068654 IMAGE Consortium	AA796683
6994 H	7 021UAMOP	ESTs, Weakly similar to Keap1 [M.musculus]	1077786 IMAGE Consortium	AA796458
6995 H	9 021UAMOP	ESTs	1068127 IMAGE Consortium	AA763658
6996 H	11 021UAMOP	ESTs, Highly similar to hypothetical protein [H.sapiens]	1066067 IMAGE Consortium	AA673972
6997 B	1 0211AMOQ	myomesin 2	1064754 IMAGE Consortium	AI594057
6998 B	3 0211AMOQ	ESTs	1066557 IMAGE Consortium	AA616077
6999 B	5 0211AMOQ	ESTs, Weakly similar to polypeptide GalNAc transferase-T1 [M.musculus]	1066733 IMAGE Consortium	AA611977
7000 B	7 0211AMOQ	ESTs	1079162 IMAGE Consortium	AA823472
7001 B	9 0211AMOQ			
7002 B	11 0211AMOQ	intelectin	1097545 IMAGE Consortium	AA869219

7003 D	1	0211AMOQ	ESTs	1066291	IMAGE Consortium	AI594378
7004 D	3	0211AMOQ	ESTs, Moderately similar to match to AA908753 [H.sapiens]	1065847	IMAGE Consortium	AA619540
7005 D	5	0211AMOQ	ESTs, Highly similar to angiotensin II AT2 receptor-interacting protein [M.musculus]	1066907	IMAGE Consortium	AA616052
7006 D	7	0211AMOQ	ESTs	1079180	IMAGE Consortium	AA823478
7007 D	9	0211AMOQ	defensin related cryptdin 5	1096212	IMAGE Consortium	AA871410
7008 D	11	0211AMOQ	Mus musculus 129 cryptdin 6 gene, complete cds	1096942	IMAGE Consortium	AA871914
7009 F	1	0211AMOQ	ESTs	1065584	IMAGE Consortium	AI594620
7010 F	3	0211AMOQ	Mest-linked imprinted transcript 1	1065892	IMAGE Consortium	AA611551
7011 F	5	0211AMOQ	mitogen regulated protein, proliferin 3	1094143	IMAGE Consortium	AA870247
7012 F	7	0211AMOQ				
7013 F	9	0211AMOQ	ESTs	1097079	IMAGE Consortium	AI641890
7014 F	11	0211AMOQ				
7015 H	1	0211AMOQ	ESTs	1067050	IMAGE Consortium	AA764290
7016 H	3	0211AMOQ	ESTs	1065901	IMAGE Consortium	AA611566
7017 H	5	0211AMOQ	ESTs	1094321	IMAGE Consortium	AI592912
7018 H	7	0211AMOQ	ESTs	1079256	IMAGE Consortium	AI593633
7019 H	9	0211AMOQ	ESTs, Weakly similar to membrane glycoprotein [M.musculus]	1095780	IMAGE Consortium	AA869082
7020 H	11	0211AMOQ	ESTs, Highly similar to P11D_MOUSE PHOSPHATIDYLINOSITOL 3-KINASE CATALYTIC SUBUNIT, DELTA ISOFORM [M.musculus]	400170	IMAGE Consortium	W77584
7021 B	1	0218AMOR	ESTs	389345	IMAGE Consortium	W71072
7022 B	3	0218AMOR	antigen identified by monoclonal antibody MRC OX-2	400842	IMAGE Consortium	AI893281
7023 B	5	0218AMOR	Rous sarcoma oncogene	402966	IMAGE Consortium	AI425806
7024 B	7	0218AMOR	ESTs, Weakly similar to COPP_RAT COATOMER BETA' SUBUNIT [R.norvegicus]	1107446	IMAGE Consortium	AA623616
7025 B	9	0218AMOR	ESTs	1107871	IMAGE Consortium	AA623716
7026 B	11	0218AMOR	ESTs	1125027	IMAGE Consortium	AA692976
7027 D	1	0218AMOR	ESTs	401315	IMAGE Consortium	AI425733
7028 D	3	0218AMOR	ESTs	402303	IMAGE Consortium	AI425772
7029 D	5	0218AMOR	ESTs, Weakly similar to transcriptional repressor Mpc2 [M.musculus]	391119	IMAGE Consortium	W75009
7030 D	7	0218AMOR	topoisomerase (DNA) I	1107479	IMAGE Consortium	AA623645
7031 D	9	0218AMOR	ESTs, Highly similar to unnamed protein product [H.sapiens]	1107943	IMAGE Consortium	AA623811
7032 D	11	0218AMOR	ESTs, Weakly similar to cell division cycle 37 homolog [M.musculus]	1125099	IMAGE Consortium	AA693061
7033 F	1	0218AMOR	enabled homolog (Drosophila)	400635	IMAGE Consortium	AI893955
7034 F	3	0218AMOR	G protein-coupled receptor 27	402354	IMAGE Consortium	W78348
7035 F	5	0218AMOR	ESTs	390464	IMAGE Consortium	AI430183
7036 F	7	0218AMOR	ESTs	1107551	IMAGE Consortium	AA623758

7037 F	9 0218AMOR	defensin related cryptdin, related sequence 2	1097493 IMAGE Consortium	AA869173
7038 F	11 0218AMOR	ESTs	1125248 IMAGE Consortium	AA693191
7039 H	1 0218AMOR	ESTs	401390 IMAGE Consortium	AI425735
7040 H	3 0218AMOR			
7041 H	5 0218AMOR	ESTs	391237 IMAGE Consortium	AI430204
7042 H	7 0218AMOR	yes-associated protein, 65 kDa	1107584 IMAGE Consortium	AA623823
7043 H	9 0218AMOR	seb4 mRNA	1108042 IMAGE Consortium	AA623909
7044 H	11 0218AMOR	ESTs	1125295 IMAGE Consortium	AA675566
7045 B	1 021FAMOS	ariadne homolog 2 (Drosophila)	1125452 IMAGE Consortium	AA675160
7046 B	3 021FAMOS	ESTs, Weakly similar to BETA-ADAPTIN [Homo sapiens; Rattus norvegicus; Bos taurus]	1134329 IMAGE Consortium	AA675608
7047 B	5 021FAMOS	DNA segment, Chr 5, ERATO Doi 708, expressed	1125878 IMAGE Consortium	AA823639
7048 B	7 021FAMOS	ESTs	1137181 IMAGE Consortium	AA689758
7049 B	9 021FAMOS	ESTs	1138151 IMAGE Consortium	AA691755
7050 B	11 021FAMOS	ESTs, Weakly similar to Inv [M.musculus]	1138536 IMAGE Consortium	AA688667
7051 D	1 021FAMOS	torsin family 1, member B	1125498 IMAGE Consortium	AA675204
7052 D	3 021FAMOS	ESTs, Weakly similar to activator protein 4 [D.melanogaster]	1125645 IMAGE Consortium	AA824156
7053 D	5 021FAMOS	UDP-N-acetyl-alpha-D-galactosamine:(N-acetylneuraminyl)-galactosyl-N-acetylglucosaminylpolypeptide-beta-1, 4-N-acetylgalactosaminyltransferase	1125908 IMAGE Consortium	AA823657
7054 D	7 021FAMOS	ESTs, Highly similar to UDP-GalNAc:polypeptide N-acetylgalactosaminyltransferase T6 [R.norvegicus]	1137254 IMAGE Consortium	AA688927
7055 D	9 021FAMOS	ESTs	1152842 IMAGE Consortium	AA644819
7056 D	11 021FAMOS	distal intestinal serine protease	1138561 IMAGE Consortium	AA688697
7057 F	1 021FAMOS	solute carrier family 29 (nucleoside transporters), member 2	1134223 IMAGE Consortium	AA675457
7058 F	3 021FAMOS	purine rich element binding protein B	1134418 IMAGE Consortium	AA681235
7059 F	5 021FAMOS	ESTs	1134936 IMAGE Consortium	AA690121
7060 F	7 021FAMOS	ESTs	1138048 IMAGE Consortium	AA691589
7061 F	9 021FAMOS			
7062 F	11 021FAMOS	Kruppel-like factor 3 (basic)	1138575 IMAGE Consortium	AA688702
7063 H	1 021FAMOS	U1 small nuclear ribonucleoprotein 70 kDa polypeptide A	1125507 IMAGE Consortium	AA675224
7064 H	3 021FAMOS	ESTs	1134457 IMAGE Consortium	AA681098
7065 H	5 021FAMOS	ESTs	1136124 IMAGE Consortium	AA762191
7066 H	7 021FAMOS	ESTs	1138058 IMAGE Consortium	AA691635
7067 H	9 021FAMOS	ESTs	1136820 IMAGE Consortium	AA688795
7068 H	11 021FAMOS	ESTs	1137887 IMAGE Consortium	AA691358
7069 B	1 021MAMOT	ESTs	1138961 IMAGE Consortium	AI536395
7070 B	3 021MAMOT	ESTs	1166694 IMAGE Consortium	AA710841
7071 B	5 021MAMOT	ESTs, Weakly similar to zinc finger protein 95 [M.musculus]	1163141 IMAGE Consortium	AA771229

7072 B	7 021MAMOT	ESTs	1163478 IMAGE Consortium	AI587721
7073 B	9 021MAMOT	ESTs	1182048 IMAGE Consortium	AA711962
7074 B	11 021MAMOT	ESTs	1195301 IMAGE Consortium	AA754903
7075 D	1 021MAMOT	intracisternal A particles	1166033 IMAGE Consortium	AA710164
7076 D	3 021MAMOT	ESTs	1166739 IMAGE Consortium	AA711008
7077 D	5 021MAMOT	homeo box A7	1162432 IMAGE Consortium	AA692071
7078 D	7 021MAMOT	myosin-binding protein H	1162842 IMAGE Consortium	AA770788
7079 D	9 021MAMOT	POU domain, class 2, transcription factor 1	1181696 IMAGE Consortium	AA711419
7080 D	11 021MAMOT	ESTs	1195316 IMAGE Consortium	AA754899
7081 F	1 021MAMOT	neuronal tyrosine/threonine phosphatase 1	1165334 IMAGE Consortium	AA709718
7082 F	3 021MAMOT	ESTs	1166823 IMAGE Consortium	AA710415
7083 F	5 021MAMOT	sialyltransferase 5	1153834 IMAGE Consortium	AA667986
7084 F	7 021MAMOT	myogenin	1162866 IMAGE Consortium	AA770807
7085 F	9 021MAMOT	ESTs	1181879 IMAGE Consortium	AA711674
7086 F	11 021MAMOT	vomer nasal organ family 1, receptor 2	1195366 IMAGE Consortium	AA754983
7087 H	1 021MAMOT	ESTs	1166118 IMAGE Consortium	AA710387
7088 H	3 021MAMOT	ESTs	1162011 IMAGE Consortium	AA815654
7089 H	5 021MAMOT	zinc-iron transporter-like	1164050 IMAGE Consortium	AA690621
7090 H	7 021MAMOT	Tax interaction protein 40	1162907 IMAGE Consortium	AA770881
7091 H	9 021MAMOT	nuclear factor I/A	1195030 IMAGE Consortium	AA690418
7092 H	11 021MAMOT	phosphodiesterase 1B1, Ca <sup>2+</sup> -calmodulin dependent, 63 kDa	1195462 IMAGE Consortium	AA756678
7093 B	1 021TAMOU	ESTs	1194992 IMAGE Consortium	AA690387
7094 B	3 021TAMOU	ESTs	387797 IMAGE Consortium	AI430644
7095 B	5 021TAMOU	ESTs	373748 IMAGE Consortium	AI430081
7096 B	7 021TAMOU	aplysia ras-related homolog N (RhoN)	385385 IMAGE Consortium	W74896
7097 B	9 021TAMOU	ESTs	1167716 IMAGE Consortium	AA689893
7098 B	11 021TAMOU	ESTs, Moderately similar to PHOSPHATIDYLINOSITOL 3-KINASE CATALYTIC SUBUNIT, BETA ISOFORM [Homo sapiens]	1178611 IMAGE Consortium	AA709957
7099 D	1 021TAMOU	ESTs, Highly similar to AF143096_1 peptidyl-prolyl cis-trans isomerase EPVH [H.sapiens]	1180304 IMAGE Consortium	AA755967
7100 D	3 021TAMOU	ESTs, Highly similar to MPPA_RAT MITOCHONDRIAL PROCESSING PEPTIDASE ALPHA SUBUNIT PRECURSOR [R.norvegicus]	387895 IMAGE Consortium	W66602
7101 D	5 021TAMOU	ESTs	375202 IMAGE Consortium	W62850
7102 D	7 021TAMOU	ESTs	385567 IMAGE Consortium	AI430494
7103 D	9 021TAMOU	mitogen activated protein kinase 9	1178040 IMAGE Consortium	AA718629
7104 D	11 021TAMOU	Mus musculus unknown mRNA	1177953 IMAGE Consortium	AA718560
7105 F	1 021TAMOU	EST	1180477 IMAGE Consortium	AI593261
7106 F	3 021TAMOU	ESTs	387991 IMAGE Consortium	AI430381
7107 F	5 021TAMOU	ESTs	373831 IMAGE Consortium	AI430084
7108 F	7 021TAMOU	vesicle-associated membrane protein 8	387062 IMAGE Consortium	W65964

7109 F	9 021TAMOU	ESTs	1178113 IMAGE Consortium	AA718696
7110 F	11 021TAMOU	ESTs	1225001 IMAGE Consortium	AA709656
7111 H	1 021TAMOU			
7112 H	3 021TAMOU	ESTs	373270 IMAGE Consortium	W57273
7113 H	5 021TAMOU	ESTs	374599 IMAGE Consortium	AI430456
7114 H	7 021TAMOU	Mus musculus mRNA for 1A13 protein	385624 IMAGE Consortium	W65200
7115 H	9 021TAMOU	ESTs, Moderately similar to beta-tubulin cofactor D [H.sapiens]	1178129 IMAGE Consortium	AA718704
7116 H	11 021TAMOU	ESTs	1225018 IMAGE Consortium	AA709788
7117 B	1 0210AMOV	TXK tyrosine kinase	1225170 IMAGE Consortium	AA762897
7118 B	3 0210AMOV	ESTs	1225559 IMAGE Consortium	AA762934
7119 B	5 0210AMOV	epidermal growth factor receptor pathway substrate 8	1195703 IMAGE Consortium	AA711869
7120 B	7 0210AMOV	ESTs	1195892 IMAGE Consortium	AA717014
7121 B	9 0210AMOV	complement component 3a receptor 1	1230537 IMAGE Consortium	AA879827
7122 B	11 0210AMOV	ESTs	1230873 IMAGE Consortium	AA880094
7123 D	1 0210AMOV	ESTs	1225193 IMAGE Consortium	AI428661
7124 D	3 0210AMOV	ESTs	1225627 IMAGE Consortium	AI428724
7125 D	5 0210AMOV	ESTs	1195754 IMAGE Consortium	AA711981
7126 D	7 0210AMOV	ESTs	1225955 IMAGE Consortium	AA739270
7127 D	9 0210AMOV	ESTs	1230552 IMAGE Consortium	AA879834
7128 D	11 0210AMOV	sialyltransferase 7 ((alpha-N-acetylneuraminy 2,3-betagalactosyl-1,3)-N-acetyl galactosaminide alpha-2,6-sialyltransferase) F	1230916 IMAGE Consortium	AA790409
7129 F	1 0210AMOV			
7130 F	3 0210AMOV	ESTs	1224970 IMAGE Consortium	AA709624
7131 F	5 0210AMOV	mastocytoma N-deacetylase/N-sulfotransferase	1195763 IMAGE Consortium	AA711997
7132 F	7 0210AMOV	ESTs	1195906 IMAGE Consortium	AA716786
7133 F	9 0210AMOV	ESTs	1230607 IMAGE Consortium	AA879870
7134 F	11 0210AMOV	ESTs	1230966 IMAGE Consortium	AA790455
7135 H	1 0210AMOV	ESTs	1226065 IMAGE Consortium	AA739381
7136 H	3 0210AMOV	a disintegrin-like and metalloprotease (reprolysin type) with thrombospondin type 1 motif, 1	1195684 IMAGE Consortium	AA711852
7137 H	5 0210AMOV			
7138 H	7 0210AMOV	ESTs, Weakly similar to APOLIPOPROTEIN B MRNA EDITING PROTEIN [Homo sapiens]	1195942 IMAGE Consortium	AA716815
7139 H	9 0210AMOV	SRY-box containing gene 10	1230665 IMAGE Consortium	AA879915
7140 H	11 0210AMOV	retinoblastoma-like 1 (p107)	1227032 IMAGE Consortium	AA738977
7141 B	1 0217AMOW	ESTs	1226576 IMAGE Consortium	AI425662
7142 B	3 0217AMOW	ESTs, Weakly similar to GLO2_RAT HYDROXYACYLGLUTATHIONE HYDROLASE [R.norvegicus]	1243679 IMAGE Consortium	AA822454
7143 B	5 0217AMOW	ESTs	1263237 IMAGE Consortium	AI509556
7144 B	7 0217AMOW	ESTs	1247515 IMAGE Consortium	AA763339

7145 B	9	0217AMOW	ESTs	1248615	IMAGE Consortium	AI465070
7146 B	11	0217AMOW	ESTs	1246415	IMAGE Consortium	AA823184
7147 D	1	0217AMOW	ESTs	1243393	IMAGE Consortium	AA789435
7148 D	3	0217AMOW	ESTs	1226954	IMAGE Consortium	AA738866
7149 D	5	0217AMOW	ESTs	1263315	IMAGE Consortium	AA856528
7150 D	7	0217AMOW	ESTs	1248271	IMAGE Consortium	AA982183
7151 D	9	0217AMOW	ESTs	1248640	IMAGE Consortium	AA759494
7152 D	11	0217AMOW	M.musculus mRNA for protein expressed at high levels in testis	1246432	IMAGE Consortium	AA823193
7153 F	1	0217AMOW	ESTs	1244161	IMAGE Consortium	AA790688
7154 F	3	0217AMOW	lurcher transcript 1	1243807	IMAGE Consortium	AA822676
7155 F	5	0217AMOW	ESTs	1248058	IMAGE Consortium	AA762233
7156 F	7	0217AMOW	ESTs	1262864	IMAGE Consortium	AA839578
7157 F	9	0217AMOW	ESTs	1246201	IMAGE Consortium	AA823010
7158 F	11	0217AMOW	ESTs, Highly similar to sushi-repeat-containing protein [M.musculus]	1247178	IMAGE Consortium	AI465834
7159 H	1	0217AMOW	ESTs	1226706	IMAGE Consortium	AA821501
7160 H	3	0217AMOW	ESTs, Moderately similar to SPRT_RAT SPROUTIN□ [R.norvegicus]	1243966	IMAGE Consortium	AA790247
7161 H	5	0217AMOW	ESTs, Weakly similar to AMMECR1 protein [H.sapiens]	1248104	IMAGE Consortium	AA762313
7162 H	7	0217AMOW	ESTs	1248296	IMAGE Consortium	AA789558
7163 H	9	0217AMOW	ESTs	1246258	IMAGE Consortium	AI466090
7164 H	11	0217AMOW	ESTs	1245774	IMAGE Consortium	AA822743
7165 B	1	021EAMOX	ESTs	1247310	IMAGE Consortium	AA839730
7166 B	3	021EAMOX	nuclear receptor co-repressor 1	1245205	IMAGE Consortium	AA821529
7167 B	5	021EAMOX	ESTs	1263376	IMAGE Consortium	AI509492
7168 B	7	021EAMOX	ESTs, Weakly similar to JC4863 homeobox protein zhx-1 - mouse□ [M.musculus]	1265051	IMAGE Consortium	AA881623
7169 B	9	021EAMOX	toll-like receptor 2	1263783	IMAGE Consortium	AA863729
7170 B	11	021EAMOX	ESTs	1264673	IMAGE Consortium	AA881395
7171 D	1	021EAMOX	ESTs	1247347	IMAGE Consortium	AA839767
7172 D	3	021EAMOX	ESTs	1246021	IMAGE Consortium	AA822429
7173 D	5	021EAMOX	ESTs, Highly similar to KIAA0169 protein [H.sapiens]	1264107	IMAGE Consortium	AI509457
7174 D	7	021EAMOX	ESTs, Weakly similar to MLZ4_MOUSE ZINC FINGER PROTEIN MLZ-4 [M.musculus]	1265071	IMAGE Consortium	AI509698
7175 D	9	021EAMOX	Public domain EST	1265241	IMAGE Consortium	
7176 D	11	021EAMOX	CD80 antigen	1263968	IMAGE Consortium	AA863833
7177 F	1	021EAMOX	ESTs	1245983	IMAGE Consortium	AI466042
7178 F	3	021EAMOX	ESTs, Weakly similar to S34198 IgE Fc receptor II, low-affinity - rat□ [R.norvegicus]	1245314	IMAGE Consortium	AI606925
7179 F	5	021EAMOX	syntaxin binding protein 4	1264173	IMAGE Consortium	AA919678
7180 F	7	021EAMOX	vinculin	1265092	IMAGE Consortium	AA881654
7181 F	9	021EAMOX	ESTs	1263889	IMAGE Consortium	AA874434

7182 F	11 021EAMOX	ESTs, Highly similar to BCL2_MOUSE APOPTOSIS REGULATOR BCL-2 [M.musculus]	1265430 IMAGE Consortium	AA867214
7183 H	1 021EAMOX	ESTs	1247428 IMAGE Consortium	AA839814
7184 H	3 021EAMOX	ESTs	1245331 IMAGE Consortium	AA821617
7185 H	5 021EAMOX	ESTs	1264210 IMAGE Consortium	AA867614
7186 H	7 021EAMOX	ESTs, Weakly similar to A53933 myosin I myr 4 - rat [R.norvegicus]	1264373 IMAGE Consortium	AA881346
7187 H	9 021EAMOX	troponin I, skeletal, slow 1	1265332 IMAGE Consortium	AA867133
7188 H	11 021EAMOX	ESTs	1265514 IMAGE Consortium	AA866989
7189 B	1 021LAMOY	ESTs	1264958 IMAGE Consortium	AA881517
7190 B	3 021LAMOY	EST	1281494 IMAGE Consortium	AI563818
7191 B	5 021LAMOY	glutamine synthetase pseudogene 1	1378629 IMAGE Consortium	AI019469
7192 B	7 021LAMOY	hypothetical protein	1380212 IMAGE Consortium	AI060442
7193 B	9 021LAMOY	laminin, beta 3	1364902 IMAGE Consortium	AI019837
7194 B	11 021LAMOY	interferon regulatory factor 1	1344724 IMAGE Consortium	AA939782
7195 D	1 021LAMOY	ESTs, Weakly similar to myeloid zinc finger protein-2 [M.musculus]	1265857 IMAGE Consortium	AA881904
7196 D	3 021LAMOY	ESTs	1281516 IMAGE Consortium	AA874408
7197 D	5 021LAMOY	ESTs	1379474 IMAGE Consortium	AI050556
7198 D	7 021LAMOY	T-box 2	1381068 IMAGE Consortium	AI035911
7199 D	9 021LAMOY	prostaglandin I receptor (IP)	1364948 IMAGE Consortium	AI019668
7200 D	11 021LAMOY	caspase 7	1344736 IMAGE Consortium	AA939788
7201 F	1 021LAMOY	ESTs	1328202 IMAGE Consortium	AA915591
7202 F	3 021LAMOY	ESTs	1281541 IMAGE Consortium	AA874467
7203 F	5 021LAMOY	karyopherin (importin) alpha 4	1378794 IMAGE Consortium	AI020155
7204 F	7 021LAMOY	myosin VIIa	1365095 IMAGE Consortium	AI019905
7205 F	9 021LAMOY	secretogranin III	1366406 IMAGE Consortium	AI021458
7206 F	11 021LAMOY	calcium/calmodulin-dependent protein kinase IV	1344831 IMAGE Consortium	AA939631
7207 H	1 021LAMOY	interleukin 2 receptor, beta chain	1328304 IMAGE Consortium	AA915712
7208 H	3 021LAMOY	colony stimulating factor 3 (granulocyte)	1314427 IMAGE Consortium	AA914420
7209 H	5 021LAMOY	prenylated SNARE protein	1378823 IMAGE Consortium	AI020451
7210 H	7 021LAMOY	excision repair cross-complementing rodent repair deficiency, complementation group 5	1380598 IMAGE Consortium	AI060824
7211 H	9 021LAMOY	ESTs	1365762 IMAGE Consortium	AI020735
7212 H	11 021LAMOY	Werner syndrome homolog (human)	1344845 IMAGE Consortium	AA939691
7213 B	1 021SAMOZ	ESTs, Weakly similar to CG10347 gene product [D.melanogaster]	1346577 IMAGE Consortium	AA958472
7214 B	3 021SAMOZ	proline-serine-threonine phosphatase-interacting protein 1	1349888 IMAGE Consortium	AA980931
7215 B	5 021SAMOZ	ESTs	1382293 IMAGE Consortium	AI037505
7216 B	7 021SAMOZ	ESTs	1382586 IMAGE Consortium	AI036362
7217 B	9 021SAMOZ	ESTs	1382986 IMAGE Consortium	AI120322
7218 B	11 021SAMOZ	antigen p97 (melanoma associated) identified by monoclonal antibodies 133.2 and 96.5	404366 IMAGE Consortium	AI322380

7219 D	1	021SAMOZ	ESTs	1345877	IMAGE Consortium	AA982084
7220 D	3	021SAMOZ	tumor necrosis factor receptor superfamily, member 17	1349977	IMAGE Consortium	AA989895
7221 D	5	021SAMOZ	Ras and a-factor-converting enzyme 1 homolog (S. cerevisiae)	1382320	IMAGE Consortium	AI037517
7222 D	7	021SAMOZ	basic leucine zipper transcription factor, ATF-like	1383316	IMAGE Consortium	AI120819
7223 D	9	021SAMOZ	IGFBP-like protein	403411	IMAGE Consortium	W82904
7224 D	11	021SAMOZ	ESTs	403657	IMAGE Consortium	W82196
7225 F	1	021SAMOZ	polyomavirus enhancer activator 3	1346663	IMAGE Consortium	AA959284
7226 F	3	021SAMOZ	cartilage derived retinoic acid sensitive protein	1382044	IMAGE Consortium	AA982842
7227 F	5	021SAMOZ	ESTs	1382360	IMAGE Consortium	AI037598
7228 F	7	021SAMOZ	ESTs	1382626	IMAGE Consortium	AI036402
7229 F	9	021SAMOZ				
7230 F	11	021SAMOZ	ESTs	404428	IMAGE Consortium	W83146
7231 H	1	021SAMOZ	calcium/calmodulin-dependent protein kinase kinase 1, alpha	1348181	IMAGE Consortium	AA940207
7232 H	3	021SAMOZ	allograft inflammatory factor 1	1382096	IMAGE Consortium	AA797956
7233 H	5	021SAMOZ	ESTs	1382402	IMAGE Consortium	AI046514
7234 H	7	021SAMOZ	ESTs	1382640	IMAGE Consortium	AI036487
7235 H	9	021SAMOZ	UDP-Gal:betaGlcNAc beta 1,4-galactosyltransferase, polypeptide 5	403139	IMAGE Consortium	AI425818
7236 H	11	021SAMOZ	ESTs	403717	IMAGE Consortium	W82292
7237 B	1	021SAMP0	ESTs	406363	IMAGE Consortium	W83451
7238 B	3	021SAMP0	ESTs	405925	IMAGE Consortium	AI425340
7239 B	5	021SAMP0	ESTs	407944	IMAGE Consortium	AI430265
7240 B	7	021SAMP0	ATP-binding cassette, sub-family B (MDR/TAP), member 3	1397130	IMAGE Consortium	AI098411
7241 B	9	021SAMP0	Paneth cell enhanced expression	1384160	IMAGE Consortium	AI117296
7242 B	11	021SAMP0	ESTs	1383726	IMAGE Consortium	AI180894
7243 D	1	021SAMP0	ESTs, Highly similar to ZINC FINGER PROTEIN 91 [Homo sapiens]	407137	IMAGE Consortium	AI893319
7244 D	3	021SAMP0	ESTs	405990	IMAGE Consortium	W83301
7245 D	5	021SAMP0	ESTs	1383826	IMAGE Consortium	AI181682
7246 D	7	021SAMP0	brain derived neurotrophic factor	1397218	IMAGE Consortium	AI098498
7247 D	9	021SAMP0	cytochrome P450, subfamily IV B, polypeptide 1	1384195	IMAGE Consortium	AI181090
7248 D	11	021SAMP0	ESTs	1383766	IMAGE Consortium	AI180925
7249 F	1	021SAMP0	amyloid beta (A4) precursor-like protein 1	406434	IMAGE Consortium	W83478
7250 F	3	021SAMP0	ESTs	406734	IMAGE Consortium	W83739
7251 F	5	021SAMP0	tubulin cofactor a	1383836	IMAGE Consortium	AI181686
7252 F	7	021SAMP0	Mus musculus NK cell receptor 2B4 mRNA, complete cds	1395874	IMAGE Consortium	AI180741
7253 F	9	021SAMP0	ESTs, Weakly similar to PLASMOLIPIN [Rattus norvegicus]	1383543	IMAGE Consortium	AI120958
7254 F	11	021SAMP0	ESTs, Moderately similar to SAST_RAT S-ACYL FATTY ACID SYNTHASE THIOESTERASE, MEDIUM CHAIN	1383784	IMAGE Consortium	AI180945



		[R.norvegicus]		
7255 H	1	021SAMP0	ESTs	407191 IMAGE Consortium W84213
7256 H	3	021SAMP0	lysyl oxidase	406747 IMAGE Consortium W83882
7257 H	5	021SAMP0	ESTs	1383851 IMAGE Consortium AI181716
7258 H	7	021SAMP0	latent transforming growth factor beta binding protein 2	1396864 IMAGE Consortium AI099009
7259 H	9	021SAMP0	platelet factor 4	1383594 IMAGE Consortium AI120986
7260 H	11	021SAMP0	ESTs, Moderately similar to AF089897_1 topoisomerase-related function protein [H.sapiens]	1430827 IMAGE Consortium AI182256
7261 B	1	021AMSI	ESTs	1433770 IMAGE Consortium AI606617
7262 B	3	021AMSI	CCAAT/enhancer binding protein alpha (C/EBP), related sequence 1	1446242 IMAGE Consortium AI151993
7263 B	5	021AMSI	claudin 5	1430427 IMAGE Consortium AI182191
7264 B	7	021AMSI	ESTs, Weakly similar to KIAA0874 protein [H.sapiens]	1400720 IMAGE Consortium AI121535
7265 B	9	021AMSI	ESTs, Highly similar to unnamed protein product [H.sapiens]	1399550 IMAGE Consortium AI121359
7266 B	11	021AMSI	ESTs	1448906 IMAGE Consortium AI573593
7267 D	1	021AMSI	ESTs	1434610 IMAGE Consortium AI877327
7268 D	3	021AMSI	interferon gamma inducing factor binding protein	1445869 IMAGE Consortium AI122010
7269 D	5	021AMSI	ESTs	1430524 IMAGE Consortium AI182827
7270 D	7	021AMSI	ESTs	1429022 IMAGE Consortium AI183169
7271 D	9	021AMSI	microrchidia	1428933 IMAGE Consortium AI183160
7272 D	11	021AMSI	ESTs	1448951 IMAGE Consortium AI573630
7273 F	1	021AMSI	ESTs	1433922 IMAGE Consortium AA840161
7274 F	3	021AMSI	ESTs	1429924 IMAGE Consortium AI181398
7275 F	5	021AMSI	ESTs, Highly similar to unnamed protein product [H.sapiens]	1401003 IMAGE Consortium AI122049
7276 F	7	021AMSI	ESTs, Weakly similar to myosin-Va [R.norvegicus]	1400864 IMAGE Consortium AI121707
7277 F	9	021AMSI	ESTs	1399799 IMAGE Consortium AI115773
7278 F	11	021AMSI	ESTs	1449767 IMAGE Consortium AI153139
7279 H	1	021AMSI	ESTs	1433927 IMAGE Consortium AI608003
7280 H	3	021AMSI	ESTs, Moderately similar to KIAA1139 protein [H.sapiens]	1430006 IMAGE Consortium AI122219
7281 H	5	021AMSI	orosomuroid 1	1400397 IMAGE Consortium AI117779
7282 H	7	021AMSI	ESTs, Weakly similar to putative type III alcohol dehydrogenase [D.melanogaster]	1400920 IMAGE Consortium AI121911
7283 H	9	021AMSI	apyrase 1, homolog (C. lectularius)	1399959 IMAGE Consortium AI121485
7284 H	11	021AMSI	baculoviral IAP repeat-containing 1a	1449810 IMAGE Consortium AI153148
7285 B	1	0216AMP2	histocompatibility 2, class II, locus Mb2	1446110 IMAGE Consortium AI157238
7286 B	3	0216AMP2	ESTs, Weakly similar to matrin cyclophilin [R.norvegicus]	1448063 IMAGE Consortium AI154653
7287 B	5	0216AMP2	ELK1, member of ETS oncogene family	1449058 IMAGE Consortium AI663752
7288 B	7	0216AMP2	ESTs	1478098 IMAGE Consortium AI152690
7289 B	9	0216AMP2	ESTs	1478433 IMAGE Consortium AI152678
7290 B	11	0216AMP2	ESTs	1478742 IMAGE Consortium AI157691
7291 D	1	0216AMP2	cAMP responsive element binding protein 1	1447005 IMAGE Consortium AI158037

7292 D	3 0216AMP2	ESTs, Weakly similar to ZF37_MOUSE ZINC FINGER PROTEIN 37 [M.musculus]	1447421 IMAGE Consortium	AI154242
7293 D	5 0216AMP2	chemokine (C-C) receptor 7	1446936 IMAGE Consortium	AI158536
7294 D	7 0216AMP2	ESTs	1478174 IMAGE Consortium	AI152405
7295 D	9 0216AMP2	hematopoietic cell transcript 1	1478461 IMAGE Consortium	AI152865
7296 D	11 0216AMP2	ESTs	1478798 IMAGE Consortium	AI197473
7297 F	1 0216AMP2			
7298 F	3 0216AMP2	stimulated by retinoic acid gene 6	1448168 IMAGE Consortium	AI158798
7299 F	5 0216AMP2	ESTs	1449138 IMAGE Consortium	AI153732
7300 F	7 0216AMP2	ESTs, Weakly similar to AF158182_1 preprocathepsin P [M.musculus]	1477580 IMAGE Consortium	AI155314
7301 F	9 0216AMP2	caspase 9	1477823 IMAGE Consortium	AI155222
7302 F	11 0216AMP2	ESTs	1494997 IMAGE Consortium	AI157392
7303 H	1 0216AMP2	ESTs, Moderately similar to unknown protein IT12 [H.sapiens]	1447151 IMAGE Consortium	AI158359
7304 H	3 0216AMP2	interferon gamma receptor	1448192 IMAGE Consortium	AI158816
7305 H	5 0216AMP2	ESTs	1448539 IMAGE Consortium	AI573352
7306 H	7 0216AMP2	ESTs	1478305 IMAGE Consortium	AI152538
7307 H	9 0216AMP2	raf-related oncogene	1477850 IMAGE Consortium	AI155246
7308 H	11 0216AMP2	homeo box B7	1479853 IMAGE Consortium	AI156326
7309 B	1 021DAMP3	homeo box C4	1749576 IMAGE Consortium	AI047090
7310 B	3 021DAMP3	ESTs, Weakly similar to XLR_MOUSE X-LINKED LYMPHOCYTE-REGULATED PROTEIN PM1 [M.musculus]	1746104 IMAGE Consortium	AI036904
7311 B	5 021DAMP3	ESTs, Weakly similar to F55A12.9 [C.elegans]	1746471 IMAGE Consortium	AI037699
7312 B	7 021DAMP3	ESTs, Moderately similar to similar to putative product coded in C.elegans cosmid C01C10. [H.sapiens]	1749181 IMAGE Consortium	AI047920
7313 B	9 021DAMP3	brain protein	1617134 IMAGE Consortium	AA967857
7314 B	11 021DAMP3	ESTs	1617321 IMAGE Consortium	AA982464
7315 D	1 021DAMP3	ESTs	1749607 IMAGE Consortium	AI047122
7316 D	3 021DAMP3	ESTs	1746193 IMAGE Consortium	AI037647
7317 D	5 021DAMP3	ESTs, Moderately similar to DIAC_RAT DI-N-ACETYLCHITOBIASE PRECURSOR [R.norvegicus]	1745849 IMAGE Consortium	AI789630
7318 D	7 021DAMP3	Mus musculus clone L3 variable group of 2-cell-stage gene family mRNA, complete cds	1749470 IMAGE Consortium	AI046759
7319 D	9 021DAMP3	ESTs	1617185 IMAGE Consortium	AA968016
7320 D	11 021DAMP3	tumor necrosis factor receptor superfamily, member 9	1497753 IMAGE Consortium	AI664286
7321 F	1 021DAMP3	LIM homeobox protein 1	1749654 IMAGE Consortium	AI046793
7322 F	3 021DAMP3	ESTs	1745549 IMAGE Consortium	AI173036
7323 F	5 021DAMP3	ESTs	1745887 IMAGE Consortium	AI789680
7324 F	7 021DAMP3	6-phosphofructo-2-kinase/fructose-2,6-biphosphatase 1	1511542 IMAGE Consortium	AI663904
7325 F	9 021DAMP3	chitinase 5 (oviductin)	1498222 IMAGE Consortium	AI226220
7326 F	11 021DAMP3	placental growth factor	1617488 IMAGE Consortium	AA982549

7327 H	1 021DAMP3	crystallin, gamma D	1749741 IMAGE Consortium	AI048379
7328 H	3 021DAMP3			
7329 H	5 021DAMP3	testis specific gene 1	1746656 IMAGE Consortium	AI046576
7330 H	7 021DAMP3	ESTs, Weakly similar to KIAA0475 protein [H.sapiens]	1510918 IMAGE Consortium	AI227453
7331 H	9 021DAMP3	nucleosome assembly protein 1-like 2	1617201 IMAGE Consortium	AA982396
7332 H	11 021DAMP3	ESTs	1617508 IMAGE Consortium	AA982569
7333 B	1 021KAMP4	heart and neural crest derivatives expressed transcript 1	1851169 IMAGE Consortium	AI197413
7334 B	3 021KAMP4	glial cell line derived neurotrophic factor family receptor alpha 2	1764307 IMAGE Consortium	AI047785
7335 B	5 021KAMP4	deoxycytidine kinase	537045 IMAGE Consortium	AA117124
7336 B	7 021KAMP4			
7337 B	9 021KAMP4	solute carrier family 2 (facilitated glucose transporter), member 1	536794 IMAGE Consortium	AA114539
7338 B	11 021KAMP4	potassium channel modulatory factor DEBT-91	537755 IMAGE Consortium	AI550132
7339 D	1 021KAMP4	tolloid-like	1851223 IMAGE Consortium	AI197263
7340 D	3 021KAMP4	serine/threonine kinase 10	1764541 IMAGE Consortium	AI115517
7341 D	5 021KAMP4	Mus musculus E-cadherin binding protein E7 mRNA, complete cds	537060 IMAGE Consortium	AA114742
7342 D	7 021KAMP4	sema domain, seven thrombospondin repeats (type 1 and type 1-like), transmembrane domain (TM) and short cytoplasmic domain, (semaphorin) 5B	537250 IMAGE Consortium	AA118106
7343 D	9 021KAMP4	NY-REN-18 antigen	537544 IMAGE Consortium	AI550116
7344 D	11 021KAMP4	ESTs, Highly similar to APOLIPOPROTEIN B-100 PRECURSOR [Homo sapiens]	537847 IMAGE Consortium	AA120586
7345 F	1 021KAMP4	ESTs	1851394 IMAGE Consortium	AI194360
7346 F	3 021KAMP4			
7347 F	5 021KAMP4			
7348 F	7 021KAMP4	ESTs	537266 IMAGE Consortium	AI550101
7349 F	9 021KAMP4	transcriptional regulator, SIN3A (yeast)	537589 IMAGE Consortium	AI550122
7350 F	11 021KAMP4	G protein-coupled receptor kinase 6	538067 IMAGE Consortium	AA120287
7351 H	1 021KAMP4			
7352 H	3 021KAMP4			
7353 H	5 021KAMP4	ESTs, Weakly similar to AF220193_1 uncharacterized hypothalamus protein HT007 [H.sapiens]	537108 IMAGE Consortium	AA117961
7354 H	7 021KAMP4	ESTs, Moderately similar to MAK16 PROTEIN [Saccharomyces cerevisiae]	537328 IMAGE Consortium	AI325769
7355 H	9 021KAMP4	DNA segment, Chr 16, Wayne State University 109, expressed	537595 IMAGE Consortium	AA118151
7356 H	11 021KAMP4	ESTs, Moderately similar to ZINC FINGER Y-CHROMOSOMAL PROTEIN 1 [Xenopus laevis]	538090 IMAGE Consortium	AI325780
7357 B	1 021RAMP5	ESTs	538217 IMAGE Consortium	AI550157
7358 B	3 021RAMP5	ESTs	538460 IMAGE Consortium	AI550173
7359 B	5 021RAMP5			

7360 B	7 021RAMP5	serine/arginine-rich protein specific kinase 1	988849 IMAGE Consortium	AA638416
7361 B	9 021RAMP5	serum-inducible kinase	403206 IMAGE Consortium	W81741
7362 B	11 021RAMP5	protein phosphatase 5, catalytic subunit	404009 IMAGE Consortium	AI894334
7363 D	1 021RAMP5	ESTs, Moderately similar to unnamed protein product [H.sapiens]	538252 IMAGE Consortium	AA116636
7364 D	3 021RAMP5	ESTs, Weakly similar to KIAA0616 protein [H.sapiens]	538506 IMAGE Consortium	AA117368
7365 D	5 021RAMP5	ESTs, Highly similar to URIDYLATE KINASE [Saccharomyces cerevisiae]	537955 IMAGE Consortium	AI892472
7366 D	7 021RAMP5	Shc SH2-domain binding protein 1	2101969 IMAGE Consortium	AW210261
7367 D	9 021RAMP5	aconitase 1	403275 IMAGE Consortium	W82672
7368 D	11 021RAMP5	bromodomain-containing 2	400539 IMAGE Consortium	AI425711
7369 F	1 021RAMP5	ESTs	538296 IMAGE Consortium	AI550163
7370 F	3 021RAMP5	ESTs, Weakly similar to unknown [S.cerevisiae]	538544 IMAGE Consortium	AA116816
7371 F	5 021RAMP5	heterogeneous nuclear ribonucleoprotein A2/B1	537968 IMAGE Consortium	AI550144
7372 F	7 021RAMP5	SIK similar protein	977310 IMAGE Consortium	AA560326
7373 F	9 021RAMP5	ESTs, Highly similar to P2UR_MOUSE P2U PURINOCEPTOR 1 [M.musculus]	405832 IMAGE Consortium	W83063
7374 F	11 021RAMP5	CD37 antigen	403475 IMAGE Consortium	AI893967
7375 H	1 021RAMP5	ESTs, Moderately similar to AF196304_1 SUMO-1-specific protease [H.sapiens]	538308 IMAGE Consortium	AI550164
7376 H	3 021RAMP5	ESTs	537835 IMAGE Consortium	AI550135
7377 H	5 021RAMP5	integrin beta 1 (fibronectin receptor beta)	538729 IMAGE Consortium	AA123128
7378 H	7 021RAMP5			
7379 H	9 021RAMP5	ESTs	401568 IMAGE Consortium	AI425744
7380 H	11 021RAMP5	huntingtin interacting protein 1 related	404331 IMAGE Consortium	W82687
7381 B	1 021YAMP6	uracil-DNA glycosylase	406824 IMAGE Consortium	W83782
7382 B	3 021YAMP6	chaperonin subunit 7 (eta)	407165 IMAGE Consortium	W84206
7383 B	5 021YAMP6	Mus musculus COP9 complex subunit 4 (COPS4) mRNA, complete cds {IMAGE:438210}	438210 IMAGE Consortium	AA008135
7384 B	7 021YAMP6	SEC15 homolog (S. cerevisiae)	423053 IMAGE Consortium	W97996
7385 B	9 021YAMP6	solute carrier family 27 (fatty acid transporter), member 5	440520 IMAGE Consortium	AI552452
7386 B	11 021YAMP6	tight junction protein 3	444761 IMAGE Consortium	AA016336
7387 D	1 021YAMP6	interferon (alpha and beta) receptor	407979 IMAGE Consortium	W84257
7388 D	3 021YAMP6	SWI/SNF related, matrix associated, actin dependent regulator of chromatin, subfamily b, member 1	407181 IMAGE Consortium	W84225
7389 D	5 021YAMP6	IMAGE EST {IMAGE:438749}	438749 IMAGE Consortium	AA051080
7390 D	7 021YAMP6	myotubularin related protein 7	426021 IMAGE Consortium	AA000871
7391 D	9 021YAMP6	fibroblast growth factor receptor 2	444034 IMAGE Consortium	AA014912
7392 D	11 021YAMP6	tumor-suppressing subchromosomal transferable fragment 3	441969 IMAGE Consortium	AA014727
7393 F	1 021YAMP6	transcription factor UBF	406843 IMAGE Consortium	AI893975
7394 F	3 021YAMP6	Paraoxonase 2 {IMAGE:423266}	423266 IMAGE Consortium	W98586
7395 F	5 021YAMP6	guanylate kinase membrane-associated inverted 1	426596 IMAGE Consortium	AA002500

7396 F	7 021YAMP6	histone deacetylase 3	427571 IMAGE Consortium	AA002536
7397 F	9 021YAMP6	centromere autoantigen A	439365 IMAGE Consortium	AA004134
7398 F	11 021YAMP6	GPI anchor attachment protein 1	439057 IMAGE Consortium	AA008266
7399 H	1 021YAMP6	NADH dehydrogenase (ubiquinone) 1 alpha subcomplex 2 (8 kDa)	406190 IMAGE Consortium	W83909
7400 H	3 021YAMP6	Myelin transcription factor 1-like {IMAGE:423114}	423114 IMAGE Consortium	W97257
7401 H	5 021YAMP6	ATPase, Ca <sup>++</sup> transporting, cardiac muscle, slow twitch 2	437239 IMAGE Consortium	AA003458
7402 H	7 021YAMP6	glucose phosphate isomerase 1 complex	438517 IMAGE Consortium	AA050875
7403 H	9 021YAMP6	cathepsin J	442312 IMAGE Consortium	AA013726
7404 H	11 021YAMP6	leukemia/lymphoma related factor	442939 IMAGE Consortium	AA014909
7405 B	1 0215AMP7	Chaperonin subunit 4 (delta) {IMAGE:459668}	459668 IMAGE Consortium	AA027583
7406 B	3 0215AMP7	phosphatidylinositol 3-kinase, regulatory subunit, polypeptide 2 (p85 beta)	464298 IMAGE Consortium	AA033042
7407 B	5 0215AMP7	IMAGE EST {IMAGE:479471}	479471 IMAGE Consortium	AA049350
7408 B	7 0215AMP7	calmodulin	479619 IMAGE Consortium	AI606650
7409 B	9 0215AMP7	IMAGE EST {IMAGE:481630}	481630 IMAGE Consortium	AA060255
7410 B	11 0215AMP7	fragile X mental retardation gene, autosomal homolog	484191 IMAGE Consortium	AA080092
7411 D	1 0215AMP7	Myeloperoxidase {IMAGE:466693}	466693 IMAGE Consortium	AA031015
7412 D	3 0215AMP7	leukotriene B4 receptor 1	458727 IMAGE Consortium	AA023970
7413 D	5 0215AMP7	Beta-spectrin 3 {IMAGE:478354}	478354 IMAGE Consortium	AA049581
7414 D	7 0215AMP7	DiGeorge syndrome gene c	479657 IMAGE Consortium	AA051453
7415 D	9 0215AMP7	Tubulin, beta 2 {IMAGE:482602}	482602 IMAGE Consortium	AA061838
7416 D	11 0215AMP7	Mus musculus APC-binding protein EB2 mRNA, partial cds	482240 IMAGE Consortium	AI323100
7417 F	1 0215AMP7	Public domain EST	459075 IMAGE Consortium	
7418 F	3 0215AMP7	Snail homolog, (Drosophila) {IMAGE:469080}	469080 IMAGE Consortium	AA036380
7419 F	5 0215AMP7	ESTs, Moderately similar to GTP:AMP PHOSPHOTRANSFERASE MITOCHONDRIAL [R.norvegicus] {IMAGE:479750}	479750 IMAGE Consortium	AA051422
7420 F	7 0215AMP7	proteasome (prosome, macropain) subunit, beta type 2	479711 IMAGE Consortium	AA051665
7421 F	9 0215AMP7	Paired-like homeodomain transcription factor 3 {IMAGE:482871}	482871 IMAGE Consortium	AA062140
7422 F	11 0215AMP7	Harvey rat sarcoma virus oncogene	483791 IMAGE Consortium	AI323132
7423 H	1 0215AMP7	nuclear transcription factor-Y alpha	459572 IMAGE Consortium	AA027394
7424 H	3 0215AMP7			
7425 H	5 0215AMP7	Mus musculus branched chain alpha ketoacid dehydrogenase kinase (bckd-kinase) mRNA, nuclear gene encoding mitochondrial protein, complete cds {IMAGE:480575}	480575 IMAGE Consortium	AA059497
7426 H	7 0215AMP7	apolipoprotein A-II	479777 IMAGE Consortium	AA051684
7427 H	9 0215AMP7	IMAGE EST {IMAGE:482001}	482001 IMAGE Consortium	AA059886
7428 H	11 0215AMP7	apolipoprotein B editing complex 2	482571 IMAGE Consortium	AA061902
7429 B	1 021CAMP8	Mus musculus calpain Lp82 mRNA, complete cds	574491 IMAGE Consortium	AA117626

		{IMAGE:574491}		
7430 B	3	021CAMP8	ESTs {IMAGE:577739}	577739 IMAGE Consortium AA139223
7431 B	5	021CAMP8	Mus musculus aldehyde dehydrogenase Ahd-2-like mRNA, complete cds {IMAGE:596348}	596348 IMAGE Consortium AA122814
7432 B	7	021CAMP8	malate dehydrogenase, soluble	635526 IMAGE Consortium AA165713
7433 B	9	021CAMP8	hnRNP-associated with lethal yellow	636522 IMAGE Consortium AA185293
7434 B	11	021CAMP8	catenin alpha 2	658649 IMAGE Consortium AA221924
7435 D	1	021CAMP8	Mus musculus TIA mRNA, complete cds {IMAGE:575920}	575920 IMAGE Consortium AA117605
7436 D	3	021CAMP8		
7437 D	5	021CAMP8	SEC61, gamma subunit ( <i>S. cerevisiae</i> )	597242 IMAGE Consortium AA145254
7438 D	7	021CAMP8	N-myc downstream regulated 2	634855 IMAGE Consortium AA163503
7439 D	9	021CAMP8	ESTs, Weakly similar to HYPOTHETICAL PROTEIN F44G4.1 IN CHROMOSOME II [ <i>C.elegans</i> ] {IMAGE:658190}	658190 IMAGE Consortium AA198227
7440 D	11	021CAMP8	LIM only 4	658726 IMAGE Consortium AA222003
7441 F	1	021CAMP8	deoxyribonuclease I	570643 IMAGE Consortium AA108495
7442 F	3	021CAMP8	promyelocytic leukemia	579866 IMAGE Consortium AA116994
7443 F	5	021CAMP8	glutamate-cysteine ligase, catalytic subunit	598795 IMAGE Consortium AA163042
7444 F	7	021CAMP8	thyroid hormone receptor-associated protein 100 kDa	634879 IMAGE Consortium AA163511
7445 F	9	021CAMP8	Cadherin 2 {IMAGE:671834}	671834 IMAGE Consortium AA242226
7446 F	11	021CAMP8	ribosomal protein S6 kinase polypeptide 1	671944 IMAGE Consortium AA244933
7447 H	1	021CAMP8	G protein-coupled receptor 33	575313 IMAGE Consortium AI386076
7448 H	3	021CAMP8	epidermal growth factor	580236 IMAGE Consortium AI326499
7449 H	5	021CAMP8	DNA Segment, Chr 15 Massachusetts Institute of Technology 260	597026 IMAGE Consortium AA145461
7450 H	7	021CAMP8	Ngfi-A binding protein 1	621117 IMAGE Consortium AI592274
7451 H	9	021CAMP8	Trophoblast specific protein {IMAGE:659899}	659899 IMAGE Consortium AA220024
7452 H	11	021CAMP8	YY1 transcription factor	669833 IMAGE Consortium AA230457
7453 B	1	021JAMP9	Mus musculus Masp-2 mRNA for mannose-binding lectin associated serine protease-2, complete cds {IMAGE:680232}	680232 IMAGE Consortium AA238260
7454 B	3	021JAMP9	sperm specific antigen 1	681433 IMAGE Consortium AA237943
7455 B	5	021JAMP9	Carbon catabolite repression 4 homolog ( <i>S. cerevisiae</i> ) {IMAGE:717197}	717197 IMAGE Consortium AA404014
7456 B	7	021JAMP9	TRANSDUCIN-LIKE ENHANCER PROTEIN 3 {IMAGE:722501}	722501 IMAGE Consortium AA253995
7457 B	9	021JAMP9	RAS protein-specific guanine nucleotide-releasing factor 1	720305 IMAGE Consortium AA261374
7458 B	11	021JAMP9	Glycosylation dependent cell adhesion molecule 1 {IMAGE:749605}	749605 IMAGE Consortium AA288467
7459 D	1	021JAMP9	ESTs, Weakly similar to BROWN PROTEIN [ <i>D.melanogaster</i> ] {IMAGE:680198}	680198 IMAGE Consortium AA237916
7460 D	3	021JAMP9	cytochrome P450, 1a2, aromatic compound inducible	680035 IMAGE Consortium AA242360
7461 D	5	021JAMP9	cytochrome P450, 2a12	692245 IMAGE Consortium AA238327

7462 D	7 021JAMP9	Transforming growth factor, beta induced, 68 kDa {IMAGE:734101}	734101 IMAGE Consortium	AA268592
7463 D	9 021JAMP9	alpha 1 microglobulin/bikunin	720375 IMAGE Consortium	AA261468
7464 D	11 021JAMP9	Fibroblast growth factor inducible 15 {IMAGE:750018}	750018 IMAGE Consortium	AA388985
7465 F	1 021JAMP9	Mus musculus D9 splice variant 2 mRNA, complete cds {IMAGE:679322}	679322 IMAGE Consortium	AA244657
7466 F	3 021JAMP9	ribosomal protein S6	679374 IMAGE Consortium	AI390326
7467 F	5 021JAMP9	calbindin-28K	697469 IMAGE Consortium	AA245046
7468 F	7 021JAMP9	Laminin, alpha 4 {IMAGE:736616}	736616 IMAGE Consortium	AA270669
7469 F	9 021JAMP9	hemolytic complement	719686 IMAGE Consortium	AA254990
7470 F	11 021JAMP9	interleukin 12b	750641 IMAGE Consortium	AA288235
7471 H	1 021JAMP9	interleukin 1 receptor accessory protein	678770 IMAGE Consortium	AA239486
7472 H	3 021JAMP9	small GTPase, homolog (S. cerevisiae)	681035 IMAGE Consortium	AA250086
7473 H	5 021JAMP9			
7474 H	7 021JAMP9	interleukin 2 receptor, gamma chain	721711 IMAGE Consortium	AA260777
7475 H	9 021JAMP9			
7476 H	11 021JAMP9	coagulation factor V	747430 IMAGE Consortium	AA272843
7477 B	1 0213AMPA	microfibrillar-associated protein 2	736871 IMAGE Consortium	AA271914
7478 B	3 0213AMPA	ESTs {IMAGE:790569}	790569 IMAGE Consortium	AA290020
7479 B	5 0213AMPA	cyclic nucleotide phosphodiesterase 1	777624 IMAGE Consortium	AA276865
7480 B	7 0213AMPA	heat shock protein cognate 70, testis	846935 IMAGE Consortium	AA433639
7481 B	9 0213AMPA	related to Drosophila groucho gene	317192 IMAGE Consortium	W13670
7482 B	11 0213AMPA	adipocyte complement related protein of 30 kDa	864276 IMAGE Consortium	AA458072
7483 D	1 0213AMPA	vanilloid receptor-like protein 1	747109 IMAGE Consortium	AA274980
7484 D	3 0213AMPA	Ecotropic viral integration site 5 {IMAGE:791360}	791360 IMAGE Consortium	AA415421
7485 D	5 0213AMPA	apoptosis inhibitory 6	764624 IMAGE Consortium	AA274023
7486 D	7 0213AMPA	sialyltransferase 7 ((alpha-N-acetylneuraminyl 2,3-beta-galactosyl-1,3)-N-acetyl galactosaminide alpha-2,6-sialyltransferase) B	831797 IMAGE Consortium	AA444639
7487 D	9 0213AMPA	Tousled-like kinase (Arabidopsis)	317858 IMAGE Consortium	W12907
7488 D	11 0213AMPA	protein phosphatase 1, regulatory (inhibitor) subunit 5	870856 IMAGE Consortium	AA462037
7489 F	1 0213AMPA	SH2-B PH domain containing signaling mediator 1	747914 IMAGE Consortium	AA260854
7490 F	3 0213AMPA	catenin (cadherin-associated protein), delta 2 (neural plakophilin-related arm-repeat protein)	765554 IMAGE Consortium	AA275119
7491 F	5 0213AMPA	microtubule-associated protein 7	791737 IMAGE Consortium	AA414664
7492 F	7 0213AMPA	creatine kinase, mitochondrial 1, ubiquitous	849182 IMAGE Consortium	AA469799
7493 F	9 0213AMPA	pyridoxal (pyridoxine, vitamin B6) kinase	317479 IMAGE Consortium	W13533
7494 F	11 0213AMPA	hexokinase 2	870931 IMAGE Consortium	AA462165
7495 H	1 0213AMPA	ATP synthase, H+ transporting, mitochondrial FO complex, subunit g	747976 IMAGE Consortium	AA260966
7496 H	3 0213AMPA	meprin 1 alpha	790399 IMAGE Consortium	AA290091
7497 H	5 0213AMPA	Catenin alpha 1 {IMAGE:818508}	818508 IMAGE Consortium	AA434955
7498 H	7 0213AMPA	septin 4	849273 IMAGE Consortium	AA469889

7499 H	9 0213AMPA	Mus musculus ras-related protein (rab18) mRNA, complete cds {IMAGE:851551}	851551 IMAGE Consortium	AA461854
7500 H	11 0213AMPA	regulatory protein, T lymphocyte 1	871756 IMAGE Consortium	AI614435
7501 B	1 021AAMPB	DNA-binding protein A	871974 IMAGE Consortium	AA473431
7502 B	3 021AAMPB	Mus musculus mRNA for tissue kallikrein, complete cds	889863 IMAGE Consortium	AA498574
7503 B	5 021AAMPB	SRY-box containing gene 15	963268 IMAGE Consortium	AA671659
7504 B	7 021AAMPB	retinal degeneration B2 homolog (Drosophila)	1108822 IMAGE Consortium	AA647446
7505 B	9 021AAMPB	3-hydroxy-3-methylglutaryl-Coenzyme A reductase	1195828 IMAGE Consortium	AA712088
7506 B	11 021AAMPB			
7507 D	1 021AAMPB	neutral amino acid transporter	875568 IMAGE Consortium	AA475589
7508 D	3 021AAMPB	serine/threonine kinase 25 (Ste20, yeast homolog)	902505 IMAGE Consortium	AA518501
7509 D	5 021AAMPB	calpain 8	959635 IMAGE Consortium	AA546660
7510 D	7 021AAMPB	caspase 12	1096376 IMAGE Consortium	AA871539
7511 D	9 021AAMPB	ATPase, Ca <sup>++</sup> transporting, cardiac muscle, fast twitch 1	1195881 IMAGE Consortium	AA717019
7512 D	11 021AAMPB	procollagen C-proteinase enhancer protein	1162727 IMAGE Consortium	AA770854
7513 F	1 021AAMPB	Mus musculus cGMP phosphodiesterase (PDE9A*1) mRNA, complete cds {IMAGE:889753}	889753 IMAGE Consortium	AA498410
7514 F	3 021AAMPB	house-keeping protein 1	903250 IMAGE Consortium	AA518970
7515 F	5 021AAMPB	phosphodiesterase 8	1066664 IMAGE Consortium	AA611596
7516 F	7 021AAMPB	interleukin 1 beta	1139544 IMAGE Consortium	AA667815
7517 F	9 021AAMPB	neuropilin	1166099 IMAGE Consortium	AA710379
7518 F	11 021AAMPB	thyroid hormone responsive SPOT14 homolog (Rattus)	1195446 IMAGE Consortium	AA756672
7519 H	1 021AAMPB	Histocompatibility 2, D region locus 1 {IMAGE:920425}	920425 IMAGE Consortium	AA538511
7520 H	3 021AAMPB	ets variant gene 6 (TEL oncogene)	888471 IMAGE Consortium	AA497592
7521 H	5 021AAMPB	solute carrier family 11 (proton-coupled divalent metal ion transporters), member 2	1055033 IMAGE Consortium	AA619114
7522 H	7 021AAMPB	basic transcription factor 3	1138873 IMAGE Consortium	AA688716
7523 H	9 021AAMPB	solute carrier family 25 (mitochondrial carrier; adenine nucleotide translocator), member 4	1195917 IMAGE Consortium	AA716803
7524 H	11 021AAMPB	oxidative stress induced	1195498 IMAGE Consortium	AA690750
7525 B	1 021HAMPC	cullin 1	1178842 IMAGE Consortium	AA734523
7526 B	3 021HAMPC	ESTs	1245757 IMAGE Consortium	AA798849
7527 B	5 021HAMPC	periaxin	1332330 IMAGE Consortium	AI159496
7528 B	7 021HAMPC	thymoma viral proto-oncogene 3	1330420 IMAGE Consortium	AA929481
7529 B	9 021HAMPC	ESTs	1328535 IMAGE Consortium	AA915566
7530 B	11 021HAMPC	cyclin G	1382644 IMAGE Consortium	AI036489
7531 D	1 021HAMPC	imprinted and ancient	1162926 IMAGE Consortium	AA770890
7532 D	3 021HAMPC	cathepsin E	1247224 IMAGE Consortium	AA839435
7533 D	5 021HAMPC	endothelial differentiation, G-protein-coupled receptor 6	1345422 IMAGE Consortium	AI158682
7534 D	7 021HAMPC	protein tyrosine phosphatase, receptor type, S	1328214 IMAGE Consortium	AA915604
7535 D	9 021HAMPC	epoxide hydrolase 2, cytoplasmic	1382381 IMAGE Consortium	AI046504
7536 D	11 021HAMPC	septin 9	1382655 IMAGE Consortium	AI036505
7537 F	1 021HAMPC	methyltransferase-like 1 (S. cerevisiae)	1162952 IMAGE Consortium	AA770911



7538 F	3 021HAMPC	SA rat hypertension-associated homolog	1230605 IMAGE Consortium	AA879869
7539 F	5 021HAMPC	carboxyl ester lipase	1332400 IMAGE Consortium	AI876997
7540 F	7 021HAMPC	katanin p60 (ATPase-containing) subunit A1	1331956 IMAGE Consortium	AA958649
7541 F	9 021HAMPC	peptidyl arginine deiminase, type IV	1380273 IMAGE Consortium	AI060554
7542 F	11 021HAMPC	histocompatibility 2, class II, locus DMa	1382670 IMAGE Consortium	AI036062
7543 H	1 021HAMPC	troponin T3, skeletal, fast	1162963 IMAGE Consortium	AA770924
7544 H	3 021HAMPC	hematopoietic cell signal transducer	1243813 IMAGE Consortium	AA822679
7545 H	5 021HAMPC	RAR-related orphan receptor gamma	1331769 IMAGE Consortium	AI181104
7546 H	7 021HAMPC	germ cell-less homolog (Drosophila)	1345749 IMAGE Consortium	AI154560
7547 H	9 021HAMPC	parathyroid hormone receptor	1382420 IMAGE Consortium	AI046522
7548 H	11 021HAMPC	ESTs, Moderately similar to AF170316_1 glycoprotein IIb [M.musculus]	1366675 IMAGE Consortium	AI021345
7549 B	1 021OAMPD	ribosomal protein L41	1382985 IMAGE Consortium	AI120332
7550 B	3 021OAMPD	Rho-associated coiled-coil forming kinase 2	1396474 IMAGE Consortium	AI882290
7551 B	5 021OAMPD	ESTs, Weakly similar to COMP_RAT CARTILAGE OLIGOMERIC MATRIX PROTEIN PRECURSOR [R.norvegicus]	1383744 IMAGE Consortium	AI180914
7552 B	7 021OAMPD	epidermal growth factor receptor pathway substrate 15, related sequence	1480139 IMAGE Consortium	AI156265
7553 B	9 021OAMPD	filaggrin	1429608 IMAGE Consortium	AI877033
7554 B	11 021OAMPD	N-myc downstream regulated 1	330628 IMAGE Consortium	W14224
7555 D	1 021OAMPD	solute carrier family 34 (sodium phosphate), member 2	1383027 IMAGE Consortium	AI120403
7556 D	3 021OAMPD	syndecan 4	1383401 IMAGE Consortium	AI120899
7557 D	5 021OAMPD	fumarylacetoacetate hydrolase	1383759 IMAGE Consortium	AI180932
7558 D	7 021OAMPD	ATP-binding cassette, sub-family F (GCN20), member 1	1493967 IMAGE Consortium	AI156605
7559 D	9 021OAMPD	discs, large homolog 1 (Drosophila)	1430499 IMAGE Consortium	AI182816
7560 D	11 021OAMPD	kinesin-associated protein 3	331395 IMAGE Consortium	W14953
7561 F	1 021OAMPD	trans-acting transcription factor 3	1399769 IMAGE Consortium	AI115748
7562 F	3 021OAMPD	lysozyme	1382758 IMAGE Consortium	AI036879
7563 F	5 021OAMPD	protein kinase, AMP-activated, gamma 1 non-catalytic subunit	1383781 IMAGE Consortium	AI180954
7564 F	7 021OAMPD			
7565 F	9 021OAMPD	acetyl-Coenzyme A dehydrogenase, short chain	1447896 IMAGE Consortium	AI159342
7566 F	11 021OAMPD	high mobility group nucleosomal binding domain 1	331716 IMAGE Consortium	W14418
7567 H	1 021OAMPD	Pim-1 associated protein	1383157 IMAGE Consortium	AI120655
7568 H	3 021OAMPD	serum/glucocorticoid regulated kinase	1383570 IMAGE Consortium	AI120968
7569 H	5 021OAMPD	calcium binding protein Kip 2	1447282 IMAGE Consortium	AI158564
7570 H	7 021OAMPD	period homolog (Drosophila)	1434438 IMAGE Consortium	AA866683
7571 H	9 021OAMPD	fatty acid Coenzyme A ligase, long chain 2	1495812 IMAGE Consortium	AI789976
7572 H	11 021OAMPD	Mpv17 transgene, kidney disease mutant	332371 IMAGE Consortium	W08716
7573 B	1 021VAMPE	brain expressed X-linked 2	1852387 IMAGE Consortium	AI256623
7574 B	3 021VAMPE	GLI-Kruppel family member GLI2	1851402 IMAGE Consortium	AI194364
7575 B	5 021VAMPE	ESTs {IMAGE:334906}	334906 IMAGE Consortium	W17831

7576 B	7 021VAMPE	thioether S-methyltransferase	334658 IMAGE Consortium	W18463
7577 B	9 021VAMPE	guanylate cyclase activator 2 (guanylin 2, intestinal, heatstable)	333377 IMAGE Consortium	W15890
7578 B	11 021VAMPE	ESTs, Moderately similar to motor protein [H.sapiens] {IMAGE:336021}	336021 IMAGE Consortium	W36382
7579 D	1 021VAMPE	lysophospholipase 2	1749789 IMAGE Consortium	AI048448
7580 D	3 021VAMPE	ESTs, Highly similar to unnamed protein product [M.musculus]	1746549 IMAGE Consortium	AI037719
7581 D	5 021VAMPE	integrin beta 4	332870 IMAGE Consortium	W15610
7582 D	7 021VAMPE	glucosidase, alpha, acid	335005 IMAGE Consortium	W17911
7583 D	9 021VAMPE	thioredoxin-like 2	334167 IMAGE Consortium	W16243
7584 D	11 021VAMPE	cytochrome P450, 2d9	335442 IMAGE Consortium	AI893895
7585 F	1 021VAMPE	transition protein 1	1746201 IMAGE Consortium	AI037649
7586 F	3 021VAMPE	tumor protein D52-like 1	1495918 IMAGE Consortium	AI790090
7587 F	5 021VAMPE	heat shock protein, 84 kDa 1	332999 IMAGE Consortium	W08790
7588 F	7 021VAMPE	carnitine palmitoyltransferase 1, liver	335258 IMAGE Consortium	W17627
7589 F	9 021VAMPE	Ena-vasodilator stimulated phosphoprotein	334941 IMAGE Consortium	W37035
7590 F	11 021VAMPE	myc box dependent interacting protein 1	335797 IMAGE Consortium	W34122
7591 H	1 021VAMPE	pleiotrophin	1749974 IMAGE Consortium	AI047160
7592 H	3 021VAMPE	ESTs, Weakly similar to HYPOTHETICAL 103.9 KD PROTEIN ZK370.3 IN CHROMOSOME III [Caenorhabditis elegans] {IMAGE:333511}	333511 IMAGE Consortium	W33982
7593 H	5 021VAMPE	glycosylphosphatidylinositol specific phospholipase D1	333314 IMAGE Consortium	AI324398
7594 H	7 021VAMPE	carbonyl reductase 2	334479 IMAGE Consortium	W36635
7595 H	9 021VAMPE	calpain 2	333504 IMAGE Consortium	AI391241
7596 H	11 021VAMPE	ESTs	336457 IMAGE Consortium	AI893899
7597 B	1 0212AMPF	RGD containing protein	335141 IMAGE Consortium	AA052411
7598 B	3 0212AMPF	actinin alpha 2	335781 IMAGE Consortium	W34282
7599 B	5 0212AMPF	plakophilin 1	349510 IMAGE Consortium	W30104
7600 B	7 0212AMPF	IMAGE EST {IMAGE:367765}	367765 IMAGE Consortium	W53959
7601 B	9 0212AMPF	ATP-binding cassette, sub-family F (GCN20), member 3	368660 IMAGE Consortium	W56930
7602 B	11 0212AMPF	Mus musculus apoptosis activator Mtd (Mtd) mRNA, complete cds {IMAGE:386559}	386559 IMAGE Consortium	W65005
7603 D	1 0212AMPF			
7604 D	3 0212AMPF	brain expressed X-linked 3	335816 IMAGE Consortium	W34128
7605 D	5 0212AMPF	programmed cell death 2	350139 IMAGE Consortium	W34637
7606 D	7 0212AMPF	MARCKS-like protein	367133 IMAGE Consortium	W54626
7607 D	9 0212AMPF	split hand/foot deleted gene 1	373019 IMAGE Consortium	W59554
7608 D	11 0212AMPF	lysosomal-associated protein transmembrane 4A	374302 IMAGE Consortium	W65214
7609 F	1 0212AMPF	eukaryotic translation initiation factor 3, subunit 8 (110 kDa)	335310 IMAGE Consortium	W17453
7610 F	3 0212AMPF	t-complex testis expressed 1	335845 IMAGE Consortium	W34143
7611 F	5 0212AMPF	ninein	350211 IMAGE Consortium	W35926

7612 F	7 0212AMPF	glutathione S-transferase, theta 1	367583 IMAGE Consortium	W54467
7613 F	9 0212AMPF	nuclear protein GANP	367240 IMAGE Consortium	AI427376
7614 F	11 0212AMPF	endothelial PAS domain protein 1	386658 IMAGE Consortium	W64505
7615 H	1 0212AMPF	actin related protein 2/3 complex, subunit 1A (41 kDa)	335406 IMAGE Consortium	W17709
7616 H	3 0212AMPF	ESTs, Highly similar to LMG2_MOUSE LAMININ GAMMA-2 CHAIN PRECURSOR [M.musculus]	335886 IMAGE Consortium	AI390151
7617 H	5 0212AMPF	Friedreich ataxia	353108 IMAGE Consortium	W36030
7618 H	7 0212AMPF	serine hydroxymethyl transferase 1 (soluble)	368208 IMAGE Consortium	W53917
7619 H	9 0212AMPF	DNA segment, Chr 17, ERATO Doi 592, expressed	368082 IMAGE Consortium	W53680
7620 H	11 0212AMPF	tubulin alpha 2	387455 IMAGE Consortium	AI893237
7621 B	1 0211AJ7N	Internal_Control_E	Internal_Control_E	Controls
7622 B	3 0211AJ7N	Internal_Control_F	Internal_Control_F	Controls
7623 B	5 0211AJ7N	Internal_Control_G	Internal_Control_G	Controls
7624 B	7 0211AJ7N	Internal_Control_H	Internal_Control_H	Controls
7625 B	9 0211AJ7N	Internal_Control_I	Internal_Control_I	Controls
7626 B	11 0211AJ7N	Internal_Control_J	Internal_Control_J	Controls
7627 D	1 0211AJ7N	Internal_Control_K	Internal_Control_K	Controls
7628 D	3 0211AJ7N	Internal_Control_L	Internal_Control_L	Controls
7629 D	5 0211AJ7N	Internal_Control_M	Internal_Control_M	Controls
7630 D	7 0211AJ7N	Internal_Control_N	Internal_Control_N	Controls
7631 D	9 0211AJ7N	Internal_Control_O	Internal_Control_O	Controls
7632 D	11 0211AJ7N	Internal_Control_P	Internal_Control_P	Controls
7633 F	1 0211AJ7N	Internal_Control_Q	Internal_Control_Q	Controls
7634 F	3 0211AJ7N	Internal_Control_R	Internal_Control_R	Controls
7635 F	5 0211AJ7N	Internal_Control_S	Internal_Control_S	Controls
7636 F	7 0211AJ7N	Internal_Control_T	Internal_Control_T	Controls
7637 F	9 0211AJ7N	Internal_Control_U	Internal_Control_U	Controls
7638 F	11 0211AJ7N	Internal_Control_V	Internal_Control_V	Controls
7639 H	1 0211AJ7N	Internal_Control_W	Internal_Control_W	Controls
7640 H	3 0211AJ7N	Internal_Control_X	Internal_Control_X	Controls
7641 H	5 0211AJ7N	Internal_Control_Y	Internal_Control_Y	Controls
7642 H	7 0211AJ7N	Internal_Control_Z	Internal_Control_Z	Controls
7643 H	9 0211AJ7N	Internal_Control_AA	Internal_Control_AA	Controls
7644 H	11 0211AJ7N	Internal_Control_AB	Internal_Control_AB	Controls
7651 B	2 0217AJ7T	Control: Ratio 1:3 (Cy3:Cy5)	Ratio_A	Controls
7652 B	4 0217AJ7T	Control: Ratio 3:1 (Cy3:Cy5)	Ratio_B	Controls
7653 B	6 0217AJ7T	Control: Ratio 1:10 (Cy3:Cy5)	Ratio_C	Controls
7654 B	8 0217AJ7T	Control: Ratio 10:1 (Cy3:Cy5)	Ratio_D	Controls
7655 B	10 0217AJ7T	Control: Ratio 1:25 (Cy3:Cy5)	Ratio_E	Controls
7656 B	12 0217AJ7T	Control: Ratio 25:1 (Cy3:Cy5)	Ratio_F	Controls
7657 D	2 0217AJ7T	Control: Sensitivity 2pg	Sensitivity_A	Controls
7658 D	4 0217AJ7T	Control: Sensitivity 20pg	Sensitivity_B	Controls

7659 D	6 0217AJ7T	Control: Sensitivity 200pg	Sensitivity_C	Controls	
7660 D	8 0217AJ7T	Control: Sensitivity 2000pg	Sensitivity_D	Controls	
7661 D	10 0217AJ7T	Control: Buffer Only	Buffer_Only_Control	Controls	
7662 D	12 0217AJ7T	Control: Buffer Only	Buffer_Only_Control	Controls	
7663 F	2 0217AJ7T	Control: Internal	Internal_Control_A	Controls	
7664 F	4 0217AJ7T	Control: Buffer Only	Buffer_Only_Control	Controls	
7665 F	6 0217AJ7T	Control: Internal	Internal_Control_B	Controls	
7666 F	8 0217AJ7T	Control: Buffer Only	Buffer_Only_Control	Controls	
7667 F	10 0217AJ7T	Control: Internal	Internal_Control_C	Controls	
7668 F	12 0217AJ7T	Control: Internal	Internal_Control_D	Controls	
7669 H	2 0217AJ7T	Control: Complex Target (Mus musculus)	RNA_Quality_MM	Controls	
7670 H	4 0217AJ7T	Control: Complex Target (Mus musculus)	RNA_Quality_MM	Controls	
7671 H	6 0217AJ7T	Control: Complex Target (Mus musculus)	RNA_Quality_MM	Controls	
7672 H	8 0217AJ7T	Control: Housekeeping (Alpha Tubulin)	Housekeeping_B	Controls	
7673 H	10 0217AJ7T	Control: Housekeeping (Ribosomal S9)	Housekeeping_A	Controls	
7674 H	12 0217AJ7T	Control: Housekeeping (23kD HBP)	Housekeeping_C	Controls	
7675 B	2 021JAMMK	heparan sulfate-N-deacetylase/N-sulfotransferase	303720	IMAGE Consortium	W09271
7676 B	4 021JAMMK	ESTs, Highly similar to TEB4 protein [H.sapiens]	315121	IMAGE Consortium	AI605705
7677 B	6 021JAMMK	ESTs, Weakly similar to SECIS binding protein 2 [R.norvegicus]	315651	IMAGE Consortium	W09948
7678 B	8 021JAMMK	ESTs	426351	IMAGE Consortium	AA003183
7679 B	10 021JAMMK	ESTs	437736	IMAGE Consortium	AA007866
7680 B	12 021JAMMK	FXYD domain-containing ion transport regulator 1	426150	IMAGE Consortium	AA002987
7681 D	2 021JAMMK	ESTs	313359	IMAGE Consortium	AI605602
7682 D	4 021JAMMK	myelin basic protein	315274	IMAGE Consortium	AI605718
7683 D	6 021JAMMK	ESTs	315804	IMAGE Consortium	AI605757
7684 D	8 021JAMMK	eyes absent 2 homolog (Drosophila)	419144	IMAGE Consortium	W88004
7685 D	10 021JAMMK	potassium voltage-gated channel, subfamily Q, member 2	438522	IMAGE Consortium	AA051041
7686 D	12 021JAMMK	ESTs	539813	IMAGE Consortium	AA123481
7687 F	2 021JAMMK	small proline-rich protein 2A	314112	IMAGE Consortium	AI605648
7688 F	4 021JAMMK	inhibin alpha	314568	IMAGE Consortium	W10710
7689 F	6 021JAMMK	EST	315857	IMAGE Consortium	AI414825
7690 F	8 021JAMMK	bone morphogenetic protein 1	425827	IMAGE Consortium	AA000813
7691 F	10 021JAMMK	ESTs	438618	IMAGE Consortium	AA051010
7692 F	12 021JAMMK	ESTs, Highly similar to hypothetical protein [H.sapiens]	539830	IMAGE Consortium	AA119991
7693 H	2 021JAMMK	ESTs	314209	IMAGE Consortium	AI605650
7694 H	4 021JAMMK	S100 calcium binding protein A5	315361	IMAGE Consortium	AI327447
7695 H	6 021JAMMK	ESTs, Weakly similar to similar to tyrosyl-tRNA synthetase. [C.elegans]	317387	IMAGE Consortium	AI536305
7696 H	8 021JAMMK	hyaluronic acid-binding protein 2	426700	IMAGE Consortium	AA002504
7697 H	10 021JAMMK	ESTs	421377	IMAGE Consortium	W97017
7698 H	12 021JAMMK	E4F transcription factor 1	539903	IMAGE Consortium	AI324466

7699 B	2 021QAMML	retinoblastoma binding protein 7	540283 IMAGE Consortium	AI550560
7700 B	4 021QAMML	guanine nucleotide binding protein, alpha 11	540744 IMAGE Consortium	AA162211
7701 B	6 021QAMML	ESTs	539773 IMAGE Consortium	AI550277
7702 B	8 021QAMML	ESTs	539268 IMAGE Consortium	AA116688
7703 B	10 021QAMML	ESTs, Weakly similar to C33G8.2 [C.elegans]	539407 IMAGE Consortium	AA123134
7704 B	12 021QAMML	ESTs	538940 IMAGE Consortium	AA117283
7705 D	2 021QAMML	ESTs	540358 IMAGE Consortium	AA155138
7706 D	4 021QAMML	ESTs	540816 IMAGE Consortium	AA162226
7707 D	6 021QAMML	cathepsin Z	539055 IMAGE Consortium	AI892501
7708 D	8 021QAMML	ESTs	539276 IMAGE Consortium	AA117449
7709 D	10 021QAMML	Mus musculus Nit protein 2 (Nit2) mRNA, complete cds	539428 IMAGE Consortium	AA123136
7710 D	12 021QAMML	ESTs, Highly similar to ZINC FINGER PROTEIN 43 [Homo sapiens]	538942 IMAGE Consortium	AI892497
7711 F	2 021QAMML	ESTs	540403 IMAGE Consortium	AA155243
7712 F	4 021QAMML	ESTs	539443 IMAGE Consortium	AA117388
7713 F	6 021QAMML	ESTs	539087 IMAGE Consortium	AA117355
7714 F	8 021QAMML	ESTs, Weakly similar to CBFA_MOUSE CCAAT-BINDING TRANSCRIPTION FACTOR SUBUNIT A [M.musculus]	539291 IMAGE Consortium	AA116833
7715 F	10 021QAMML	cerberus 1 homolog (Xenopus laevis)	538769 IMAGE Consortium	AA120122
7716 F	12 021QAMML	transcription factor A, mitochondrial	539693 IMAGE Consortium	AI324462
7717 H	2 021QAMML	ESTs	540459 IMAGE Consortium	AA162023
7718 H	4 021QAMML	ESTs	539512 IMAGE Consortium	AA153179
7719 H	6 021QAMML	ESTs, Moderately similar to HYPOTHETICAL 90.0 KD PROTEIN T20B12.1 IN CHROMOSOME III [Caenorhabditis elegans]	539103 IMAGE Consortium	AA116405
7720 H	8 021QAMML	progesterone receptor membrane component	539300 IMAGE Consortium	AA117455
7721 H	10 021QAMML	reticulocalbin	539525 IMAGE Consortium	AI550245
7722 H	12 021QAMML			
7723 B	2 021XAMMM	mutS homolog 6 (E. coli)	541252 IMAGE Consortium	AI324481
7724 B	4 021XAMMM	ESTs	540785 IMAGE Consortium	AI550587
7725 B	6 021XAMMM	thyrotropin releasing hormone	559851 IMAGE Consortium	AA208912
7726 B	8 021XAMMM	ESTs	805917 IMAGE Consortium	AA399879
7727 B	10 021XAMMM	ESTs	560224 IMAGE Consortium	AA210541
7728 B	12 021XAMMM	DNA segment, Chr 6, Wayne State University 113, expressed	560375 IMAGE Consortium	AA210558
7729 D	2 021XAMMM	CDC28 protein kinase 1	540561 IMAGE Consortium	AA162721
7730 D	4 021XAMMM	ESTs	540790 IMAGE Consortium	AA154636
7731 D	6 021XAMMM	ESTs	559861 IMAGE Consortium	AA211956
7732 D	8 021XAMMM	ESTs, Weakly similar to weak similarity to Bacillus stearothermophilus 30S ribosomal protein S21 [C.elegans]	805946 IMAGE Consortium	AA399907
7733 D	10 021XAMMM	ESTs	560244 IMAGE Consortium	AA210228

7734 D	12 021XAMMM			
7735 F	2 021XAMMM	pre-B-cell colony-enhancing factor	541305 IMAGE Consortium	AA162217
7736 F	4 021XAMMM	ESTs, Moderately similar to TUMOR-ASSOCIATED ANTIGEN L6 HOMOLOG [Mesocricetus auratus]	540848 IMAGE Consortium	AI550591
7737 F	6 021XAMMM			
7738 F	8 021XAMMM	pyruvate dehydrogenase 2	805980 IMAGE Consortium	AA399773
7739 F	10 021XAMMM	polyglutamine binding protein 1	560289 IMAGE Consortium	AA210238
7740 F	12 021XAMMM	platelet derived growth factor receptor, alpha polypeptide	560391 IMAGE Consortium	AA210424
7741 H	2 021XAMMM	ribosomal protein S24	540616 IMAGE Consortium	AI892534
7742 H	4 021XAMMM	translocase of inner mitochondrial membrane 8 homolog b (yeast)	540886 IMAGE Consortium	AI550596
7743 H	6 021XAMMM	stromal cell derived factor 4	560002 IMAGE Consortium	AA265136
7744 H	8 021XAMMM	H2-K region expressed gene 4	805994 IMAGE Consortium	AA399940
7745 H	10 021XAMMM	ESTs	560295 IMAGE Consortium	AA210240
7746 H	12 021XAMMM	ESTs, Highly similar to DEOXYHYPUSINE SYNTHASE [Homo sapiens]	560396 IMAGE Consortium	AA210560
7747 B	2 0214AMMN	ESTs	560584 IMAGE Consortium	AA212102
7748 B	4 0214AMMN	growth hormone releasing hormone	806117 IMAGE Consortium	AA432501
7749 B	6 0214AMMN	ESTs	822755 IMAGE Consortium	AI549676
7750 B	8 0214AMMN	DNA segment, Chr 15, ERATO Doi 405, expressed	865061 IMAGE Consortium	AI614740
7751 B	10 0214AMMN	ESTs	822463 IMAGE Consortium	AI549644
7752 B	12 0214AMMN	f-box and WD-40 domain protein 2	821881 IMAGE Consortium	AA413165
7753 D	2 0214AMMN	ESTs, Moderately similar to hypothetical protein [H.sapiens]	560781 IMAGE Consortium	AA212057
7754 D	4 0214AMMN	gap junction membrane channel protein beta 2	806143 IMAGE Consortium	AI597531
7755 D	6 0214AMMN	reduced expression 2	822780 IMAGE Consortium	AA474825
7756 D	8 0214AMMN	peroxiredoxin 1	865156 IMAGE Consortium	AA450452
7757 D	10 0214AMMN	ESTs	822478 IMAGE Consortium	AA474160
7758 D	12 0214AMMN	ESTs	821937 IMAGE Consortium	AA474047
7759 F	2 0214AMMN	forkhead box A2	560873 IMAGE Consortium	AA212144
7760 F	4 0214AMMN	programmed cell death 10	822588 IMAGE Consortium	AI549654
7761 F	6 0214AMMN	adenosine A2b receptor	822833 IMAGE Consortium	AI549683
7762 F	8 0214AMMN	Mus musculus mRNA for a stretch regulated skeletal muscle protein (Usmg3 gene)	865225 IMAGE Consortium	AA450767
7763 F	10 0214AMMN	glutamate oxaloacetate transaminase 2, mitochondrial	865300 IMAGE Consortium	AA450847
7764 F	12 0214AMMN	pinin	821962 IMAGE Consortium	AA474053
7765 H	2 0214AMMN	eukaryotic translation elongation factor 1 alpha 1	560899 IMAGE Consortium	AA212150
7766 H	4 0214AMMN	polo-like kinase homolog, (Drosophila)	822624 IMAGE Consortium	AA474228
7767 H	6 0214AMMN	hypothetical protein	822889 IMAGE Consortium	AA438130
7768 H	8 0214AMMN	peroxiredoxin 5	865299 IMAGE Consortium	AA450855
7769 H	10 0214AMMN	predicted gene ICRFP703B1614Q5.3	865396 IMAGE Consortium	AA444590
7770 H	12 0214AMMN	core binding factor beta	821338 IMAGE Consortium	AA437837
7771 B	2 021BAMMO	platelet-activating factor acetylhydrolase, isoform 1b,	821411 IMAGE Consortium	AA438069

		alpha2 subunit		
7772 B	4 021BAMMO	Bcl2-associated X protein	821740 IMAGE Consortium	AA413214
7773 B	6 021BAMMO	Mus musculus brain cDNA, clone MNCb-0663, similar to Mus musculus augmenter of liver regeneration (Alr) mRNA	865066 IMAGE Consortium	AA469648
7774 B	8 021BAMMO	ESTs, Highly similar to Unknown [H.sapiens]	865158 IMAGE Consortium	AA450453
7775 B	10 021BAMMO	ESTs, Highly similar to unnamed protein product [H.sapiens]	865211 IMAGE Consortium	AA450533
7776 B	12 021BAMMO	transcriptional intermediary factor 1, beta	865281 IMAGE Consortium	AA450841
7777 D	2 021BAMMO	ESTs, Highly similar to GTPASE-ACTIVATING PROTEIN [Homo sapiens]	806343 IMAGE Consortium	AA432613
7778 D	4 021BAMMO	widely-interspaced zinc finger motifs	806443 IMAGE Consortium	AA470287
7779 D	6 021BAMMO	RAD23b homolog (S. cerevisiae)	865071 IMAGE Consortium	AA469658
7780 D	8 021BAMMO	splicing factor, arginine/serine-rich 2 (SC-35)	865160 IMAGE Consortium	AA450454
7781 D	10 021BAMMO	proteasome (prosome, macropain) 26S subunit, non-ATPase, 13	865217 IMAGE Consortium	AA450536
7782 D	12 021BAMMO	general transcription factor IIH, polypeptide 4	865289 IMAGE Consortium	AA450843
7783 F	2 021BAMMO	ESTs, Highly similar to PSD5_HUMAN 26S PROTEASOME SUBUNIT S5B [H.sapiens]	821658 IMAGE Consortium	AI552358
7784 F	4 021BAMMO	karyopherin (importin) alpha 2	821786 IMAGE Consortium	AI549610
7785 F	6 021BAMMO	tumor rejection antigen gp96	865089 IMAGE Consortium	AA469667
7786 F	8 021BAMMO	RNA binding motif protein 3	865166 IMAGE Consortium	AA450457
7787 F	10 021BAMMO	protein kinase C substrate 80K-H	865221 IMAGE Consortium	AA450538
7788 F	12 021BAMMO	ESTs, Highly similar to KIAA0645 protein [H.sapiens]	865305 IMAGE Consortium	AA450857
7789 H	2 021BAMMO	ribosomal protein L13a	821676 IMAGE Consortium	AI552361
7790 H	4 021BAMMO	Mus musculus brain cDNA, clone MNCb-0783	821803 IMAGE Consortium	AI549614
7791 H	6 021BAMMO	ESTs, Highly similar to HSPC021 [H.sapiens]	865109 IMAGE Consortium	AA469722
7792 H	8 021BAMMO	ESTs, Weakly similar to CG12404 gene product [D.melanogaster]	865171 IMAGE Consortium	AA450467
7793 H	10 021BAMMO	ESTs	865229 IMAGE Consortium	AA450768
7794 H	12 021BAMMO	GPI-anchored membrane protein 1	865312 IMAGE Consortium	AA450851
7795 B	2 021IAMMP	ESTs	865352 IMAGE Consortium	AI643193
7796 B	4 021IAMMP	ESTs, Moderately similar to DRPLA [H.sapiens]	865481 IMAGE Consortium	AI643212
7797 B	6 021IAMMP	ESTs, Moderately similar to unnamed protein product [H.sapiens]	865863 IMAGE Consortium	AI552950
7798 B	8 021IAMMP	ESTs	905860 IMAGE Consortium	AA529422
7799 B	10 021IAMMP	ornithine aminotransferase	907047 IMAGE Consortium	AI551961
7800 B	12 021IAMMP	ESTs, Highly similar to TRANSLOCON-ASSOCIATED PROTEIN, GAMMA SUBUNIT [Rattus norvegicus]	906703 IMAGE Consortium	AA522311
7801 D	2 021IAMMP	heat shock 70kD protein 5 (glucose-regulated protein, 78kD)	865361 IMAGE Consortium	AA450922
7802 D	4 021IAMMP	ESTs, Highly similar to unnamed protein product [H.sapiens]	865562 IMAGE Consortium	AI552932
7803 D	6 021IAMMP	ESTs	865875 IMAGE Consortium	AI552954

7804 D	8 021IAMMP	ESTs	905914 IMAGE Consortium	AA529485
7805 D	10 021IAMMP	ESTs, Moderately similar to KIAA0875 protein [H.sapiens]	907211 IMAGE Consortium	AI551973
7806 D	12 021IAMMP	solute carrier family 7 (cationic amino acid transporter, y+ system), member 7	907469 IMAGE Consortium	AA588984
7807 F	2 021IAMMP	glucose regulated protein, 58 kDa	865364 IMAGE Consortium	AA450918
7808 F	4 021IAMMP	ESTs	865629 IMAGE Consortium	AA445075
7809 F	6 021IAMMP	ESTs, Weakly similar to BRCA1-associated RING domain protein [M.musculus]	906158 IMAGE Consortium	AA529326
7810 F	8 021IAMMP	ESTs	866000 IMAGE Consortium	AA473497
7811 F	10 021IAMMP	ESTs	906511 IMAGE Consortium	AA522221
7812 F	12 021IAMMP	ESTs	907474 IMAGE Consortium	AA588976
7813 H	2 021IAMMP	heat shock protein, 60 kDa	865378 IMAGE Consortium	AA444576
7814 H	4 021IAMMP	ribosomal protein L6	865653 IMAGE Consortium	AI552938
7815 H	6 021IAMMP	stromal cell derived factor receptor 2	906232 IMAGE Consortium	AA529439
7816 H	8 021IAMMP	Mus musculus Creb-rp, Tnx and Cyp21 genes for cAMP response element binding protein-related protein, tenascin-X and steroid 21-hydroxylase, partial cds and complete sequences	866044 IMAGE Consortium	AA473564
7817 H	10 021IAMMP	phospholipase C, beta 3	906524 IMAGE Consortium	AA522219
7818 H	12 021IAMMP	ESTs, Highly similar to AF112204_1 Vacuolar proton pump subunit SFD alpha isoform [H.sapiens]	906758 IMAGE Consortium	AA522340
7819 B	2 021PAMMQ	zinc finger protein 54	907601 IMAGE Consortium	AA590535
7820 B	4 021PAMMQ	upstream regulatory element binding protein 1	908767 IMAGE Consortium	AA591500
7821 B	6 021PAMMQ	Mus musculus TCR beta locus from bases 1 to 250611 (section 1 of 3) of the complete sequence	513886 IMAGE Consortium	AA061775
7822 B	8 021PAMMQ	RAS protein activator like 1 (GAP1 like)	514044 IMAGE Consortium	AA062153
7823 B	10 021PAMMQ	RuvB-like protein 1	515287 IMAGE Consortium	AA064425
7824 B	12 021PAMMQ	ESTs, Moderately similar to Pro-Pol-dUTPase polyprotein [M.musculus]	907617 IMAGE Consortium	AI551391
7825 D	2 021PAMMQ	homeo box B1	908792 IMAGE Consortium	AA591510
7826 D	4 021PAMMQ	ESTs	514634 IMAGE Consortium	AI595170
7827 D	6 021PAMMQ	ESTs	514069 IMAGE Consortium	AI427888
7828 D	8 021PAMMQ	ESTs	515106 IMAGE Consortium	AI595289
7829 D	10 021PAMMQ	ESTs	515297 IMAGE Consortium	AI595608
7830 D	12 021PAMMQ	ESTs	907660 IMAGE Consortium	AA555657
7831 F	2 021PAMMQ	myosin Ib	514415 IMAGE Consortium	AI595148
7832 F	4 021PAMMQ	acyl-Coenzyme A thioesterase 3, mitochondrial	513931 IMAGE Consortium	AI595099
7833 F	6 021PAMMQ	hypothetical protein	514152 IMAGE Consortium	AI595121
7834 F	8 021PAMMQ	Rhesus blood group-associated C glycoprotein	515131 IMAGE Consortium	AA064433
7835 F	10 021PAMMQ	microtubule associated testis specific serine/threonine protein kinase		
7836 F	12 021PAMMQ	golgi autoantigen, golgin subfamily a, 3	514588 IMAGE Consortium	AA063782
7837 H	2 021PAMMQ	RAN, member RAS oncogene family	906945 IMAGE Consortium	AI551951



7838 H	4 021PAMMQ	ESTs	513707 IMAGE Consortium	AI595228
7839 H	6 021PAMMQ	uroporphyrinogen III synthase	513942 IMAGE Consortium	AA061691
7840 H	8 021PAMMQ	testicular serine protease 2	514204 IMAGE Consortium	AI595126
7841 H	10 021PAMMQ	ESTs	515134 IMAGE Consortium	AI595291
7842 H	12 021PAMMQ	ESTs, Moderately similar to KIAA1211 protein [H.sapiens]	515333 IMAGE Consortium	AI595615
7843 B	2 021WAMMR	ESTs, Highly similar to unnamed protein product [H.sapiens]	515439 IMAGE Consortium	AA064333
7844 B	4 021WAMMR	ESTs, Highly similar to protein serine/threonine phosphatase 4 regulatory subunit 1 [H.sapiens]	514867 IMAGE Consortium	AA064022
7845 B	6 021WAMMR	serine/threonine kinase 22A (spermiogenesis associated)	515563 IMAGE Consortium	AA065467
7846 B	8 021WAMMR	ESTs	515615 IMAGE Consortium	AI595021
7847 B	10 021WAMMR	ESTs	515858 IMAGE Consortium	AI595533
7848 B	12 021WAMMR	kidney androgen regulated protein	517210 IMAGE Consortium	AI892236
7849 D	2 021WAMMR	endothelial monocyte activating polypeptide 2	515462 IMAGE Consortium	AI594998
7850 D	4 021WAMMR	signaling intermediate in Toll pathway-evolutionarily conserved	514929 IMAGE Consortium	AI595271
7851 D	6 021WAMMR	ESTs, Weakly similar to AF167405_1 ADAM 27 [M.musculus]	516394 IMAGE Consortium	AA110044
7852 D	8 021WAMMR	ESTs, Weakly similar to nolp [H.sapiens]	515653 IMAGE Consortium	AI595023
7853 D	10 021WAMMR	Mus musculus mdgl-1 mRNA, complete cds	515861 IMAGE Consortium	AI595535
7854 D	12 021WAMMR	ESTs	517230 IMAGE Consortium	AA062425
7855 F	2 021WAMMR	ATP synthase, H <sup>+</sup> transporting, mitochondrial F0 complex, subunit c (subunit 9), isoform 1	515500 IMAGE Consortium	AA065942
7856 F	4 021WAMMR	ESTs	514947 IMAGE Consortium	AA061196
7857 F	6 021WAMMR	Mouse (clone BALB11N) mRNA, complete cds of unknown function	516004 IMAGE Consortium	AI326539
7858 F	8 021WAMMR	ESTs, Highly similar to RGC-32 [R.norvegicus]	515663 IMAGE Consortium	AA066016
7859 F	10 021WAMMR	ESTs	515894 IMAGE Consortium	AI595537
7860 F	12 021WAMMR	origin recognition complex, subunit 2 homolog (S. cerevisiae)	517253 IMAGE Consortium	AA062361
7861 H	2 021WAMMR	Sin3-associated polypeptide 18	515505 IMAGE Consortium	AI595008
7862 H	4 021WAMMR	cleavage and polyadenylation specific factor 4, 30kD subunit	515000 IMAGE Consortium	AA064044
7863 H	6 021WAMMR	protein phosphatase 2 (formerly 2A), regulatory subunit B" (PR72), alpha isoform and (PR 130), beta isoform	516049 IMAGE Consortium	AA120741
7864 H	8 021WAMMR	proteasome (prosome, macropain) subunit, beta type 5	515685 IMAGE Consortium	AA066023
7865 H	10 021WAMMR	coactivator of 150 kD	515926 IMAGE Consortium	AA062401
7866 H	12 021WAMMR	phosphatidylcholine transfer protein	517288 IMAGE Consortium	AA066072
7867 B	2 0213AMMS	ESTs, Highly similar to NARDILYSIN PRECURSOR [Rattus norvegicus]	517338 IMAGE Consortium	AI892237
7868 B	4 0213AMMS	glycerol kinase	517468 IMAGE Consortium	AA059599
7869 B	6 0213AMMS	SAR1a gene homolog (S. cerevisiae)	517657 IMAGE Consortium	AA066667
7870 B	8 0213AMMS	ESTs	517600 IMAGE Consortium	AA066568

7871 B	10 0213AMMS	ESTs, Highly similar to IF36_HUMAN EUKARYOTIC TRANSLATION INITIATION FACTOR 3 SUBUNIT 6 [H.sapiens]	517785 IMAGE Consortium	AA066651
7872 B	12 0213AMMS	ESTs	519291 IMAGE Consortium	AI597343
7873 D	2 0213AMMS			
7874 D	4 0213AMMS	single Ig IL-1 receptor related protein	517477 IMAGE Consortium	AI597102
7875 D	6 0213AMMS	ESTs, Highly similar to B-MYC TRANSFORMING PROTEIN [Rattus norvegicus]	516935 IMAGE Consortium	AI892231
7876 D	8 0213AMMS	aldolase 1, A isoform	517620 IMAGE Consortium	AI892240
7877 D	10 0213AMMS	gene rich cluster, C10 gene	517797 IMAGE Consortium	AA066684
7878 D	12 0213AMMS	solute carrier family 16 (monocarboxylic acid transporters), member 7	519315 IMAGE Consortium	AA106512
7879 F	2 0213AMMS	Kruppel-like factor 3 (basic)	517397 IMAGE Consortium	AA062455
7880 F	4 0213AMMS	ESTs, Moderately similar to hypothetical protein [H.sapiens]	517486 IMAGE Consortium	AI597101
7881 F	6 0213AMMS	cytochrome c, somatic	517705 IMAGE Consortium	AI892244
7882 F	8 0213AMMS	transcobalamin 2	517628 IMAGE Consortium	AA066473
7883 F	10 0213AMMS			
7884 F	12 0213AMMS	ESTs, Highly similar to A57514 RNA helicase HEL117 - rat [R.norvegicus]	518681 IMAGE Consortium	AA105953
7885 H	2 0213AMMS	DiGeorge syndrome chromosome region 6	517410 IMAGE Consortium	AA062498
7886 H	4 0213AMMS	fragile histidine triad gene	517492 IMAGE Consortium	AI597104
7887 H	6 0213AMMS	Kruppel-like factor 9	517752 IMAGE Consortium	AA066598
7888 H	8 0213AMMS	cytochrome c oxidase, subunit VIc	517641 IMAGE Consortium	AI892241
7889 H	10 0213AMMS	ESTs, Moderately similar to AF231110_1 muscleblind [M.musculus]	519074 IMAGE Consortium	AA106833
7890 H	12 0213AMMS	phosphatidylinositol-4-phosphate 5-kinase, type 1 beta	519411 IMAGE Consortium	AA106993
7891 B	2 021AAMMT	ESTs, Weakly similar to scaffold attachment factor B [R.norvegicus]	518947 IMAGE Consortium	AA106772
7892 B	4 021AAMMT	Mus musculus strain 129/Ola papillary renal cell carcinoma-associated protein (Prcc) mRNA, complete cds	518813 IMAGE Consortium	AA089181
7893 B	6 021AAMMT	ESTs, Highly similar to The KIAA0145 gene product is related to diacylglycerol kinase. [H.sapiens]	519404 IMAGE Consortium	AA106557
7894 B	8 021AAMMT	ESTs	519991 IMAGE Consortium	AA109804
7895 B	10 021AAMMT	hypothetical protein, clone 2-6	520300 IMAGE Consortium	AA110890
7896 B	12 021AAMMT	ESTs, Highly similar to macrophage maturation-associated transcript dd3f protein [H.sapiens]	521097 IMAGE Consortium	AA066818
7897 D	2 021AAMMT	methylmalonyl-Coenzyme A mutase	518969 IMAGE Consortium	AA106902
7898 D	4 021AAMMT	integral membrane protein 2 B	518856 IMAGE Consortium	AA106337
7899 D	6 021AAMMT	ESTs, Weakly similar to AF182426_1 arylacetamide deacetylase [R.norvegicus]	519438 IMAGE Consortium	AI597355
7900 D	8 021AAMMT	mepripin 1 beta	520031 IMAGE Consortium	AI324410
7901 D	10 021AAMMT	galactokinase	520324 IMAGE Consortium	AA110914
7902 D	12 021AAMMT	ESTs, Highly similar to NADH-UBIQUINONE	521117 IMAGE Consortium	AA066803

7903 F	2 021AAMMT	OXIDOREDUCTASE SUBUNIT B14.5B [Bos taurus]	518971 IMAGE Consortium	AI892747
7904 F	4 021AAMMT	glutathione S-transferase, alpha 2 (Yc2)	518880 IMAGE Consortium	AA106360
7905 F	6 021AAMMT	signal recognition particle 14 kDa (homologous Alu RNA-binding protein)	519512 IMAGE Consortium	AA106935
7906 F	8 021AAMMT	ESTs	520082 IMAGE Consortium	AA109835
7907 F	10 021AAMMT	ESTs, Weakly similar to unknown [M.musculus]	520353 IMAGE Consortium	AA109873
7908 F	12 021AAMMT	ESTs, Moderately similar to unnamed protein product [H.sapiens]	520415 IMAGE Consortium	AA105796
7909 H	2 021AAMMT	itchy	518990 IMAGE Consortium	AA106154
7910 H	4 021AAMMT	ESTs, Weakly similar to ACDS_MOUSE ACYL-COA DEHYDROGENASE, SHORT-CHAIN SPECIFIC PRECURSOR [M.musculus]	518883 IMAGE Consortium	AA106365
7911 H	6 021AAMMT	cytochrome P450, 4a14	519535 IMAGE Consortium	
7912 H	8 021AAMMT	Public domain EST	520098 IMAGE Consortium	AA107637
7913 H	10 021AAMMT	ESTs	520355 IMAGE Consortium	AI324412
7914 H	12 021AAMMT	UDP-glucuronosyltransferase 2 family, member 5	521166 IMAGE Consortium	AA066981
7915 B	2 021HAMMU	ESTs, Highly similar to SERB_HUMAN L-3-PHOSPHOSERINE PHOSPHATASE [H.sapiens]	520578 IMAGE Consortium	AI552694
7916 B	4 021HAMMU	peroxisome proliferative activated receptor, gamma, coactivator 1	520904 IMAGE Consortium	AI552710
7917 B	6 021HAMMU	ESTs	442866 IMAGE Consortium	AA014199
7918 B	8 021HAMMU	ESTs	439946 IMAGE Consortium	AA008471
7919 B	10 021HAMMU	t-complex-associated testis expressed 1		
7920 B	12 021HAMMU	ESTs, Highly similar to AF161509_1 HSPC160 [H.sapiens]	522814 IMAGE Consortium	AA087508
7921 D	2 021HAMMU	esterase 10	520613 IMAGE Consortium	AA066534
7922 D	4 021HAMMU	transcription factor like 4	520937 IMAGE Consortium	AA065608
7923 D	6 021HAMMU	alpha-N-acetylglucosaminidase (Sanfilippo disease IIIB)	444344 IMAGE Consortium	AA015216
7924 D	8 021HAMMU	galanin	440230 IMAGE Consortium	AI894042
7925 D	10 021HAMMU	ESTs, Highly similar to unnamed protein product [H.sapiens]	523230 IMAGE Consortium	AI614576
7926 D	12 021HAMMU	ESTs, Weakly similar to PROBABLE PEPTIDYL-TRNA HYDROLASE [Bacillus subtilis]	522843 IMAGE Consortium	AA087651
7927 F	2 021HAMMU	phosphatidylinositol 3-kinase, C2 domain containing, gamma polypeptide	520661 IMAGE Consortium	AA066552
7928 F	4 021HAMMU	hydroxysteroid 17-beta dehydrogenase 1	520995 IMAGE Consortium	AI326554
7929 F	6 021HAMMU	caudal type homeo box 2	444481 IMAGE Consortium	AI894056
7930 F	8 021HAMMU	ESTs, Weakly similar to PROTEIN KINASE SPK1 [Saccharomyces cerevisiae]	441316 IMAGE Consortium	AA009198
7931 F	10 021HAMMU	ESTs	523258 IMAGE Consortium	AI614583
7932 F	12 021HAMMU	four and a half LIM domains 1	522876 IMAGE Consortium	AI326574
7933 H	2 021HAMMU	ESTs	520708 IMAGE Consortium	AA066390
7934 H	4 021HAMMU	lipopolysaccharide inducible C-C chemokine receptor	442765 IMAGE Consortium	AA034646

		related		
7935 H	6 021HAMMU	WW domain-containing protein 3	443932 IMAGE Consortium	AA014158
7936 H	8 021HAMMU	ESTs, Highly similar to unnamed protein product [H.sapiens]	523133 IMAGE Consortium	AI552875
7937 H	10 021HAMMU	selectin, endothelial cell, ligand	522554 IMAGE Consortium	AA067657
7938 H	12 021HAMMU	ESTs, Highly similar to HETEROGENOUS NUCLEAR RIBONUCLEOPROTEIN U [Homo sapiens]	522922 IMAGE Consortium	AI552837
7939 B	2 021OAMMV	kinectin 1	523087 IMAGE Consortium	AA067264
7940 B	4 021OAMMV	potassium channel, subfamily K, member 1	521905 IMAGE Consortium	AI552753
7941 B	6 021OAMMV	ESTs, Highly similar to FIBRILLIN 1 PRECURSOR [Homo sapiens]	522100 IMAGE Consortium	AI386241
7942 B	8 021OAMMV	ESTs, Weakly similar to CG11802 gene product [D.melanogaster]	523199 IMAGE Consortium	AA067107
7943 B	10 021OAMMV	ESTs	523402 IMAGE Consortium	AA080173
7944 B	12 021OAMMV	nuclear factor I/B	522926 IMAGE Consortium	AA065880
7945 D	2 021OAMMV			
7946 D	4 021OAMMV	ESTs	521952 IMAGE Consortium	AA106871
7947 D	6 021OAMMV	apolipoprotein E	521479 IMAGE Consortium	AI325603
7948 D	8 021OAMMV	ESTs, Moderately similar to hepatocellular carcinoma-associated antigen 59 [H.sapiens]	523268 IMAGE Consortium	AA067123
7949 D	10 021OAMMV	CDC-like kinase 2	523408 IMAGE Consortium	AA080256
7950 D	12 021OAMMV	hydroxysteroid dehydrogenase-4, delta<5>-3-beta	522929 IMAGE Consortium	AI552842
7951 F	2 021OAMMV	ribosomal protein L13	522399 IMAGE Consortium	AA067628
7952 F	4 021OAMMV	ESTs, Weakly similar to predicted using Genefinder [C.elegans]	521988 IMAGE Consortium	AA106709
7953 F	6 021OAMMV	acidic ribosomal phosphoprotein PO	521549 IMAGE Consortium	AI386059
7954 F	8 021OAMMV	keratin complex 1, gene 3	523282 IMAGE Consortium	AA079939
7955 F	10 021OAMMV	ESTs, Highly similar to KIAA0560 protein [H.sapiens]	523412 IMAGE Consortium	AA080258
7956 F	12 021OAMMV	glycoprotein galactosyltransferase alpha 1, 3	523683 IMAGE Consortium	AA067151
7957 H	2 021OAMMV	ESTs	521765 IMAGE Consortium	AI427560
7958 H	4 021OAMMV	ESTs, Weakly similar to T08D2.6 [C.elegans]	521133 IMAGE Consortium	AA066915
7959 H	6 021OAMMV	solute carrier family 27 (fatty acid transporter), member 2	521620 IMAGE Consortium	AA066694
7960 H	8 021OAMMV	myotubularin related protein 1	523285 IMAGE Consortium	AI614592
7961 H	10 021OAMMV	ESTs, Highly similar to MAJOR CENTROMERE AUTOANTIGEN B [Mus musculus]	523440 IMAGE Consortium	AI892313
7962 H	12 021OAMMV	hypoxanthine guanine phosphoribosyl transferase	523992 IMAGE Consortium	AA072014
7963 B	2 021VAMMW	ESTs, Highly similar to ER23_HUMAN ER LUMEN PROTEIN RETAINING RECEPTOR 3 [H.sapiens]	524067 IMAGE Consortium	AA066130
7964 B	4 021VAMMW	ESTs, Moderately similar to G01522 acidic 82 kDa protein - human [H.sapiens]	524284 IMAGE Consortium	AI614800
7965 B	6 021VAMMW	tissue inhibitor of metalloproteinase 2	523876 IMAGE Consortium	AA067331
7966 B	8 021VAMMW	syndecan binding protein	524562 IMAGE Consortium	AA068481
7967 B	10 021VAMMW	ESTs	524751 IMAGE Consortium	AA067780

7968 B	12 021VAMMW	ESTs, Highly similar to AF161432_1 HSPC314 [H.sapiens]	524081 IMAGE Consortium	AA066251
7969 D	2 021VAMMW	thioredoxin-like (32kD)	524115 IMAGE Consortium	AA066178
7970 D	4 021VAMMW			
7971 D	6 021VAMMW	Mus musculus mRNA for a stretch regulated skeletal muscle protein (Usmg5 gene)	523934 IMAGE Consortium	AA067380
7972 D	8 021VAMMW			
7973 D	10 021VAMMW	ESTs	524776 IMAGE Consortium	AA068613
7974 D	12 021VAMMW	acetyl-Coenzyme A transporter	524101 IMAGE Consortium	AI614011
7975 F	2 021VAMMW	ESTs, Moderately similar to JNK-binding protein JNKBP1 [M.musculus]	524141 IMAGE Consortium	AA066191
7976 F	4 021VAMMW	ESTs, Highly similar to AF161432_1 HSPC314 [H.sapiens]	524357 IMAGE Consortium	AA065718
7977 F	6 021VAMMW	neural precursor cell expressed, developmentally down-regulated gene 4	523964 IMAGE Consortium	AA067727
7978 F	8 021VAMMW	lithium-sensitive myo-inositol monophosphatase A1	524589 IMAGE Consortium	AA068556
7979 F	10 021VAMMW	ESTs	524812 IMAGE Consortium	AA097002
7980 F	12 021VAMMW	DNA segment, Chr 1, University of California at Los Angeles 3	524108 IMAGE Consortium	AA066256
7981 H	2 021VAMMW	ESTs, Moderately similar to dJ622L5.8.1 [H.sapiens]	524186 IMAGE Consortium	AA066197
7982 H	4 021VAMMW	tumor susceptibility gene 101	524382 IMAGE Consortium	AA065752
7983 H	6 021VAMMW	ESTs, Highly similar to T2DB_HUMAN TRANSCRIPTION INITIATION FACTOR TFIID 18 KD SUBUNIT [H.sapiens]	523974 IMAGE Consortium	AI386247
7984 H	8 021VAMMW	ESTs, Moderately similar to Y101_HUMAN HYPOTHETICAL PROTEIN KIAA0101 [H.sapiens]	524613 IMAGE Consortium	AA068562
7985 H	10 021VAMMW	ESTs	524861 IMAGE Consortium	AI614817
7986 H	12 021VAMMW	RW1 protein	524126 IMAGE Consortium	AA066176
7987 B	2 0212AMMX	alcohol dehydrogenase 3 complex	524271 IMAGE Consortium	AA065690
7988 B	4 0212AMMX	ESTs	524474 IMAGE Consortium	AA067814
7989 B	6 0212AMMX	membrane component, surface marker 1	525061 IMAGE Consortium	AI643097
7990 B	8 0212AMMX	epithelial membrane protein 1	524845 IMAGE Consortium	AA097101
7991 B	10 0212AMMX	cytochrome P450, 2j5	524967 IMAGE Consortium	AA096985
7992 B	12 0212AMMX	ESTs, Weakly similar to DPP4_MOUSE DIPEPTIDYL PEPTIDASE IV [M.musculus]	533418 IMAGE Consortium	AA104524
7993 D	2 0212AMMX	ESTs	524275 IMAGE Consortium	AA067894
7994 D	4 0212AMMX	ESTs	524499 IMAGE Consortium	AA067823
7995 D	6 0212AMMX	ESTs, Weakly similar to membrane glycoprotein CIG30 [M.musculus]	533113 IMAGE Consortium	AI614843
7996 D	8 0212AMMX	ESTs, Moderately similar to dJ187J11.1 [H.sapiens]	524848 IMAGE Consortium	AA088935
7997 D	10 0212AMMX			
7998 D	12 0212AMMX	ESTs, Highly similar to BETA-ADRENERGIC RECEPTOR KINASE 1 [Bos taurus]	534159 IMAGE Consortium	AI386289
7999 F	2 0212AMMX	ESTs, Moderately similar to unknown [H.sapiens]	524293 IMAGE Consortium	AI614805
8000 F	4 0212AMMX	ect2 oncogene	524595 IMAGE Consortium	AA068464

8001 F	6 0212AMMX	pre-mRNA protein kinase	525180 IMAGE Consortium	AA096763
8002 F	8 0212AMMX	ESTs, Moderately similar to zinc-finger protein MCG4 [H.sapiens]	532880 IMAGE Consortium	AI614832
8003 F	10 0212AMMX	SEC8 (S. cerevisiae)	534063 IMAGE Consortium	AA072569
8004 F	12 0212AMMX	peroxisomal membrane protein 3, 35 kDa	534171 IMAGE Consortium	AI326609
8005 H	2 0212AMMX	ESTs, Highly similar to HYPOTHETICAL 31.1 KD PROTEIN B0228.7 IN CHROMOSOME II [Caenorhabditis elegans]	524297 IMAGE Consortium	AA067899
8006 H	4 0212AMMX	SPARC-like 1 (mast9, hev1)	524644 IMAGE Consortium	AA068603
8007 H	6 0212AMMX	translocase of inner mitochondrial membrane 10 homolog (yeast)	525184 IMAGE Consortium	AA096899
8008 H	8 0212AMMX	ESTs, Moderately similar to SAS_HUMAN SARCOMA AMPLIFIED SEQUENCE [H.sapiens]	524877 IMAGE Consortium	AA088966
8009 H	10 0212AMMX	ESTs, Highly similar to PHOSPHOENOLPYRUVATE CARBOXYKINASE, CYTOSOLIC [Rattus norvegicus]	533202 IMAGE Consortium	AI551975
8010 H	12 0212AMMX	ESTs	533455 IMAGE Consortium	AI551995
8011 B	2 0219AMMY	ESTs, Weakly similar to AF165114_1 Borg4 [M.musculus]	534292 IMAGE Consortium	AI552060
8012 B	4 0219AMMY	ESTs, Highly similar to MICROSOMAL SIGNAL PEPTIDASE 21 KD SUBUNIT [Canis familiaris]	550564 IMAGE Consortium	AI386300
8013 B	6 0219AMMY	RAD9 homolog (S. pombe)	536565 IMAGE Consortium	AI325746
8014 B	8 0219AMMY	ATP synthase, H <sup>+</sup> transporting, mitochondrial F1 complex, alpha subunit, isoform 1	535257 IMAGE Consortium	AI892400
8015 B	10 0219AMMY	ESTs, Moderately similar to unnamed protein product [H.sapiens]	535685 IMAGE Consortium	AA106222
8016 B	12 0219AMMY	U2 small nuclear ribonucleoprotein auxiliary factor (U2AF), 35 kDa, related sequence 1	476501 IMAGE Consortium	AA050756
8017 D	2 0219AMMY	ESTs	532892 IMAGE Consortium	AA068951
8018 D	4 0219AMMY	ESTs	550614 IMAGE Consortium	AA097941
8019 D	6 0219AMMY	transglutaminase 2, C polypeptide	536592 IMAGE Consortium	AI892437
8020 D	8 0219AMMY	ESTs	535262 IMAGE Consortium	AI552098
8021 D	10 0219AMMY	nucleoplasmin 3	534981 IMAGE Consortium	AA073445
8022 D	12 0219AMMY	lysyl oxidase-like	478041 IMAGE Consortium	AI322695
8023 F	2 0219AMMY	ESTs, Weakly similar to AF163095_1 HRAS suppressor [M.musculus]	533604 IMAGE Consortium	AI552004
8024 F	4 0219AMMY	EST AA407494	536175 IMAGE Consortium	AA073401
8025 F	6 0219AMMY	heme oxygenase (decycling) 1	536616 IMAGE Consortium	AI324438
8026 F	8 0219AMMY	ESTs, Highly similar to PUTATIVE RECEPTOR PROTEIN [Homo sapiens]	535320 IMAGE Consortium	AI892401
8027 F	10 0219AMMY	DNA segment, human D4S114	535754 IMAGE Consortium	AI325723
8028 F	12 0219AMMY	transforming growth factor, beta 1	478105 IMAGE Consortium	AA049522
8029 H	2 0219AMMY	aminolevulinic acid synthase 1	533794 IMAGE Consortium	AI892379
8030 H	4 0219AMMY	ESTs	536197 IMAGE Consortium	AI549977
8031 H	6 0219AMMY	guanine nucleotide binding protein, alpha stimulating	536637 IMAGE Consortium	AI550047
8032 H	8 0219AMMY	cyclin-dependent kinase 4	535389 IMAGE Consortium	AI892404

8033 H	10 0219AMMY	ESTs, Highly similar to 5-FORMYLTETRAHYDROFOLATE CYCLO-LIGASE [Oryctolagus cuniculus]	535906 IMAGE Consortium	AI325728
8034 H	12 0219AMMY	ESTs, Highly similar to BETA-ARRESTIN 2 [Rattus norvegicus]	478158 IMAGE Consortium	AI322696
8035 B	2 021GAMMZ	Kruppel-like factor 1 (erythroid)	478493 IMAGE Consortium	AI323045
8036 B	4 021GAMMZ	hypermethylated in cancer 1	478844 IMAGE Consortium	AA048805
8037 B	6 021GAMMZ	ESTs	572936 IMAGE Consortium	AA111601
8038 B	8 021GAMMZ	ESTs	571524 IMAGE Consortium	AI587781
8039 B	10 021GAMMZ	ESTs, Moderately similar to AF155115_1 NY-REN-58 antigen [H.sapiens]	480223 IMAGE Consortium	AA058086
8040 B	12 021GAMMZ	ESTs	482303 IMAGE Consortium	AA063757
8041 D	2 021GAMMZ	growth associated protein 43	479228 IMAGE Consortium	AI323051
8042 D	4 021GAMMZ	homeo box A5	475369 IMAGE Consortium	AI322633
8043 D	6 021GAMMZ	ESTs	573816 IMAGE Consortium	AI530459
8044 D	8 021GAMMZ	ESTs	573038 IMAGE Consortium	AI587876
8045 D	10 021GAMMZ	retinol binding protein 2, cellular	481058 IMAGE Consortium	AI385457
8046 D	12 021GAMMZ	ESTs, Moderately similar to PTERIN-4-ALPHA-CARBINOLAMINE DEHYDRATASE [Homo sapiens; Rattus norvegicus; Mus musculus]	480924 IMAGE Consortium	AA064249
8047 F	2 021GAMMZ	synuclein, alpha	477819 IMAGE Consortium	AI510034
8048 F	4 021GAMMZ	G protein-coupled receptor 19	475460 IMAGE Consortium	AA041917
8049 F	6 021GAMMZ	ESTs, Weakly similar to S28183 aryl sulfotransferase [M.musculus]	492984 IMAGE Consortium	AI595193
8050 F	8 021GAMMZ	ESTs	572363 IMAGE Consortium	AI587834
8051 F	10 021GAMMZ	thymus cell antigen 1, theta	480473 IMAGE Consortium	AA064119
8052 F	12 021GAMMZ	neutrophilic granule protein	481675 IMAGE Consortium	AA060282
8053 H	2 021GAMMZ	ESTs	477876 IMAGE Consortium	AA048648
8054 H	4 021GAMMZ	ESTs, Highly similar to KIAA0622 protein [H.sapiens]	476216 IMAGE Consortium	AI509974
8055 H	6 021GAMMZ	ESTs	571294 IMAGE Consortium	AI587759
8056 H	8 021GAMMZ	ESTs, Weakly similar to /prediction	572479 IMAGE Consortium	AI587839
8057 H	10 021GAMMZ	nucleoside diphosphate kinase (nm23-M4)	480596 IMAGE Consortium	AI892129
8058 H	12 021GAMMZ	ESTs, Weakly similar to KRUC_HUMAN KERATIN, ULTRA HIGH-SULFUR MATRIX PROTEIN [H.sapiens]	481703 IMAGE Consortium	AI594897
8059 B	2 021GAMNO	ESTs	483391 IMAGE Consortium	AI552476
8060 B	4 021GAMNO	DEAD (aspartate-glutamate-alanine-aspartate) box polypeptide, Y chromosome	573643 IMAGE Consortium	AI510645
8061 B	6 021GAMNO	ESTs	573937 IMAGE Consortium	AA119395
8062 B	8 021GAMNO	eosinophil-associated ribonuclease 3	575740 IMAGE Consortium	AA124831
8063 B	10 021GAMNO	cytochrome P450, subfamily IVF, polypeptide 14 (leukotriene B4 omega hydroxylase)	987518 IMAGE Consortium	AA793220
8064 B	12 021GAMNO	hemopoietic cell phosphatase	1001376 IMAGE Consortium	AA771630
8065 D	2 021GAMNO	lymphoblastomic leukemia	482694 IMAGE Consortium	AI323106

8066 D	4 021GAMN0	ESTs	573775 IMAGE Consortium	AA119030
8067 D	6 021GAMN0	EST	575427 IMAGE Consortium	AI447546
8068 D	8 021GAMN0	estrogen receptor 1 (alpha)	576624 IMAGE Consortium	AI386103
8069 D	10 021GAMN0	dihydropyrimidinase	987631 IMAGE Consortium	AA575393
8070 D	12 021GAMN0	ESTs, Highly similar to RSP5 PROTEIN [Saccharomyces cerevisiae]	1001390 IMAGE Consortium	AA771635
8071 F	2 021GAMN0	ESTs	483420 IMAGE Consortium	AI552477
8072 F	4 021GAMN0	ESTs	575207 IMAGE Consortium	AI530648
8073 F	6 021GAMN0	synaptotagmin 4	574704 IMAGE Consortium	AI323217
8074 F	8 021GAMN0	ESTs	575921 IMAGE Consortium	AA117609
8075 F	10 021GAMN0	insulin-like growth factor 1	1001007 IMAGE Consortium	AI604642
8076 F	12 021GAMN0	muscleblind-like (Drosophila)	1001424 IMAGE Consortium	AA666535
8077 H	2 021GAMN0	protoporphyrinogen oxidase	482868 IMAGE Consortium	AI323109
8078 H	4 021GAMN0	ESTs	573823 IMAGE Consortium	AA119061
8079 H	6 021GAMN0	ESTs	575453 IMAGE Consortium	AI530679
8080 H	8 021GAMN0	protein tyrosine phosphatase, receptor type, A	987763 IMAGE Consortium	AA572365
8081 H	10 021GAMN0			
8082 H	12 021GAMN0	ESTs, Highly similar to RO52_MOUSE 52 KD RO PROTEIN [M.musculus]	1001437 IMAGE Consortium	AA666549
8083 B	2 021NAMN1	ESTs	1001633 IMAGE Consortium	AI587969
8084 B	4 021NAMN1	DNA segment, Chr 6, ERATO Doi 352, expressed	1002544 IMAGE Consortium	AA596742
8085 B	6 021NAMN1	ESTs, Moderately similar to MO25 PROTEIN [Mus musculus]	1002811 IMAGE Consortium	AA600488
8086 B	8 021NAMN1	ESTs	1020655 IMAGE Consortium	AI594161
8087 B	10 021NAMN1	ESTs	1002065 IMAGE Consortium	AA684314
8088 B	12 021NAMN1	prostaglandin E receptor EP4 subtype	1002310 IMAGE Consortium	AA596456
8089 D	2 021NAMN1	ESTs, Highly similar to hypothetical SBBI03 protein [H.sapiens]	1001668 IMAGE Consortium	AA666641
8090 D	4 021NAMN1	ESTs	1002567 IMAGE Consortium	AI606342
8091 D	6 021NAMN1	serine protease inhibitor 4	1002830 IMAGE Consortium	AA600496
8092 D	8 021NAMN1	ESTs, Moderately similar to trg [R.norvegicus]	1020761 IMAGE Consortium	AI503211
8093 D	10 021NAMN1	ESTs, Moderately similar to SIGNAL RECOGNITION PARTICLE 19 KD PROTEIN [Homo sapiens]	1002088 IMAGE Consortium	AA684367
8094 D	12 021NAMN1	ESTs, Highly similar to spliceosomal protein SAP 155 [H.sapiens]	1002357 IMAGE Consortium	AA596555
8095 F	2 021NAMN1	ESTs	1001675 IMAGE Consortium	AI587976
8096 F	4 021NAMN1	latexin	1002571 IMAGE Consortium	AA596753
8097 F	6 021NAMN1	ESTs, Moderately similar to unnamed protein product [H.sapiens]	1002862 IMAGE Consortium	AA600515
8098 F	8 021NAMN1	heat shock 70 kDa protein 4	1020794 IMAGE Consortium	AA645750
8099 F	10 021NAMN1			
8100 F	12 021NAMN1	ESTs, Highly similar to AF155353_1 ankyrin repeat-containing protein Asb-2 [M.musculus]	1002370 IMAGE Consortium	AA596484



8101 H	2 021NAMN1	glycine transporter 1	1000982 IMAGE Consortium	AA771321
8102 H	4 021NAMN1	ESTs	1002593 IMAGE Consortium	AI606345
8103 H	6 021NAMN1	ESTs, Weakly similar to AF182443_1 F-box protein FBL2 [R.norvegicus]	1021095 IMAGE Consortium	AA645952
8104 H	8 021NAMN1	histocompatibility 2, D region locus 1	1020808 IMAGE Consortium	AI594174
8105 H	10 021NAMN1	ESTs	1002157 IMAGE Consortium	AA608101
8106 H	12 021NAMN1	numb gene homolog (Drosophila)	1002402 IMAGE Consortium	AA596565
8107 B	2 021UAMN2	Mus musculus AKR T-cell receptor gamma chain V gamma 1-J gamma 4-C gamma 4 region mRNA, partial cds	1002501 IMAGE Consortium	AA596594
8108 B	4 021UAMN2	ubiquitously expressed transcript	1021151 IMAGE Consortium	AA646227
8109 B	6 021UAMN2	ESTs, Highly similar to SPECTRIN ALPHA CHAIN, NON-ERYTHROID [Rattus norvegicus]	1038868 IMAGE Consortium	AA624568
8110 B	8 021UAMN2	secreted modular calcium-binding protein 2	1023630 IMAGE Consortium	AA799008
8111 B	10 021UAMN2	Ste-20 related kinase	1039234 IMAGE Consortium	AA624649
8112 B	12 021UAMN2	ESTs, Highly similar to unnamed protein product [H.sapiens]	1038979 IMAGE Consortium	AA624868
8113 D	2 021UAMN2	ESTs	1001803 IMAGE Consortium	AI587997
8114 D	4 021UAMN2	ESTs, Moderately similar to unnamed protein product [H.sapiens]	1038630 IMAGE Consortium	AA671177
8115 D	6 021UAMN2	Mus musculus mRNA for ADP-ribosylation factor 1 GTPase activating protein (ARF1 GAP gene)	1038898 IMAGE Consortium	AA624702
8116 D	8 021UAMN2	mitogen activated protein kinase kinase kinase 1	1023720 IMAGE Consortium	AA799087
8117 D	10 021UAMN2	islet mitochondrial antigen, 38 kD	1039348 IMAGE Consortium	AA624795
8118 D	12 021UAMN2	ESTs	1039836 IMAGE Consortium	AA615196
8119 F	2 021UAMN2			
8120 F	4 021UAMN2	ESTs, Highly similar to AF132947_1 CGI-13 protein [H.sapiens]	1038652 IMAGE Consortium	AA671264
8121 F	6 021UAMN2	procollagen, type V, alpha 2	1038921 IMAGE Consortium	AA624724
8122 F	8 021UAMN2	ESTs	1023777 IMAGE Consortium	AA799142
8123 F	10 021UAMN2	troponin T2, cardiac	1038690 IMAGE Consortium	AA671284
8124 F	12 021UAMN2	ESTs	1039995 IMAGE Consortium	AA611343
8125 H	2 021UAMN2	ESTs	1002696 IMAGE Consortium	AA607443
8126 H	4 021UAMN2	RNA polymerase 1-3 (16 kDa subunit)	1038678 IMAGE Consortium	AA671280
8127 H	6 021UAMN2	tubby-like protein 3	1038942 IMAGE Consortium	AA624839
8128 H	8 021UAMN2	vesicle transport through interaction with t-SNAREs 1b homolog	1023848 IMAGE Consortium	AA866934
8129 H	10 021UAMN2	hydroxyacyl-Coenzyme A dehydrogenase, type II	1038789 IMAGE Consortium	AA671508
8130 H	12 021UAMN2	Vav2 oncogene	1040019 IMAGE Consortium	AA612119
8131 B	2 0211AMN3	DNA segment, Chr 15, ERATO Doi 785, expressed	1040192 IMAGE Consortium	AA615287
8132 B	4 0211AMN3	keratin complex 1, acidic, gene 18	1077443 IMAGE Consortium	AA727960
8133 B	6 0211AMN3	alcohol dehydrogenase family 3, subfamily A2	1077617 IMAGE Consortium	AA821976
8134 B	8 0211AMN3	UDP-glucuronosyltransferase 1 family, member 1	1077767 IMAGE Consortium	AA822117

8135 B	10 0211AMN3	serine protease inhibitor 2-2	1077628 IMAGE Consortium	AA821980
8136 B	12 0211AMN3	Ral-interacting protein 1	1040585 IMAGE Consortium	AA612374
8137 D	2 0211AMN3	acyl-Coenzyme A oxidase	1040259 IMAGE Consortium	AA612012
8138 D	4 0211AMN3	Harvey rat sarcoma oncogene, subgroup R	1077495 IMAGE Consortium	AA821900
8139 D	6 0211AMN3	sterol carrier protein 2, liver	1077666 IMAGE Consortium	AA822085
8140 D	8 0211AMN3	ESTs	1040879 IMAGE Consortium	AA607765
8141 D	10 0211AMN3	hemopexin	1077673 IMAGE Consortium	AA822009
8142 D	12 0211AMN3	uridine monophosphate kinase	1040596 IMAGE Consortium	AA612418
8143 F	2 0211AMN3	ESTs	1040336 IMAGE Consortium	AA612154
8144 F	4 0211AMN3	endoglin	1077502 IMAGE Consortium	AI594824
8145 F	6 0211AMN3	homocysteine-inducible, endoplasmic reticulum stress-inducible, ubiquitin-like domain member 1	1077687 IMAGE Consortium	AA822015
8146 F	8 0211AMN3	mesenchyme homeobox 1	1040887 IMAGE Consortium	AA607781
8147 F	10 0211AMN3	apolipoprotein A-I	1077744 IMAGE Consortium	AA822098
8148 F	12 0211AMN3	ESTs, Moderately similar to AF225902_1 separation anxiety protein [D.melanogaster]	1040610 IMAGE Consortium	AA612423
8149 H	2 0211AMN3	ESTs, Highly similar to CALCIUM-DEPENDENT PROTEASE, SMALL [Sus scrofa]	1040479 IMAGE Consortium	AI594512
8150 H	4 0211AMN3	acyl-Coenzyme A dehydrogenase, very long chain	1077517 IMAGE Consortium	AA821917
8151 H	6 0211AMN3	serine protease inhibitor 1-1	1077714 IMAGE Consortium	AA822027
8152 H	8 0211AMN3	amylase 2, pancreatic	1077480 IMAGE Consortium	AA821884
8153 H	10 0211AMN3	endothelin 1	1040154 IMAGE Consortium	AA792278
8154 H	12 0211AMN3	extracellular matrix protein 1	1040661 IMAGE Consortium	AA606242
8155 B	2 0218AMN4	ESTs	1040757 IMAGE Consortium	AA606735
8156 B	4 0218AMN4	heparanase	1092868 IMAGE Consortium	AA674378
8157 B	6 0218AMN4	ESTs, Weakly similar to CG15403 gene product [D.melanogaster]	1092245 IMAGE Consortium	AA755396
8158 B	8 0218AMN4	ESTs, Moderately similar to hypothetical protein [H.sapiens]	1148738 IMAGE Consortium	AA791669
8159 B	10 0218AMN4	ESTs, Weakly similar to Similarity to Entamoeba elongation factor 1-alpha [C.elegans]	1148425 IMAGE Consortium	AA646290
8160 B	12 0218AMN4			
8161 D	2 0218AMN4	ESTs, Weakly similar to B39066 proline-rich protein 15 - rat [R.norvegicus]	1040794 IMAGE Consortium	AA607583
8162 D	4 0218AMN4	ESTs	1092883 IMAGE Consortium	AA674395
8163 D	6 0218AMN4	ESTs, Weakly similar to retinal short-chain dehydrogenase/reductase [M.musculus]	1092281 IMAGE Consortium	AA822480
8164 D	8 0218AMN4	flotillin 1	1148762 IMAGE Consortium	AA673706
8165 D	10 0218AMN4	small inducible cytokine A9	1148463 IMAGE Consortium	AA646314
8166 D	12 0218AMN4			
8167 F	2 0218AMN4	elastase 2	1093168 IMAGE Consortium	AA675084
8168 F	4 0218AMN4	lectin, galactose binding, soluble 9	1092968 IMAGE Consortium	AA674445
8169 F	6 0218AMN4	ESTs, Weakly similar to AF121862_1 sorting nexin 13	1139761 IMAGE Consortium	AA645707

8170 F	8 0218AMN4	[H.sapiens] ESTs, Weakly similar to AF183144_1 left-right dynein [M.musculus]	1148768 IMAGE Consortium	AA673709
8171 F	10 0218AMN4	ESTs	1148478 IMAGE Consortium	AI591812
8172 F	12 0218AMN4			
8173 H	2 0218AMN4	ESTs, Highly similar to COMPLEMENT C8 GAMMA CHAIN PRECURSOR [Homo sapiens]	1091732 IMAGE Consortium	AA670737
8174 H	4 0218AMN4	histocompatibility 2, class II antigen E beta	1092018 IMAGE Consortium	AA792785
8175 H	6 0218AMN4	DNA segment, Chr 4, ERATO Doi 765, expressed	1093058 IMAGE Consortium	AA674655
8176 H	8 0218AMN4	DNA segment, Chr 12, ERATO Doi 604, expressed	1148797 IMAGE Consortium	AA673741
8177 H	10 0218AMN4	ESTs, Weakly similar to cathepsin F [M.musculus]	1139752 IMAGE Consortium	AA645686
8178 H	12 0218AMN4			
8179 B	2 021FAMN5			
8180 B	4 021FAMN5			
8181 B	6 021FAMN5			
8182 B	8 021FAMN5	neuronal protein 15.6	1149003 IMAGE Consortium	AA655648
8183 B	10 021FAMN5	ESTs, Weakly similar to hypothetical protein [H.sapiens]	1149095 IMAGE Consortium	AA655909
8184 B	12 021FAMN5	ESTs	1149633 IMAGE Consortium	AI644974
8185 D	2 021FAMN5			
8186 D	4 021FAMN5			
8187 D	6 021FAMN5	ESTs	1148828 IMAGE Consortium	AA717156
8188 D	8 021FAMN5	ESTs	1149015 IMAGE Consortium	AA655653
8189 D	10 021FAMN5	ESTs, Highly similar to N-ACETYLGUCOSAMINE-6- SULFATASE PRECURSOR [Homo sapiens]	1149103 IMAGE Consortium	AA655762
8190 D	12 021FAMN5	clathrin, light polypeptide (Lca)	1149658 IMAGE Consortium	AA656055
8191 F	2 021FAMN5			
8192 F	4 021FAMN5	mast cell protease 2	1149578 IMAGE Consortium	AA655910
8193 F	6 021FAMN5	BCR downstream signaling 1	1148848 IMAGE Consortium	AA717170
8194 F	8 021FAMN5	ESTs, Highly similar to PROTEIN PHOSPHATASE PP2A, 55 KD REGULATORY SUBUNIT, ALPHA ISOFORM [Homo sapiens; Oryctolagus cuniculus]	1149029 IMAGE Consortium	AA655671
8195 F	10 021FAMN5	B-cell translocation gene 1, anti-proliferative	1149109 IMAGE Consortium	AA655765
8196 F	12 021FAMN5	ESTs	1149694 IMAGE Consortium	AA656126
8197 H	2 021FAMN5			
8198 H	4 021FAMN5	prion protein	1149672 IMAGE Consortium	AA656117
8199 H	6 021FAMN5	ESTs	1148870 IMAGE Consortium	AA717184
8200 H	8 021FAMN5	ESTs, Weakly similar to HYPOTHETICAL 86.9 KD PROTEIN ZK945.3 IN CHROMOSOME II [Caenorhabditis elegans]	1149037 IMAGE Consortium	AA655675
8201 H	10 021FAMN5	ESTs, Weakly similar to HYPOTHETICAL 91.2 KD PROTEIN IN RPS7A-SCH9 INTERGENIC REGION [Saccharomyces cerevisiae]	1149138 IMAGE Consortium	AA655779
8202 H	12 021FAMN5	sarcoglycan, epsilon	1149778 IMAGE Consortium	AA674109

8203 B	2	021MAMN6	ESTs	1150307	IMAGE Consortium	AA930043
8204 B	4	021MAMN6	annexin A5	1150770	IMAGE Consortium	AA791819
8205 B	6	021MAMN6	ganglioside-induced differentiation-associated-protein 10	1193214	IMAGE Consortium	AA718465
8206 B	8	021MAMN6	ESTs, Highly similar to INTERFERON-ACTIVATABLE PROTEIN 204 [Mus musculus]	1193534	IMAGE Consortium	AA727023
8207 B	10	021MAMN6	lectin, galactose binding, soluble 1	1193680	IMAGE Consortium	AA726983
8208 B	12	021MAMN6	M.musculus ART3 gene	1197070	IMAGE Consortium	AA733607
8209 D	2	021MAMN6	ESTs	1150172	IMAGE Consortium	AA929831
8210 D	4	021MAMN6	Mus musculus brain cDNA, clone MNCb-4134	1150897	IMAGE Consortium	AA795714
8211 D	6	021MAMN6	S100 calcium binding protein A1	1193218	IMAGE Consortium	AA718467
8212 D	8	021MAMN6	Bcl2-interacting killer-like	1193544	IMAGE Consortium	AA726901
8213 D	10	021MAMN6	ESTs, Highly similar to T17338 hypothetical protein DKFZp434O125.1 - human [H.sapiens]	1193691	IMAGE Consortium	AA727007
8214 D	12	021MAMN6	complement component 3	1197111	IMAGE Consortium	AA733629
8215 F	2	021MAMN6	spermiogenesis gene	1150294	IMAGE Consortium	AA930023
8216 F	4	021MAMN6	ESTs	1150932	IMAGE Consortium	AA791426
8217 F	6	021MAMN6	ESTs	1193242	IMAGE Consortium	AA718487
8218 F	8	021MAMN6	ESTs, Highly similar to unnamed protein product [H.sapiens]	1193574	IMAGE Consortium	AA726919
8219 F	10	021MAMN6	protein tyrosine kinase 2 beta	1193699	IMAGE Consortium	AA727011
8220 F	12	021MAMN6	ESTs, Moderately similar to skin-specific protein [H.sapiens]	1196439	IMAGE Consortium	AA794174
8221 H	2	021MAMN6	ESTs	1151022	IMAGE Consortium	AA791685
8222 H	4	021MAMN6	ATPase, H+ transporting, lysosomal (vacuolar proton pump) noncatalytic accessory protein 1 (110/160 kDa)	1150963	IMAGE Consortium	AA791459
8223 H	6	021MAMN6	FXYD domain-containing ion transport regulator 5	1193443	IMAGE Consortium	AA718314
8224 H	8	021MAMN6	ESTs	1193602	IMAGE Consortium	AA726942
8225 H	10	021MAMN6	ESTs	1193705	IMAGE Consortium	AA727013
8226 H	12	021MAMN6	ESTs	1196453	IMAGE Consortium	AA794190
8227 B	2	021TAMN7	Mus musculus tubby super-family protein (Tusp) mRNA, complete cds	1198059	IMAGE Consortium	AA727130
8228 B	4	021TAMN7	ESTs, Moderately similar to unnamed protein product [H.sapiens]	1196895	IMAGE Consortium	AA733685
8229 B	6	021TAMN7				
8230 B	8	021TAMN7	ESTs, Moderately similar to bicaudal-D [M.musculus]	1210084	IMAGE Consortium	AA727595
8231 B	10	021TAMN7	ESTs	1223074	IMAGE Consortium	AA733936
8232 B	12	021TAMN7	transient receptor protein 2	1224264	IMAGE Consortium	AA709475
8233 D	2	021TAMN7	Public domain EST	1197360	IMAGE Consortium	
8234 D	4	021TAMN7	ESTs	1196911	IMAGE Consortium	AI429050
8235 D	6	021TAMN7	RuvB-like protein 2	1209448	IMAGE Consortium	AA726368
8236 D	8	021TAMN7	ESTs	1210206	IMAGE Consortium	AA727248
8237 D	10	021TAMN7	DNA fragmentation factor, alpha subunit	1223103	IMAGE Consortium	AA733970
8238 D	12	021TAMN7	organic cation transporter	1224272	IMAGE Consortium	AA709478

8239 F	2 021TAMN7	3'-phosphoadenosine 5'-phosphosulfate synthase 1	1197363 IMAGE Consortium	AA681705
8240 F	4 021TAMN7			
8241 F	6 021TAMN7	target of myb1 homolog (chicken)	1209881 IMAGE Consortium	AA727299
8242 F	8 021TAMN7	crystallin, beta A4	1210228 IMAGE Consortium	AA727265
8243 F	10 021TAMN7	opioid receptor, sigma 1	1223128 IMAGE Consortium	AA734030
8244 F	12 021TAMN7	Cbp/p300-interacting transactivator with Glu/Asp-rich carboxy-terminal domain 1	1224295 IMAGE Consortium	AA709508
8245 H	2 021TAMN7	androgen regulated protein 1	1197400 IMAGE Consortium	AA794206
8246 H	4 021TAMN7	ESTs, Weakly similar to NED4_MOUSE NEDD-4 PROTEIN [M.musculus]	1209128 IMAGE Consortium	AA726060
8247 H	6 021TAMN7	ESTs, Highly similar to ACPH_RAT ACYLAMINO-ACID-RELEASING ENZYME [R.norvegicus]	1209929 IMAGE Consortium	AA727365
8248 H	8 021TAMN7	ESTs, Weakly similar to granuphilin-a [M.musculus]	1210251 IMAGE Consortium	AA727330
8249 H	10 021TAMN7	ESTs, Highly similar to AF024631_1 ANG2 [H.sapiens]	1223155 IMAGE Consortium	AA734063
8250 H	12 021TAMN7	ESTs	1224339 IMAGE Consortium	AA709774
8251 B	2 0210AMN8	ESTs	1223942 IMAGE Consortium	AA667677
8252 B	4 0210AMN8	carboxyl terminal LIM domain protein 1	1222560 IMAGE Consortium	AA815681
8253 B	6 0210AMN8	ESTs	1210557 IMAGE Consortium	AA733740
8254 B	8 0210AMN8	complement component 1, q subcomponent, alpha polypeptide	597157 IMAGE Consortium	AA145122
8255 B	10 0210AMN8	zinc finger protein 62	596642 IMAGE Consortium	AA146501
8256 B	12 0210AMN8	myosin light chain, phosphorylatable, cardiac ventricles	1224259 IMAGE Consortium	AA795319
8257 D	2 0210AMN8	cell adhesion kinase	1210652 IMAGE Consortium	AA733791
8258 D	4 0210AMN8	troponin I, cardiac	1222665 IMAGE Consortium	AA667301
8259 D	6 0210AMN8	ESTs, Moderately similar to ubiquitin protein ligase [M.musculus]	1211293 IMAGE Consortium	AA656303
8260 D	8 0210AMN8	adenylate cyclase 7	596455 IMAGE Consortium	AI324336
8261 D	10 0210AMN8	ESTs	597372 IMAGE Consortium	AA145919
8262 D	12 0210AMN8	ESTs, Highly similar to I5P1_HUMAN TYPE I INOSITOL-1,4,5-TRISPHOSPHATE 5-PHOSPHATASE [H.sapiens]	1224454 IMAGE Consortium	AA762709
8263 F	2 0210AMN8	a disintegrin and metalloprotease domain 11	1210678 IMAGE Consortium	AA733811
8264 F	4 0210AMN8	ESTs, Highly similar to ALDEHYDE DEHYDROGENASE, MITOCHONDRIAL X PRECURSOR [Homo sapiens]	1222766 IMAGE Consortium	AA667542
8265 F	6 0210AMN8	ankyrin repeat hooked to zinc finger motif	1210640 IMAGE Consortium	AA733787
8266 F	8 0210AMN8	TYRO protein tyrosine kinase binding protein	597257 IMAGE Consortium	AA138406
8267 F	10 0210AMN8	regulatory factor X-associated ankyrin-containing protein	596672 IMAGE Consortium	AA146531
8268 F	12 0210AMN8	chondroitin sulfate proteoglycan 2	1224492 IMAGE Consortium	AA755868
8269 H	2 0210AMN8	ESTs	1211588 IMAGE Consortium	AA815813
8270 H	4 0210AMN8	histone deacetylase 6	1222882 IMAGE Consortium	AA667900
8271 H	6 0210AMN8	testis specific protein kinase 1	596247 IMAGE Consortium	AA138787
8272 H	8 0210AMN8	ESTs	597261 IMAGE Consortium	AA138288
8273 H	10 0210AMN8	small inducible cytokine subfamily B (Cys-X-Cys), member 13	596683 IMAGE Consortium	AA138390

8274 H	12 0210AMN8	ESTs, Highly similar to KIAA0544 protein [H.sapiens]	1224502 IMAGE Consortium	AA755873
8275 B	2 0217AMN9	matrix metalloproteinase 2	1224700 IMAGE Consortium	AA756126
8276 B	4 0217AMN9	low density lipoprotein receptor related protein, associated protein 1	1227387 IMAGE Consortium	AA739111
8277 B	6 0217AMN9	serine protease inhibitor, Kunitz type 1	1227398 IMAGE Consortium	AA739106
8278 B	8 0217AMN9	ESTs	1227625 IMAGE Consortium	AA760067
8279 B	10 0217AMN9	DNA segment, Chr 15, ERATO Doi 785, expressed	1228029 IMAGE Consortium	AA760117
8280 B	12 0217AMN9	hematological and neurological expressed sequence 1	1228290 IMAGE Consortium	AA755023
8281 D	2 0217AMN9	EST	1224713 IMAGE Consortium	AI642716
8282 D	4 0217AMN9	ESTs, Moderately similar to hypothetical protein [H.sapiens]	1227432 IMAGE Consortium	AA738741
8283 D	6 0217AMN9	ESTs	1227431 IMAGE Consortium	AA739142
8284 D	8 0217AMN9	ESTs, Moderately similar to AF182076_1 glioma tumor suppressor candidate region protein 2 [H.sapiens]	1227653 IMAGE Consortium	AA760088
8285 D	10 0217AMN9	ESTs, Highly similar to AF073344_1 ubiquitin-specific protease 3 [H.sapiens]	1228110 IMAGE Consortium	AA760172
8286 D	12 0217AMN9	intergral membrane protein 1	1227964 IMAGE Consortium	AA760017
8287 F	2 0217AMN9	3-phosphoglycerate dehydrogenase	1224743 IMAGE Consortium	AI428595
8288 F	4 0217AMN9	solute carrier family 1, member 7	1227481 IMAGE Consortium	AA738833
8289 F	6 0217AMN9	gap junction membrane channel protein alpha 7	1227445 IMAGE Consortium	AA738755
8290 F	8 0217AMN9	ESTs, Weakly similar to cyclic nucleotide-gated channel beta subunit [R.norvegicus]	1227722 IMAGE Consortium	AA760131
8291 F	10 0217AMN9	carbonic anhydrase 6	1228119 IMAGE Consortium	AA760185
8292 F	12 0217AMN9	ESTs, Highly similar to apoptosis specific protein [H.sapiens]	1227978 IMAGE Consortium	AA760023
8293 H	2 0217AMN9	small nuclear ribonucleoprotein polypeptide A	1224758 IMAGE Consortium	AA756168
8294 H	4 0217AMN9	proteasome (prosome, macropain) subunit, alpha type 3	1227154 IMAGE Consortium	AA738642
8295 H	6 0217AMN9	malonyl-CoA decarboxylase	1227468 IMAGE Consortium	AA738765
8296 H	8 0217AMN9	ESTs, Highly similar to Unknown [H.sapiens]	1227749 IMAGE Consortium	AA760161
8297 H	10 0217AMN9	solute carrier family 31, member 2	1228159 IMAGE Consortium	AA760255
8298 H	12 0217AMN9	keratin complex 1, gene 2	1227995 IMAGE Consortium	AA760050
8299 B	2 021RAMNA	ESTs	1229850 IMAGE Consortium	AA791196
8300 B	4 021RAMNA	selectin, lymphocyte	1243158 IMAGE Consortium	AA798766
8301 B	6 021RAMNA	ESTs, Weakly similar to homolog of the Aspergillus nidulans sudD gene product [H.sapiens]	1261555 IMAGE Consortium	AA856137
8302 B	8 021RAMNA	ESTs	1262035 IMAGE Consortium	AA874087
8303 B	10 021RAMNA	actinin alpha 2 associated LIM protein	1260116 IMAGE Consortium	AA840053
8304 B	12 021RAMNA	methyl CpG binding protein 2	1260447 IMAGE Consortium	AI464023
8305 D	2 021RAMNA	inhibin beta-B	1229916 IMAGE Consortium	AA791253
8306 D	4 021RAMNA	ESTs, Moderately similar to CLATHRIN LIGHT CHAIN B [Bos taurus]	1243201 IMAGE Consortium	AA822256
8307 D	6 021RAMNA	Fyn proto-oncogene	1261710 IMAGE Consortium	AA874687
8308 D	8 021RAMNA	tumor-associated calcium signal transducer 2	1262045 IMAGE Consortium	AA874101

8309 D	10 021RAMNA	ESTs, Moderately similar to plakophilin 2a [H.sapiens]	1260165 IMAGE Consortium	AI463984
8310 D	12 021RAMNA	MyoD family inhibitor	1261202 IMAGE Consortium	AA838992
8311 F	2 021RAMNA	ESTs	1229991 IMAGE Consortium	AA789387
8312 F	4 021RAMNA	odorant-binding protein Ia	1243239 IMAGE Consortium	AA822346
8313 F	6 021RAMNA	interferon regulatory factor 6	1261835 IMAGE Consortium	AA873977
8314 F	8 021RAMNA	hypothetical protein MNCb-2778	1262054 IMAGE Consortium	AA874095
8315 F	10 021RAMNA	pleckstrin homology, Sec7 and coiled/coil domains 2	1260264 IMAGE Consortium	AI463922
8316 F	12 021RAMNA	adenomatosis polyposis coli	1260508 IMAGE Consortium	AA855420
8317 H	2 021RAMNA	cut (Drosophila)-like 1	1230097 IMAGE Consortium	AA798919
8318 H	4 021RAMNA	inhibitor of DNA binding 4	1243263 IMAGE Consortium	AA822366
8319 H	6 021RAMNA	ESTs	1261980 IMAGE Consortium	AA874133
8320 H	8 021RAMNA	biglycan	1261370 IMAGE Consortium	AA855996
8321 H	10 021RAMNA	ESTs	1260277 IMAGE Consortium	AI562143
8322 H	12 021RAMNA	butyrate response factor 1	1260690 IMAGE Consortium	AA855522
8323 B	2 021YAMNB	WW domain binding protein 2	1262354 IMAGE Consortium	AA880520
8324 B	4 021YAMNB	DNA segment, Chr 8, ERATO Doi 354, expressed	1276994 IMAGE Consortium	AI509860
8325 B	6 021YAMNB	insulin-like growth factor binding protein 2	1277469 IMAGE Consortium	AA879643
8326 B	8 021YAMNB	eyes absent 1 homolog (Drosophila)	318888 IMAGE Consortium	W34432
8327 B	10 021YAMNB	ESTs	330367 IMAGE Consortium	AI390811
8328 B	12 021YAMNB	ESTs	317322 IMAGE Consortium	AI536300
8329 D	2 021YAMNB	ESTs, Moderately similar to Apg12 [H.sapiens]	1262377 IMAGE Consortium	AI561507
8330 D	4 021YAMNB	ESTs	1262120 IMAGE Consortium	AA840249
8331 D	6 021YAMNB	surfactant associated protein D	1276786 IMAGE Consortium	AA880738
8332 D	8 021YAMNB	TEA domain family member 1	330475 IMAGE Consortium	W15025
8333 D	10 021YAMNB	homeo box D10	331130 IMAGE Consortium	W14783
8334 D	12 021YAMNB	ESTs, Highly similar to PROGASTRICSIN PRECURSOR [Rattus norvegicus]	317475 IMAGE Consortium	AI536309
8335 F	2 021YAMNB	ESTs	1262536 IMAGE Consortium	AA839176
8336 F	4 021YAMNB	fractured callus expressed transcript 1	1262144 IMAGE Consortium	AA880421
8337 F	6 021YAMNB			
8338 F	8 021YAMNB	differential display and activated by p53	329504 IMAGE Consortium	AI536384
8339 F	10 021YAMNB	ESTs	330413 IMAGE Consortium	W14850
8340 F	12 021YAMNB	ESTs	318253 IMAGE Consortium	AI536345
8341 H	2 021YAMNB	Mus musculus brain cDNA, clone MNCb-3925	1262558 IMAGE Consortium	AA839194
8342 H	4 021YAMNB	hydroxyprostaglandin dehydrogenase 15 (NAD)	1262159 IMAGE Consortium	AA880440
8343 H	6 021YAMNB	jagged 1	1277574 IMAGE Consortium	AA880220
8344 H	8 021YAMNB	ESTs	329599 IMAGE Consortium	AI451752
8345 H	10 021YAMNB	ESTs	331171 IMAGE Consortium	AI390865
8346 H	12 021YAMNB	ESTs	318268 IMAGE Consortium	AI536346
8347 B	2 0215AMNC	ESTs	318680 IMAGE Consortium	W14583
8348 B	4 0215AMNC			
8349 B	6 0215AMNC			

8350 B	8 0215AMNC			
8351 B	10 0215AMNC	ubiquitin-like 1	1279789 IMAGE Consortium	AA896203
8352 B	12 0215AMNC	ESTs, Moderately similar to DNA-DIRECTED RNA POLYMERASES I, II, AND III 7.3 KD POLYPEPTIDE [Schizosaccharomyces pombe]	1280741 IMAGE Consortium	AA790619
8353 D	2 0215AMNC	ESTs, Weakly similar to HYPOTHETICAL 29.8 KD PROTEIN IN THIC-HEME INTERGENIC REGION [Escherichia coli]	318705 IMAGE Consortium	W34412
8354 D	4 0215AMNC			
8355 D	6 0215AMNC			
8356 D	8 0215AMNC			
8357 D	10 0215AMNC	pre B-cell leukemia transcription factor 3	1279915 IMAGE Consortium	AA896541
8358 D	12 0215AMNC	ring finger protein 1	1280868 IMAGE Consortium	AA798181
8359 F	2 0215AMNC	ESTs, Weakly similar to FORM_MOUSE FORMIN [M.musculus]	318765 IMAGE Consortium	AI536364
8360 F	4 0215AMNC			
8361 F	6 0215AMNC			
8362 F	8 0215AMNC			
8363 F	10 0215AMNC	ESTs, Weakly similar to acid sphingomyelinase-like phosphodiesterase [M.musculus]	1280101 IMAGE Consortium	AA895031
8364 F	12 0215AMNC	ESTs	1279508 IMAGE Consortium	AA981240
8365 H	2 0215AMNC			
8366 H	4 0215AMNC			
8367 H	6 0215AMNC			
8368 H	8 0215AMNC			
8369 H	10 0215AMNC	involucrin	1280261 IMAGE Consortium	AA798100
8370 H	12 0215AMNC	transcytosis associated protein p115	1279593 IMAGE Consortium	AA981335
8371 B	2 021CAMND	ATP-binding cassette, sub-family C (CFTR/MRP), member 6	1279028 IMAGE Consortium	AA896664
8372 B	4 021CAMND	protein tyrosine phosphatase, non-receptor type 9	1277850 IMAGE Consortium	AA882203
8373 B	6 021CAMND	Mus musculus TCR beta locus from bases 1 to 250611 (section 1 of 3) of the complete sequence	1350074 IMAGE Consortium	AA982254
8374 B	8 021CAMND	ESTs	598695 IMAGE Consortium	AA163993
8375 B	10 021CAMND	ESTs	598166 IMAGE Consortium	AI592084
8376 B	12 021CAMND	interleukin 4	597674 IMAGE Consortium	AI324371
8377 D	2 021CAMND	ESTs, Weakly similar to Tbc1 [M.musculus]	1278317 IMAGE Consortium	AA895142
8378 D	4 021CAMND	axin	1278596 IMAGE Consortium	AA895724
8379 D	6 021CAMND	ESTs, Moderately similar to SM4D_MOUSE SEMAPHORIN 4D PRECURSOR [M.musculus]	1367026 IMAGE Consortium	AI020170
8380 D	8 021CAMND	N-acylsphingosine amidohydrolase 2	598741 IMAGE Consortium	AI592172
8381 D	10 021CAMND	ESTs	597530 IMAGE Consortium	AI447967
8382 D	12 021CAMND	ESTs, Highly similar to POL POLYPROTEIN [Squirrel monkey retrovirus]	597699 IMAGE Consortium	AA153229



8383 F	2	021CAMND	ESTs, Weakly similar to Evi-5 [M.musculus]	1279086	IMAGE Consortium	AA980986
8384 F	4	021CAMND	ESTs, Moderately similar to AF161427_1 HSPC309 [H.sapiens]	1279327	IMAGE Consortium	AA981134
8385 F	6	021CAMND	ESTs, Moderately similar to unnamed protein product [H.sapiens]	1367110	IMAGE Consortium	AI020512
8386 F	8	021CAMND	ESTs, Weakly similar to alpha glucosidase II, alpha subunit [M.musculus]	598775	IMAGE Consortium	AI592178
8387 F	10	021CAMND	ESTs	598292	IMAGE Consortium	AA155533
8388 F	12	021CAMND	ESTs, Moderately similar to S12207 hypothetical protein [M.musculus]	598401	IMAGE Consortium	AI592125
8389 H	2	021CAMND	ESTs	1279138	IMAGE Consortium	AA981032
8390 H	4	021CAMND	fucosyltransferase 8	1279348	IMAGE Consortium	AA981143
8391 H	6	021CAMND	DNA segment, Chr 17, human D6S56E 5	1367225	IMAGE Consortium	AI020407
8392 H	8	021CAMND	ESTs	598802	IMAGE Consortium	AI592182
8393 H	10	021CAMND	ESTs, Highly similar to hypothetical protein [M.musculus]	597592	IMAGE Consortium	AA144461
8394 H	12	021CAMND	ESTs	598439	IMAGE Consortium	AA155097
8395 B	2	021JAMNE	ESTs	598561	IMAGE Consortium	AI592148
8396 B	4	021JAMNE	ESTs, Weakly similar to unknown [H.sapiens]	775164	IMAGE Consortium	AA386874
8397 B	6	021JAMNE	Mus musculus strain C3H/HeN mRNA preferentially expressed in LPS-normoresponsive macrophages	775278	IMAGE Consortium	AA387131
8398 B	8	021JAMNE	ESTs	775439	IMAGE Consortium	AA386927
8399 B	10	021JAMNE	ESTs	789813	IMAGE Consortium	AA388171
8400 B	12	021JAMNE	ESTs, Highly similar to AF182293_1 U6 snRNA-associated Sm-like protein LSm7 [H.sapiens]	805191	IMAGE Consortium	AI551634
8401 D	2	021JAMNE	ESTs, Moderately similar to unnamed protein product [H.sapiens]	597874	IMAGE Consortium	AI592059
8402 D	4	021JAMNE	Msx-interacting-zinc finger	775174	IMAGE Consortium	AA386903
8403 D	6	021JAMNE	ESTs	775297	IMAGE Consortium	AA387017
8404 D	8	021JAMNE	ESTs	775452	IMAGE Consortium	AA387387
8405 D	10	021JAMNE	ESTs, Weakly similar to nuclear protein np95 [M.musculus]	790005	IMAGE Consortium	AA387951
8406 D	12	021JAMNE	ESTs, Weakly similar to AF154120_1 sorting nexin 1 [M.musculus]	805206	IMAGE Consortium	AI551635
8407 F	2	021JAMNE	ESTs	597881	IMAGE Consortium	AI592063
8408 F	4	021JAMNE	p53 apoptosis-associated target	775186	IMAGE Consortium	AA387163
8409 F	6	021JAMNE	neuroblastoma, suppression of tumorigenicity 1	775309	IMAGE Consortium	AA387115
8410 F	8	021JAMNE	ESTs, Highly similar to KIAA0783 protein [H.sapiens]	775473	IMAGE Consortium	AA388212
8411 F	10	021JAMNE	RAD1 homolog (S. pombe)	790054	IMAGE Consortium	AA387891
8412 F	12	021JAMNE	surfeit gene 4	805238	IMAGE Consortium	AA473296
8413 H	2	021JAMNE	colony stimulating factor 1 (macrophage)	597903	IMAGE Consortium	AA162153
8414 H	4	021JAMNE	ESTs	775195	IMAGE Consortium	AA387217
8415 H	6	021JAMNE	protein phosphatase 3, catalytic subunit, alpha isoform	775374	IMAGE Consortium	AA387372
8416 H	8	021JAMNE	ESTs, Highly similar to SEC63 [H.sapiens]	775477	IMAGE Consortium	AA388214

8417 H	10 021JAMNE	BANP homolog	790099 IMAGE Consortium	AA387967
8418 H	12 021JAMNE	ESTs	804515 IMAGE Consortium	AI551535
8419 B	2 021QAMNF	ESTs	805408 IMAGE Consortium	AA473209
8420 B	4 021QAMNF	ESTs	789992 IMAGE Consortium	AI608228
8421 B	6 021QAMNF	ESTs	819770 IMAGE Consortium	AA437542
8422 B	8 021QAMNF	wild-type p53-induced gene 1	819967 IMAGE Consortium	AA437794
8423 B	10 021QAMNF	tripeptidyl peptidase II	805528 IMAGE Consortium	AA397154
8424 B	12 021QAMNF	ESTs	834194 IMAGE Consortium	AI596512
8425 D	2 021QAMNF	ESTs	805453 IMAGE Consortium	AA474586
8426 D	4 021QAMNF	enhancer of polycomb homolog 1, ( <i>Drosophila</i> )	790017 IMAGE Consortium	AA387931
8427 D	6 021QAMNF	squamous cell carcinoma antigen recognized by T-cells 3	819797 IMAGE Consortium	AA437564
8428 D	8 021QAMNF	tissue factor pathway inhibitor	819970 IMAGE Consortium	AA437788
8429 D	10 021QAMNF	coding region determinant-binding protein	805624 IMAGE Consortium	AI604680
8430 D	12 021QAMNF	ribosomal protein S3	834218 IMAGE Consortium	AI596517
8431 F	2 021QAMNF	methyl-CpG binding domain protein 2	804732 IMAGE Consortium	AA474393
8432 F	4 021QAMNF	tat-interacting protein 30 kDa	790058 IMAGE Consortium	AA388037
8433 F	6 021QAMNF	hemoglobin alpha, adult chain 1	819822 IMAGE Consortium	AA437919
8434 F	8 021QAMNF	otoconin 90	819988 IMAGE Consortium	AA437800
8435 F	10 021QAMNF	ESTs	805637 IMAGE Consortium	AI604684
8436 F	12 021QAMNF	ESTs	834220 IMAGE Consortium	AI596518
8437 H	2 021QAMNF	ESTs	804743 IMAGE Consortium	AI551680
8438 H	4 021QAMNF	ESTs	790063 IMAGE Consortium	AA388045
8439 H	6 021QAMNF	ESTs, Highly similar to AF132972_1 CGI-38 protein [H.sapiens]	819844 IMAGE Consortium	AA437934
8440 H	8 021QAMNF	kinesin heavy chain member 1A	819992 IMAGE Consortium	AA437802
8441 H	10 021QAMNF	ESTs	805651 IMAGE Consortium	AI604687
8442 H	12 021QAMNF	ESTs	834230 IMAGE Consortium	AI596520
8443 B	2 021XAMNG	stem-loop binding protein	834517 IMAGE Consortium	AI596529
8444 B	4 021XAMNG	procollagen, type XV	819789 IMAGE Consortium	AA437677
8445 B	6 021XAMNG	matrilin 1, cartilage matrix protein 1	819929 IMAGE Consortium	AA403801
8446 B	8 021XAMNG	ESTs	894167 IMAGE Consortium	AA516997
8447 B	10 021XAMNG	ESTs	894374 IMAGE Consortium	AA517432
8448 B	12 021XAMNG	budding uninhibited by benzimidazoles 1 homolog, beta ( <i>S. cerevisiae</i> )	894487 IMAGE Consortium	AA517701
8449 D	2 021XAMNG	ESTs	833810 IMAGE Consortium	AA412912
8450 D	4 021XAMNG	ESTs, Moderately similar to unknown [H.sapiens]	819808 IMAGE Consortium	AA437679
8451 D	6 021XAMNG	ligand of numb-protein X	819934 IMAGE Consortium	AA403800
8452 D	8 021XAMNG	ESTs	894219 IMAGE Consortium	AA517221
8453 D	10 021XAMNG	ESTs	894390 IMAGE Consortium	AI482042
8454 D	12 021XAMNG	polymerase (DNA directed), delta 2, regulatory subunit (50 kDa)	894532 IMAGE Consortium	AA517595
8455 F	2 021XAMNG	ESTs	833843 IMAGE Consortium	AA413015
8456 F	4 021XAMNG	neuronatin	819828 IMAGE Consortium	AA437922

8457 F	6 021XAMNG	activin receptor interacting protein 1	819956 IMAGE Consortium	AA403807
8458 F	8 021XAMNG	ESTs, Moderately similar to KIAA0061 [H.sapiens]	894261 IMAGE Consortium	AI613719
8459 F	10 021XAMNG	zinc finger protein 292	894401 IMAGE Consortium	AI613737
8460 F	12 021XAMNG	unc119 homolog (C. elegans)	893855 IMAGE Consortium	AA517193
8461 H	2 021XAMNG	ribosomal protein, mitochondrial, S7	833897 IMAGE Consortium	AA413019
8462 H	4 021XAMNG	RAN guanine nucleotide release factor	819839 IMAGE Consortium	AA437693
8463 H	6 021XAMNG	translin	819971 IMAGE Consortium	AA437796
8464 H	8 021XAMNG	spermidine synthase	894284 IMAGE Consortium	AA517328
8465 H	10 021XAMNG	ubiquitin-conjugating enzyme 5	894408 IMAGE Consortium	AA517487
8466 H	12 021XAMNG	ESTs, Highly similar to OM20_RAT MITOCHONDRIAL IMPORT RECEPTOR SUBUNIT TOM20 HOMOLOG [R.norvegicus]	894585 IMAGE Consortium	AA517720
8467 B	2 0214AMNH	ESTs, Highly similar to zinc finger protein [H.sapiens]	893029 IMAGE Consortium	AI613884
8468 B	4 0214AMNH	ESTs	835439 IMAGE Consortium	AA445838
8469 B	6 0214AMNH	ESTs	933114 IMAGE Consortium	AA538152
8470 B	8 0214AMNH	ESTs, Highly similar to AFP2_HUMAN ARFAPTIN 2 [H.sapiens]	934278 IMAGE Consortium	AA543955
8471 B	10 0214AMNH	ESTs, Highly similar to hypothetical protein [H.sapiens]	933942 IMAGE Consortium	AA545625
8472 B	12 0214AMNH	mitogen activated protein kinase 1	934460 IMAGE Consortium	AA537011
8473 D	2 0214AMNH	ESTs	892395 IMAGE Consortium	AA501315
8474 D	4 0214AMNH	Public domain EST	892973 IMAGE Consortium	
8475 D	6 0214AMNH	ESTs	934000 IMAGE Consortium	AI550934
8476 D	8 0214AMNH	cellular repressor of E1A-stimulated genes	933700 IMAGE Consortium	AA544212
8477 D	10 0214AMNH	B-cell CLL/lymphoma 7B	933965 IMAGE Consortium	AA544488
8478 D	12 0214AMNH	histone gene complex 2	934509 IMAGE Consortium	AA536636
8479 F	2 0214AMNH	ESTs	892434 IMAGE Consortium	AA501319
8480 F	4 0214AMNH	ESTs, Highly similar to leukemia/lymphoma related factor LRF [M.musculus]	834857 IMAGE Consortium	AI596550
8481 F	6 0214AMNH	ESTs, Highly similar to ALANYL-TRNA SYNTHETASE [Homo sapiens]	934074 IMAGE Consortium	AA544561
8482 F	8 0214AMNH	phospholipase D3	933757 IMAGE Consortium	AA544314
8483 F	10 0214AMNH	ESTs, Weakly similar to A48830 probable transcription regulator NT fin12 - mouse [M.musculus]	934199 IMAGE Consortium	AA544780
8484 F	12 0214AMNH	ubiquitin-like 1 (sentrin) activating enzyme subunit 2	934555 IMAGE Consortium	AA536700
8485 H	2 0214AMNH	ESTs	835195 IMAGE Consortium	AA445739
8486 H	4 0214AMNH	ESTs	834976 IMAGE Consortium	AI596555
8487 H	6 0214AMNH	programmed cell death protein 7	934088 IMAGE Consortium	AA544574
8488 H	8 0214AMNH	ESTs, Weakly similar to ACYL-COA THIOESTERASE II [Escherichia coli]	934492 IMAGE Consortium	AA536621
8489 H	10 0214AMNH	ESTs, Highly similar to AF220184_1 uncharacterized hypothalamus protein HT010 [H.sapiens]	934222 IMAGE Consortium	AA544674
8490 H	12 0214AMNH	mitogen-activated protein kinase kinase kinase kinase 4	934583 IMAGE Consortium	AA536716
8491 B	2 021BAMNI	nucleosome binding protein 1	934745 IMAGE Consortium	AA617414

8492 B	4 021BAMNI	ESTs	934865 IMAGE Consortium	AA617441
8493 B	6 021BAMNI	ESTs	617313 IMAGE Consortium	AA174340
8494 B	8 021BAMNI	ESTs	617627 IMAGE Consortium	AA174893
8495 B	10 021BAMNI	ESTs, Weakly similar to predicted using Genefinder [C.elegans]	947393 IMAGE Consortium	AA589690
8496 B	12 021BAMNI	Mus musculus mevalonate kinase mRNA, complete cds	946070 IMAGE Consortium	AI550936
8497 D	2 021BAMNI	protease (prosome, macropain) 26S subunit, ATPase 1	934769 IMAGE Consortium	AA617416
8498 D	4 021BAMNI	ESTs, Highly similar to HYPOTHETICAL 52.8 KD PROTEIN T05E11.5 IN CHROMOSOME IV [Caenorhabditis elegans]	934892 IMAGE Consortium	AA617446
8499 D	6 021BAMNI	chloride channel calcium activated 1	616626 IMAGE Consortium	AI640041
8500 D	8 021BAMNI	ESTs	616980 IMAGE Consortium	AI640079
8501 D	10 021BAMNI	ESTs, Highly similar to exportin t [H.sapiens]	947425 IMAGE Consortium	AA591421
8502 D	12 021BAMNI	nuclear factor of kappa light chain gene enhancer in B-cells inhibitor, beta	946105 IMAGE Consortium	AI550944
8503 F	2 021BAMNI	ESTs, Highly similar to U2R1_MOUSE U2 SMALL NUCLEAR RIBONUCLEOPROTEIN AUXILIARY FACTOR 35 KD SUBUNIT RELATED-PROTEIN 1 [M.musculus]	934795 IMAGE Consortium	AA617424
8504 F	4 021BAMNI	ESTs, Highly similar to RRP5_HUMAN RRP5 PROTEIN HOMOLOG [H.sapiens]	934914 IMAGE Consortium	AA617451
8505 F	6 021BAMNI	ESTs	616682 IMAGE Consortium	AI640051
8506 F	8 021BAMNI	ESTs	599032 IMAGE Consortium	AA162218
8507 F	10 021BAMNI	ESTs, Weakly similar to potassium channel, alpha subunit [R.norvegicus]	947448 IMAGE Consortium	AI558179
8508 F	12 021BAMNI	ESTs, Weakly similar to OGT1_RAT UDP-N-ACETYLGLUCOSAMINE--PEPTIDE N-ACETYLGLUCOSAMINYLTRANSFERASE 110 KDA SUBUNIT [R.norvegicus]	946116 IMAGE Consortium	AA555970
8509 H	2 021BAMNI	zyxin	934800 IMAGE Consortium	AA617310
8510 H	4 021BAMNI	DNA Segment, Chr 6, human D12S2489E	616814 IMAGE Consortium	AA174479
8511 H	6 021BAMNI	ESTs	616702 IMAGE Consortium	AI640056
8512 H	8 021BAMNI	ESTs	599101 IMAGE Consortium	AA172835
8513 H	10 021BAMNI	ESTs, Weakly similar to KRAB-zinc finger protein KZF-1 [R.norvegicus]	962769 IMAGE Consortium	AA561924
8514 H	12 021BAMNI	profilin 1	946187 IMAGE Consortium	AA555743
8515 B	2 021IAMNJ	activin receptor IIB	935005 IMAGE Consortium	AA617503
8516 B	4 021IAMNJ	guanine nucleotide binding protein, related sequence 1	935239 IMAGE Consortium	AA617215
8517 B	6 021IAMNJ	gamma-glutamyl transpeptidase	945258 IMAGE Consortium	AA545451
8518 B	8 021IAMNJ	DNA segment, Chr 17, Abbott 1e	962777 IMAGE Consortium	AA561779
8519 B	10 021IAMNJ	coxsackievirus and adenovirus receptor	977269 IMAGE Consortium	AA560303
8520 B	12 021IAMNJ	Tal1 interrupting locus	976884 IMAGE Consortium	AA562546
8521 D	2 021IAMNJ	serine protease inhibitor, Kunitz type 2	935035 IMAGE Consortium	AA617562
8522 D	4 021IAMNJ			

8523 D	6 021IAMNJ	ESTs, Moderately similar to POLYADENYLATE-BINDING PROTEIN [ <i>Xenopus laevis</i> ]	945313 IMAGE Consortium	AA546580
8524 D	8 021IAMNJ	ESTs	962798 IMAGE Consortium	AI597198
8525 D	10 021IAMNJ	spastic paraplegia 4 homolog (human)	977312 IMAGE Consortium	AA560327
8526 D	12 021IAMNJ	ESTs, Weakly similar to CG2109 gene product [ <i>D.melanogaster</i> ]	976888 IMAGE Consortium	AI616382
8527 F	2 021IAMNJ	ESTs, Highly similar to RAS-RELATED PROTEIN RAB-8 [ <i>Homo sapiens</i> ; <i>Canis familiaris</i> ]	935106 IMAGE Consortium	AA617577
8528 F	4 021IAMNJ	neuroblastoma ras oncogene	935274 IMAGE Consortium	AA617228
8529 F	6 021IAMNJ	ESTs, Weakly similar to F25H5.6 [ <i>C.elegans</i> ]	945448 IMAGE Consortium	AA546438
8530 F	8 021IAMNJ	translocase of outer mitochondrial membrane 40 homolog (yeast)	962828 IMAGE Consortium	AA561789
8531 F	10 021IAMNJ	thymidine kinase 2, mitochondrial	977348 IMAGE Consortium	AI645425
8532 F	12 021IAMNJ	suppressor of Ty 5 homolog ( <i>S. cerevisiae</i> )	976893 IMAGE Consortium	AA562559
8533 H	2 021IAMNJ	troponin I, skeletal, slow 1	935132 IMAGE Consortium	AA616983
8534 H	4 021IAMNJ	ESTs, Highly similar to KIAA1438 protein [ <i>H.sapiens</i> ]	935296 IMAGE Consortium	AA617241
8535 H	6 021IAMNJ	ESTs, Highly similar to PROBABLE PEPTIDYL-PROLYL CIS-TRANS ISOMERASE C21E11.05C [ <i>Schizosaccharomyces pombe</i> ]	945481 IMAGE Consortium	AI585414
8536 H	8 021IAMNJ	ESTs, Weakly similar to /prediction	962922 IMAGE Consortium	AA619647
8537 H	10 021IAMNJ	secreted phosphoprotein 1	976791 IMAGE Consortium	AA563324
8538 H	12 021IAMNJ	Mus musculus mRNA for epididymal secretory protein, complete cds	976916 IMAGE Consortium	AA562569
8539 B	2 021PAMNK	Ubc6p homolog	976978 IMAGE Consortium	AA562976
8540 B	4 021PAMNK	ESTs, Moderately similar to KIAA0869 protein [ <i>H.sapiens</i> ]	989257 IMAGE Consortium	AI607022
8541 B	6 021PAMNK	ESTs	988782 IMAGE Consortium	AA638368
8542 B	8 021PAMNK	ESTs	977781 IMAGE Consortium	AA562609
8543 B	10 021PAMNK	ESTs	991477 IMAGE Consortium	AI604587
8544 B	12 021PAMNK	tumor rejection antigen P1A	991714 IMAGE Consortium	AA574624
8545 D	2 021PAMNK			
8546 D	4 021PAMNK	ESTs, Highly similar to SODIUM-DEPENDENT CHOLINE TRANSPORTER [ <i>Rattus norvegicus</i> ]	989281 IMAGE Consortium	AI607024
8547 D	6 021PAMNK	eukaryotic translation initiation factor 4A2	988801 IMAGE Consortium	AA638385
8548 D	8 021PAMNK			
8549 D	10 021PAMNK	enigma homolog 1 ( <i>R. norvegicus</i> )	991488 IMAGE Consortium	AI604588
8550 D	12 021PAMNK	es64 protein	991774 IMAGE Consortium	AA574693
8551 F	2 021PAMNK	ESTs, Moderately similar to density-regulated protein [ <i>H.sapiens</i> ]	976993 IMAGE Consortium	AA562997
8552 F	4 021PAMNK	translocase of inner mitochondrial membrane 13 homolog a (yeast)	988556 IMAGE Consortium	AA638208
8553 F	6 021PAMNK	ESTs	988878 IMAGE Consortium	AA572408
8554 F	8 021PAMNK	ESTs	988334 IMAGE Consortium	AA575501
8555 F	10 021PAMNK	ESTs, Highly similar to THERMOSENSITIVE GLUCONOKINASE [ <i>Escherichia coli</i> ]	992254 IMAGE Consortium	AA589300

8556 F	12 021PAMNK	ESTs	991991 IMAGE Consortium	AA574904
8557 H	2 021PAMNK	DNA segment, Chr 14, ERATO Doi 114, expressed	977761 IMAGE Consortium	AA562600
8558 H	4 021PAMNK	ESTs, Highly similar to KIAA0370 [H.sapiens]	988627 IMAGE Consortium	AA638270
8559 H	6 021PAMNK	ESTs	988938 IMAGE Consortium	AI586361
8560 H	8 021PAMNK	G two S phase expressed protein 1	988369 IMAGE Consortium	AA571318
8561 H	10 021PAMNK	ESTs	991550 IMAGE Consortium	AA575784
8562 H	12 021PAMNK	ESTs, Weakly similar to R12E2.12 [C.elegans]	989159 IMAGE Consortium	AA571365
8563 B	2 021WAMNL	Mus musculus sphingosine-1-phosphate phosphohydrolase (Spp1) mRNA, complete cds	989837 IMAGE Consortium	AA793528
8564 B	4 021WAMNL	ESTs	990257 IMAGE Consortium	AA590120
8565 B	6 021WAMNL	ribosomal protein L44	1004124 IMAGE Consortium	AA600581
8566 B	8 021WAMNL	ESTs	1004343 IMAGE Consortium	AA600439
8567 B	10 021WAMNL	ESTs	1004630 IMAGE Consortium	AA608137
8568 B	12 021WAMNL	zinc finger protein 101	992771 IMAGE Consortium	AA607983
8569 D	2 021WAMNL	ESTs, Highly similar to PPAL_MOUSE LYOSOMAL ACID PHOSPHATASE PRECURSOR [M.musculus]	989855 IMAGE Consortium	AA793534
8570 D	4 021WAMNL	ESTs	990607 IMAGE Consortium	AA590136
8571 D	6 021WAMNL	synapsin I	1004145 IMAGE Consortium	AA600597
8572 D	8 021WAMNL	ESTs, Weakly similar to AE003406_211 symbol	1004380 IMAGE Consortium	AI606390
8573 D	10 021WAMNL	ornithine decarboxylase antizyme inhibitor	1004646 IMAGE Consortium	AA608153
8574 D	12 021WAMNL	chromatin assembly factor 1, subunit A (p150)	992839 IMAGE Consortium	AA608013
8575 F	2 021WAMNL	ESTs, Highly similar to KINESIN-II 85 KD SUBUNIT [Strongylocentrotus purpuratus]	989929 IMAGE Consortium	AA591101
8576 F	4 021WAMNL	microtubule-associated protein tau	990734 IMAGE Consortium	AA590443
8577 F	6 021WAMNL	ESTs	1004204 IMAGE Consortium	AA600617
8578 F	8 021WAMNL	casein kinase II, beta subunit	1004392 IMAGE Consortium	AA600456
8579 F	10 021WAMNL	ESTs	1004680 IMAGE Consortium	AA608285
8580 F	12 021WAMNL	ESTs	992892 IMAGE Consortium	AA608033
8581 H	2 021WAMNL			
8582 H	4 021WAMNL	ESTs	990806 IMAGE Consortium	AA590768
8583 H	6 021WAMNL	ESTs, Weakly similar to CG14549 gene product [D.melanogaster]	1004224 IMAGE Consortium	AA600628
8584 H	8 021WAMNL	hexosaminidase A	1004409 IMAGE Consortium	AA600470
8585 H	10 021WAMNL	proteasome (prosome, macropain) 26S subunit, non-ATPase, 10	1004699 IMAGE Consortium	AA608308
8586 H	12 021WAMNL	ESTs	1005017 IMAGE Consortium	AA638747
8587 B	2 0213AMNM	protein phosphatase 1, catalytic subunit, gamma isoform	1005075 IMAGE Consortium	AA608341
8588 B	4 0213AMNM	ESTs	1005141 IMAGE Consortium	AA606614
8589 B	6 0213AMNM	Public domain EST	1004840 IMAGE Consortium	
8590 B	8 0213AMNM	U2 small nuclear ribonucleoprotein B	1004936 IMAGE Consortium	AA607096
8591 B	10 0213AMNM	ESTs	1006018 IMAGE Consortium	AA607057
8592 B	12 0213AMNM	ATP synthase, H <sup>+</sup> transporting, mitochondrial F0 complex, subunit b, isoform 1	1005664 IMAGE Consortium	AA607017

8593 D	2 0213AMNM	ESTs	1005079 IMAGE Consortium	AA608342
8594 D	4 0213AMNM	ESTs	1005164 IMAGE Consortium	AA606769
8595 D	6 0213AMNM	ESTs	1004852 IMAGE Consortium	AI594242
8596 D	8 0213AMNM	ESTs	1004942 IMAGE Consortium	AA607099
8597 D	10 0213AMNM	ESTs	1006092 IMAGE Consortium	AI594301
8598 D	12 0213AMNM	basic transcription element binding protein 2	1005671 IMAGE Consortium	AA607027
8599 F	2 0213AMNM	ESTs, Highly similar to CAAX prenyl protease [H.sapiens]	1005092 IMAGE Consortium	AA608338
8600 F	4 0213AMNM	meiotic check point regulator	1004663 IMAGE Consortium	AA608165
8601 F	6 0213AMNM	ESTs	1004859 IMAGE Consortium	AI594243
8602 F	8 0213AMNM	keratin complex 2, basic, gene 8	1004944 IMAGE Consortium	AA607100
8603 F	10 0213AMNM	ESTs	1006137 IMAGE Consortium	AA607186
8604 F	12 0213AMNM	ESTs	1005693 IMAGE Consortium	AA607123
8605 H	2 0213AMNM	ESTs, Weakly similar to small GTP binding protein [M.musculus]	1005107 IMAGE Consortium	AA606591
8606 H	4 0213AMNM	PFTAIRE protein kinase 1	1004734 IMAGE Consortium	AA606561
8607 H	6 0213AMNM	ribosomal protein, mitochondrial, L7	1004868 IMAGE Consortium	AA606973
8608 H	8 0213AMNM	ESTs, Weakly similar to similar to kinensin-like protein [C.elegans]	1004951 IMAGE Consortium	AA607108
8609 H	10 0213AMNM	Public domain EST	1006182 IMAGE Consortium	
8610 H	12 0213AMNM	ESTs, Moderately similar to RNA binding protein [R.norvegicus]	1005708 IMAGE Consortium	AA607128
8611 B	2 021AAMNN	ADP-ribosylation-like factor 6 interacting protein	1005809 IMAGE Consortium	AA606940
8612 B	4 021AAMNN	ESTs	1005886 IMAGE Consortium	AA606953
8613 B	6 021AAMNN	HIV-1 Rev binding protein	1005987 IMAGE Consortium	AA607046
8614 B	8 021AAMNN	ribosomal protein L21	1005451 IMAGE Consortium	AA608351
8615 B	10 021AAMNN	ESTs, Moderately similar to HYPOTHETICAL 9.3 KD PROTEIN ZK652.1 IN CHROMOSOME III [Caenorhabditis elegans]	1005769 IMAGE Consortium	AA606656
8616 B	12 021AAMNN	cDNA sequence AB023418	620243 IMAGE Consortium	AI604201
8617 D	2 021AAMNN	DNA segment, Chr 19, Wayne State University 162, expressed	1005812 IMAGE Consortium	AA606938
8618 D	4 021AAMNN	ESTs, Highly similar to NADH-UBIQUINONE OXIDOREDUCTASE B18 SUBUNIT [Bos taurus]	1005901 IMAGE Consortium	AA606827
8619 D	6 021AAMNN	ESTs, Weakly similar to CG11591 gene product [D.melanogaster]	1005991 IMAGE Consortium	AA607048
8620 D	8 021AAMNN	ESTs	1005470 IMAGE Consortium	AA608356
8621 D	10 021AAMNN			
8622 D	12 021AAMNN	plexin 6	621001 IMAGE Consortium	AA178447
8623 F	2 021AAMNN	ESTs, Highly similar to C-1 [H.sapiens]	1005830 IMAGE Consortium	AA606677
8624 F	4 021AAMNN	ESTs, Weakly similar to PS0136 H-2 class I histocompatibility antigen T15 - mouse [M.musculus]	1005924 IMAGE Consortium	AI594294
8625 F	6 021AAMNN	metal response element binding transcription factor 2	1005997 IMAGE Consortium	AA607051
8626 F	8 021AAMNN	ADP-ribosyltransferase (NAD+; poly (ADP-ribose)	1005479 IMAGE Consortium	AA608364

		polymerase) 2		
8627 F	10 021AAMNN	ribosomal protein L37a	1005818 IMAGE Consortium	AA606671
8628 F	12 021AAMNN	ESTs	621026 IMAGE Consortium	AA178452
8629 H	2 021AAMNN	phosphatidylinositol glycan, class F	1005834 IMAGE Consortium	AA606944
8630 H	4 021AAMNN	COP9 (constitutive photomorphogenic), subunit 7a (Arabidopsis)	1005928 IMAGE Consortium	AA606961
8631 H	6 021AAMNN	ESTs, Weakly similar to AF189262_1 GABA-A receptor epsilon-like subunit [R.norvegicus]	1005382 IMAGE Consortium	AA607234
8632 H	8 021AAMNN	ring finger protein 10	1005579 IMAGE Consortium	AA606908
8633 H	10 021AAMNN	Cd63 antigen	1005939 IMAGE Consortium	AA606842
8634 H	12 021AAMNN	ESTs	620371 IMAGE Consortium	AI604216
8635 B	2 021HAMNO	ESTs	621233 IMAGE Consortium	AA177415
8636 B	4 021HAMNO	ESTs	619988 IMAGE Consortium	AI604161
8637 B	6 021HAMNO	ESTs	1022220 IMAGE Consortium	AA623356
8638 B	8 021HAMNO	transducin (beta)-like 2	1006484 IMAGE Consortium	AA624327
8639 B	10 021HAMNO	ESTs, Highly similar to warts/lats-like kinase [M.musculus]	1006896 IMAGE Consortium	AA636991
8640 B	12 021HAMNO	POU domain, class 5, transcription factor 1	1038087 IMAGE Consortium	AA624527
8641 D	2 021HAMNO	ESTs	621267 IMAGE Consortium	AI592303
8642 D	4 021HAMNO	ESTs, Weakly similar to CG8195 gene product [D.melanogaster]	1037547 IMAGE Consortium	AA623569
8643 D	6 021HAMNO	ESTs	1022273 IMAGE Consortium	AI647128
8644 D	8 021HAMNO	ESTs, Weakly similar to CG2059 gene product [D.melanogaster]	1006507 IMAGE Consortium	AA624334
8645 D	10 021HAMNO	EST AA986430	1022984 IMAGE Consortium	AA623446
8646 D	12 021HAMNO	heat shock protein, 105 kDa	1038098 IMAGE Consortium	AA624532
8647 F	2 021HAMNO	ESTs	621328 IMAGE Consortium	AI592307
8648 F	4 021HAMNO	ESTs, Highly similar to Similar to D.melanogaster parallel sister chromatids protein [H.sapiens]	1037623 IMAGE Consortium	AA638835
8649 F	6 021HAMNO	ESTs, Highly similar to PROTEIN TRANSLATION FACTOR SUI1 HOMOLOG [Anopheles gambiae]	1006295 IMAGE Consortium	AI594314
8650 F	8 021HAMNO	claudin 4	1021865 IMAGE Consortium	AI048040
8651 F	10 021HAMNO	ESTs	1006951 IMAGE Consortium	AI594132
8652 F	12 021HAMNO	ESTs	1038112 IMAGE Consortium	AA624536
8653 H	2 021HAMNO	ESTs	620736 IMAGE Consortium	AI604258
8654 H	4 021HAMNO	CDK2 (cyclin-dependent kinase 2)-asscoated protein 1	1037664 IMAGE Consortium	AA638778
8655 H	6 021HAMNO	ESTs, Highly similar to nuclear respiratory factor-1 [M.musculus]	1006311 IMAGE Consortium	AI594316
8656 H	8 021HAMNO	ESTs	1021881 IMAGE Consortium	AI048074
8657 H	10 021HAMNO	RAD21 homolog (S. pombe)	1038010 IMAGE Consortium	AI642307
8658 H	12 021HAMNO	cartilage associated protein	1038124 IMAGE Consortium	AA624677
8659 B	2 021OAMNP	vacuolar protein sorting 4b (yeast)	1037608 IMAGE Consortium	AA623599
8660 B	4 021OAMNP	RAB geranylgeranyl transferase, b subunit	1037771 IMAGE Consortium	AA638889
8661 B	6 021OAMNP	ESTs, Moderately similar to AF181992_1 dynein light	1037883 IMAGE Consortium	AA638953



		intermediate chain 1 [R.norvegicus]		
8662 B	8 021OAMNP	ESTs	1038271 IMAGE Consortium	AA671205
8663 B	10 021OAMNP			
8664 B	12 021OAMNP	ESTs	1038454 IMAGE Consortium	AA624506
8665 D	2 021OAMNP	metallothionein 1	1037652 IMAGE Consortium	AA638765
8666 D	4 021OAMNP	S-adenosylhomocysteine hydrolase	1037775 IMAGE Consortium	AA638891
8667 D	6 021OAMNP	ESTs, Highly similar to PRE-MRNA SPLICING FACTOR SRP75 [Homo sapiens]	1037913 IMAGE Consortium	AA636937
8668 D	8 021OAMNP	ESTs, Highly similar to AF151893_1 CGI-135 protein [H.sapiens]	1038288 IMAGE Consortium	AA671214
8669 D	10 021OAMNP	polycystic kidney disease 2	1038380 IMAGE Consortium	AA671410
8670 D	12 021OAMNP	ESTs, Moderately similar to TVRTNU protein-tyrosine kinase [R.norvegicus]	1038464 IMAGE Consortium	AA624583
8671 F	2 021OAMNP	aplysia ras-related homolog A2	1037660 IMAGE Consortium	AA638769
8672 F	4 021OAMNP			
8673 F	6 021OAMNP	ESTs	1037919 IMAGE Consortium	AA636939
8674 F	8 021OAMNP	ESTs	1038290 IMAGE Consortium	AA671215
8675 F	10 021OAMNP	fibroblast growth factor inducible 14	1038390 IMAGE Consortium	AA671424
8676 F	12 021OAMNP	damage specific DNA binding protein 1 (127 kDa)	1038481 IMAGE Consortium	AA624605
8677 H	2 021OAMNP	ESTs	1037682 IMAGE Consortium	AI506717
8678 H	4 021OAMNP	ESTs	1037797 IMAGE Consortium	AA638908
8679 H	6 021OAMNP	ESTs, Highly similar to RAS-RELATED PROTEIN RAP-1A [Homo sapiens; Rattus norvegicus; Bos taurus]	1037940 IMAGE Consortium	AA638960
8680 H	8 021OAMNP	programmed cell death 6 interacting protein	1038292 IMAGE Consortium	AA671216
8681 H	10 021OAMNP	ESTs, Moderately similar to KIAA0731 protein [H.sapiens]	1038408 IMAGE Consortium	AA624482
8682 H	12 021OAMNP	ESTs, Highly similar to unnamed protein product [H.sapiens]	1038483 IMAGE Consortium	AA624606
8683 B	2 021VAMNQ	ESTs, Moderately similar to SFR8_HUMAN SPLICING FACTOR, ARGININE/SERINE-RICH 8 [H.sapiens]	1038540 IMAGE Consortium	AA624740
8684 B	4 021VAMNQ	U5 small nuclear ribonucleoprotein 116 kDa	1109330 IMAGE Consortium	AA666985
8685 B	6 021VAMNQ	ring finger protein 13	1109863 IMAGE Consortium	AA823953
8686 B	8 021VAMNQ	ESTs, Weakly similar to Y117_HUMAN HYPOTHETICAL PROTEIN KIAA0117 [H.sapiens]	1111035 IMAGE Consortium	AA683900
8687 B	10 021VAMNQ	Sir2 alpha	1110786 IMAGE Consortium	AA683789
8688 B	12 021VAMNQ	lumican	2101997 IMAGE Consortium	AW210283
8689 D	2 021VAMNQ	ESTs	1038562 IMAGE Consortium	AA624873
8690 D	4 021VAMNQ	ESTs, Highly similar to MITOTIC KINESIN-LIKE PROTEIN-1 [Homo sapiens]	1109380 IMAGE Consortium	AI591702
8691 D	6 021VAMNQ	growth arrest specific 5	1109888 IMAGE Consortium	AI591729
8692 D	8 021VAMNQ	cd164 antigen	1110319 IMAGE Consortium	AA674745
8693 D	10 021VAMNQ	ribosomal protein S26	1110919 IMAGE Consortium	AA683835
8694 D	12 021VAMNQ	chloride channel 3	2102038 IMAGE Consortium	AW210300
8695 F	2 021VAMNQ	thymine DNA glycosylase	1038581 IMAGE Consortium	AA624894

8696 F	4 021VAMNQ	capping protein alpha 2	1109498 IMAGE Consortium	AA667194
8697 F	6 021VAMNQ	ESTs	1109952 IMAGE Consortium	AA795144
8698 F	8 021VAMNQ	ESTs	1111090 IMAGE Consortium	AA688746
8699 F	10 021VAMNQ	M.musculus ASF mRNA	2102209 IMAGE Consortium	AW209264
8700 F	12 021VAMNQ	M.musculus ASF mRNA	2102052 IMAGE Consortium	AW210318
8701 H	2 021VAMNQ	ESTs	1038592 IMAGE Consortium	AA624890
8702 H	4 021VAMNQ			
8703 H	6 021VAMNQ	ESTs	1110197 IMAGE Consortium	AA795561
8704 H	8 021VAMNQ	SH3 domain protein 3	1111102 IMAGE Consortium	AA688751
8705 H	10 021VAMNQ	ESTs	2102238 IMAGE Consortium	AW209273
8706 H	12 021VAMNQ	ESTs, Highly similar to hMed7 [H.sapiens]	2102054 IMAGE Consortium	AW210319
8707 B	2 0212AMNR	multiple endocrine neoplasia 1	2102154 IMAGE Consortium	AW210407
8708 B	4 0212AMNR	ESTs	2101717 IMAGE Consortium	AW210034
8709 B	6 0212AMNR	BH3 interacting domain death agonist	2101239 IMAGE Consortium	AW209623
8710 B	8 0212AMNR	IMAGE EST {IMAGE:760858}	760858 IMAGE Consortium	AA386685
8711 B	10 0212AMNR	ESTs {IMAGE:761127}	761127 IMAGE Consortium	AA387679
8712 B	12 0212AMNR	ESTs, Weakly similar to GOLIATH PROTEIN [Drosophila melanogaster] {IMAGE:804680}	804680 IMAGE Consortium	AA474374
8713 D	2 0212AMNR	ESTs	2102169 IMAGE Consortium	AW210432
8714 D	4 0212AMNR	ESTs	2101813 IMAGE Consortium	AW210121
8715 D	6 0212AMNR	myeloid/lymphoid or mixed-lineage leukemia	2101286 IMAGE Consortium	AW209656
8716 D	8 0212AMNR	ESTs {IMAGE:894169}	894169 IMAGE Consortium	AA517043
8717 D	10 0212AMNR	ESTs {IMAGE:945014}	945014 IMAGE Consortium	AA537234
8718 D	12 0212AMNR	ESTs {IMAGE:834783}	834783 IMAGE Consortium	AA467436
8719 F	2 0212AMNR			
8720 F	4 0212AMNR	poly(rC)-binding protein 4	2101848 IMAGE Consortium	AW210151
8721 F	6 0212AMNR	hemoglobin Z, beta-like embryonic chain	2101337 IMAGE Consortium	AW209707
8722 F	8 0212AMNR	ESTs {IMAGE:805496}	805496 IMAGE Consortium	AA474600
8723 F	10 0212AMNR	ESTs, Highly similar to vasopressin activated calcium mobilizing receptor-like protein [H.sapiens] {IMAGE:833817}	833817 IMAGE Consortium	AA412921
8724 F	12 0212AMNR	ESTs, Highly similar to TRANSCRIPTION FACTOR IIIA [Xenopus laevis] {IMAGE:760965}	760965 IMAGE Consortium	AA386829
8725 H	2 0212AMNR	chaperonin subunit 8 (theta)	2101513 IMAGE Consortium	AW209866
8726 H	4 0212AMNR			
8727 H	6 0212AMNR	FSHD region gene 1	2101422 IMAGE Consortium	AW209777
8728 H	8 0212AMNR	Cytochrome P450, 2e1, ethanol inducible {IMAGE:760871}	760871 IMAGE Consortium	AA386802
8729 H	10 0212AMNR	ESTs {IMAGE:833899}	833899 IMAGE Consortium	AA413020
8730 H	12 0212AMNR	ESTs {IMAGE:805386}	805386 IMAGE Consortium	AA473205
8731 B	2 0219AMNS	ESTs {IMAGE:761038}	761038 IMAGE Consortium	AA387579
8732 B	4 0219AMNS	Mus musculus phosphomannomutase Sec53p homolog mRNA, complete cds {IMAGE:775067}	775067 IMAGE Consortium	AA387369

8733 B	6 0219AMNS	ESTs, Moderately similar to AF151800_1 CGI-41 protein [H.sapiens]	619118 IMAGE Consortium	AI592533
8734 B	8 0219AMNS	ESTs	618563 IMAGE Consortium	AA175909
8735 B	10 0219AMNS	ESTs	619562 IMAGE Consortium	AI604079
8736 B	12 0219AMNS	Mus musculus tasp mRNA for testis-specific adriamycin sensitivity protein, complete cds	619810 IMAGE Consortium	AI604124
8737 D	2 0219AMNS	ESTs {IMAGE:933229}	933229 IMAGE Consortium	AA541946
8738 D	4 0219AMNS	IMAGE EST {IMAGE:774877}	774877 IMAGE Consortium	AA386758
8739 D	6 0219AMNS	ESTs, Moderately similar to LEU5_HUMAN LEUKEMIA ASSOCIATED PROTEIN 5 [H.sapiens]	619147 IMAGE Consortium	AI592541
8740 D	8 0219AMNS	ESTs	618595 IMAGE Consortium	AA170260
8741 D	10 0219AMNS	ESTs	619617 IMAGE Consortium	AA175822
8742 D	12 0219AMNS	Mus musculus lecithin retinol acyltransferase mRNA, complete cds	619832 IMAGE Consortium	AI604132
8743 F	2 0219AMNS	ESTs {IMAGE:524078}	524078 IMAGE Consortium	AA066128
8744 F	4 0219AMNS	ESTs, Highly similar to TENSIN [Gallus gallus] {IMAGE:774810}	774810 IMAGE Consortium	AA386710
8745 F	6 0219AMNS	ESTs	619195 IMAGE Consortium	AI592551
8746 F	8 0219AMNS	ESTs	619312 IMAGE Consortium	AA172358
8747 F	10 0219AMNS	ESTs	619634 IMAGE Consortium	AI604093
8748 F	12 0219AMNS	ESTs	619921 IMAGE Consortium	AI604152
8749 H	2 0219AMNS	ESTs, Highly similar to HYPOTHETICAL 44.2 KD PROTEIN IN SCO2-MRF1 INTERGENIC REGION [Saccharomyces cerevisiae] {IMAGE:524792}	524792 IMAGE Consortium	AA067828
8750 H	4 0219AMNS	IMAGE EST {IMAGE:775208}	775208 IMAGE Consortium	AA387199
8751 H	6 0219AMNS	Noxa protein	619200 IMAGE Consortium	AI592552
8752 H	8 0219AMNS	ESTs, Moderately similar to T12464 hypothetical protein DKFZp564J102.1 - human [H.sapiens]	618619 IMAGE Consortium	AA170284
8753 H	10 0219AMNS	ESTs	619641 IMAGE Consortium	AA175830
8754 H	12 0219AMNS	ESTs	619974 IMAGE Consortium	AA178028
8755 B	2 021GAMNT	ESTs, Highly similar to PRE-MRNA SPLICING FACTOR PRP6 [Saccharomyces cerevisiae] {IMAGE:535535}	535535 IMAGE Consortium	AA087611
8756 B	4 021GAMNT	IMAGE EST {IMAGE:532250}	532250 IMAGE Consortium	AA066739
8757 B	6 021GAMNT	ESTs, Moderately similar to DOLICHYL-PHOSPHATE BETA-GLUCOSYLTRANSFERASE [Saccharomyces cerevisiae] {IMAGE:657077}	657077 IMAGE Consortium	AA241340
8758 B	8 021GAMNT			
8759 B	10 021GAMNT	ESTs {IMAGE:316112}	316112 IMAGE Consortium	W12256
8760 B	12 021GAMNT	ESTs {IMAGE:330910}	330910 IMAGE Consortium	W13947
8761 D	2 021GAMNT	Heparan sulfate proteoglycan 1, cell surface-associated (fibroglycan) {IMAGE:535688}	535688 IMAGE Consortium	AA106952
8762 D	4 021GAMNT	NAD(P)H menadione oxidoreductase 1, dioxin inducible {IMAGE:532186}	532186 IMAGE Consortium	AA068375
8763 D	6 021GAMNT	IMAGE EST {IMAGE:654228}	654228 IMAGE Consortium	AA197982

8764 D	8 021GAMNT	ESTs, Weakly similar to HEM45 [H.sapiens] {IMAGE:793347}	793347 IMAGE Consortium	AA438241
8765 D	10 021GAMNT	Cytochrome P450, 2b13, phenobarbitol inducible, type c {IMAGE:316288}	316288 IMAGE Consortium	W12874
8766 D	12 021GAMNT	IMAGE EST {IMAGE:331424}	331424 IMAGE Consortium	W14960
8767 F	2 021GAMNT	ESTs, Moderately similar to biphenyl hydrolase-related protein [H.sapiens] {IMAGE:524682}	524682 IMAGE Consortium	AA068497
8768 F	4 021GAMNT	IMAGE EST {IMAGE:523965}	523965 IMAGE Consortium	AA067731
8769 F	6 021GAMNT	IMAGE EST {IMAGE:656089}	656089 IMAGE Consortium	AA239554
8770 F	8 021GAMNT	ESTs {IMAGE:779373}	779373 IMAGE Consortium	AA414831
8771 F	10 021GAMNT	ESTs {IMAGE:317778}	317778 IMAGE Consortium	W13043
8772 F	12 021GAMNT	ESTs {IMAGE:335187}	335187 IMAGE Consortium	W17598
8773 H	2 021GAMNT	ESTs, Highly similar to CHROMOSOME REGION MAINTENANCE PROTEIN 1 [Schizosaccharomyces pombe] {IMAGE:533299}	533299 IMAGE Consortium	AA105546
8774 H	4 021GAMNT	ESTs, Weakly similar to L-ASPARAGINASE [Lupinus arboreus] {IMAGE:550910}	550910 IMAGE Consortium	AA097901
8775 H	6 021GAMNT	Renin 1 structural {IMAGE:661753}	661753 IMAGE Consortium	AA207872
8776 H	8 021GAMNT	ESTs {IMAGE:779847}	779847 IMAGE Consortium	AA414733
8777 H	10 021GAMNT	IMAGE EST {IMAGE:314290}	314290 IMAGE Consortium	W12679
8778 H	12 021GAMNT	ESTs {IMAGE:337077}	337077 IMAGE Consortium	W20763
8779 B	2 021NAMNU	IMAGE EST {IMAGE:352574}	352574 IMAGE Consortium	W34404
8780 B	4 021NAMNU			
8781 B	6 021NAMNU	ESTs {IMAGE:422959}	422959 IMAGE Consortium	W97983
8782 B	8 021NAMNU	Mus musculus major histocompatibility locus class II region; Fas-binding protein Daxx (DAXX) gene, partial cds; Bing1 (BING1), tapasin (tapasin), RalGDS-like factor (RLF), KE2 (KE2), BING4 (BING4), beta1, 3-galactosyl transferase {IMAGE:425385}	425385 IMAGE Consortium	AA000107
8783 B	10 021NAMNU	Neurotrophic tyrosine kinase, receptor, type 3 {IMAGE:438183}	438183 IMAGE Consortium	AA008069
8784 B	12 021NAMNU	ESTs {IMAGE:457541}	457541 IMAGE Consortium	AA021940
8785 D	2 021NAMNU	IMAGE EST {IMAGE:373521}	373521 IMAGE Consortium	W57234
8786 D	4 021NAMNU	ESTs {IMAGE:419398}	419398 IMAGE Consortium	W88079
8787 D	6 021NAMNU	ESTs {IMAGE:426212}	426212 IMAGE Consortium	AA003144
8788 D	8 021NAMNU	ESTs {IMAGE:424617}	424617 IMAGE Consortium	AA000564
8789 D	10 021NAMNU	ESTs {IMAGE:438412}	438412 IMAGE Consortium	AA008296
8790 D	12 021NAMNU	IMAGE EST {IMAGE:458349}	458349 IMAGE Consortium	AA024360
8791 F	2 021NAMNU	IMAGE EST {IMAGE:387243}	387243 IMAGE Consortium	W65748
8792 F	4 021NAMNU	ESTs, Moderately similar to HYPOTHETICAL PROTEIN KIAA0076 [H.sapiens] {IMAGE:423035}	423035 IMAGE Consortium	W98022
8793 F	6 021NAMNU	Neurofilament, light polypeptide {IMAGE:425315}	425315 IMAGE Consortium	W99951
8794 F	8 021NAMNU	IMAGE EST {IMAGE:425578}	425578 IMAGE Consortium	AA000018

8795 F	10 021NAMNU	IMAGE EST {IMAGE:439811}	439811 IMAGE Consortium	AA008593
8796 F	12 021NAMNU	ESTs, Moderately similar to AMILORIDE-SENSITIVE AMINE OXIDASE [Rattus norvegicus] {IMAGE:456716}	456716 IMAGE Consortium	AA023491
8797 H	2 021NAMNU	ESTs {IMAGE:386218}	386218 IMAGE Consortium	W65070
8798 H	4 021NAMNU	ESTs, Highly similar to RAS-RELATED PROTEIN RAC1 [Drosophila melanogaster] {IMAGE:422182}	422182 IMAGE Consortium	W97276
8799 H	6 021NAMNU	IMAGE EST {IMAGE:424348}	424348 IMAGE Consortium	W98625
8800 H	8 021NAMNU	IMAGE EST {IMAGE:425677}	425677 IMAGE Consortium	AA000310
8801 H	10 021NAMNU	IMAGE EST {IMAGE:437594}	437594 IMAGE Consortium	AA003561
8802 H	12 021NAMNU	IMAGE EST {IMAGE:457832}	457832 IMAGE Consortium	AA021966
8803 B	2 021UAMNV	IMAGE EST {IMAGE:464134}	464134 IMAGE Consortium	AA028359
8804 B	4 021UAMNV	ESTs	621431 IMAGE Consortium	AI592319
8805 B	6 021UAMNV	ESTs	621690 IMAGE Consortium	AI591829
8806 B	8 021UAMNV	ESTs	634065 IMAGE Consortium	AI592654
8807 B	10 021UAMNV	Mus musculus kruppel-related zinc finger protein (Emzf1) mRNA, complete cds {IMAGE:477064}	477064 IMAGE Consortium	AA047965
8808 B	12 021UAMNV	Fibroblast growth factor 15 {IMAGE:479758}	479758 IMAGE Consortium	AA051675
8809 D	2 021UAMNV	IMAGE EST {IMAGE:467394}	467394 IMAGE Consortium	AA036434
8810 D	4 021UAMNV	ESTs, Weakly similar to similar to yeast heat shock protein STI1 [C.elegans]	621525 IMAGE Consortium	AI592336
8811 D	6 021UAMNV	chromobox homolog 5 (Drosophila HP1a)	621771 IMAGE Consortium	AI591837
8812 D	8 021UAMNV	ESTs	634130 IMAGE Consortium	AI666267
8813 D	10 021UAMNV	ESTs {IMAGE:476086}	476086 IMAGE Consortium	AA049861
8814 D	12 021UAMNV	ESTs {IMAGE:479271}	479271 IMAGE Consortium	AA048540
8815 F	2 021UAMNV	ESTs, Highly similar to ZINC FINGER PROTEIN ZFP-37 [Mus musculus] {IMAGE:466313}	466313 IMAGE Consortium	AA033410
8816 F	4 021UAMNV	ESTs	622276 IMAGE Consortium	AI591893
8817 F	6 021UAMNV	ESTs, Weakly similar to KIAA0512 protein [H.sapiens]	622545 IMAGE Consortium	AI591914
8818 F	8 021UAMNV	ESTs	634200 IMAGE Consortium	AI592676
8819 F	10 021UAMNV	ESTs {IMAGE:477484}	477484 IMAGE Consortium	AA048400
8820 F	12 021UAMNV	ESTs {IMAGE:483628}	483628 IMAGE Consortium	AA061286
8821 H	2 021UAMNV	IMAGE EST {IMAGE:472808}	472808 IMAGE Consortium	AA038063
8822 H	4 021UAMNV	ESTs	622283 IMAGE Consortium	AA183616
8823 H	6 021UAMNV	ESTs	621843 IMAGE Consortium	AI591847
8824 H	8 021UAMNV	ESTs	634388 IMAGE Consortium	AA185228
8825 H	10 021UAMNV	ESTs {IMAGE:478157}	478157 IMAGE Consortium	AA049610
8826 H	12 021UAMNV	CELL SURFACE GLYCOPROTEIN MAC-1 ALPHA SUBUNIT PRECURSOR {IMAGE:483591}	483591 IMAGE Consortium	AA061908
8827 B	2 0211AMNW	ESTs {IMAGE:481507}	481507 IMAGE Consortium	AA060314
8828 B	4 0211AMNW	ESTs, Weakly similar to GLUTATHIONE S-TRANSFERASE, MITOCHONDRIAL [Rattus norvegicus] {IMAGE:571334}	571334 IMAGE Consortium	AA108741
8829 B	6 0211AMNW	ESTs, Moderately similar to KIAA0253 [H.sapiens]	573609 IMAGE Consortium	AA118848

		{IMAGE:573609}		
8830 B	8 0211AMNW	ESTs {IMAGE:575285}	575285 IMAGE Consortium	AA125312
8831 B	10 0211AMNW	IMAGE EST {IMAGE:581177}	581177 IMAGE Consortium	AA138559
8832 B	12 0211AMNW	ESTs {IMAGE:598479}	598479 IMAGE Consortium	AA161823
8833 D	2 0211AMNW	ESTs {IMAGE:482652}	482652 IMAGE Consortium	AA061980
8834 D	4 0211AMNW	ESTs, Highly similar to SWI/SNF complex 170 KDa subunit [H.sapiens] {IMAGE:572000}	572000 IMAGE Consortium	AA108369
8835 D	6 0211AMNW	ESTs {IMAGE:576535}	576535 IMAGE Consortium	AA123823
8836 D	8 0211AMNW	IMAGE EST {IMAGE:576958}	576958 IMAGE Consortium	AA116447
8837 D	10 0211AMNW	ESTs, Weakly similar to Des-1 protein [D.melanogaster] {IMAGE:577698}	577698 IMAGE Consortium	AA140530
8838 D	12 0211AMNW	IMAGE EST {IMAGE:598387}	598387 IMAGE Consortium	AA155021
8839 F	2 0211AMNW	IMAGE EST {IMAGE:481234}	481234 IMAGE Consortium	AA060385
8840 F	4 0211AMNW	Mus musculus nucleolar protein (MSP58) mRNA, complete cds {IMAGE:493227}	493227 IMAGE Consortium	AA098041
8841 F	6 0211AMNW	ESTs {IMAGE:576344}	576344 IMAGE Consortium	AA120260
8842 F	8 0211AMNW	Lymphotactin {IMAGE:576815}	576815 IMAGE Consortium	AA120030
8843 F	10 0211AMNW	ESTs, Highly similar to CORONIN [Dictyostelium discoideum] {IMAGE:577627}	577627 IMAGE Consortium	AA140511
8844 F	12 0211AMNW	ESTs {IMAGE:596340}	596340 IMAGE Consortium	AA138854
8845 H	2 0211AMNW	IMAGE EST {IMAGE:481860}	481860 IMAGE Consortium	AA059783
8846 H	4 0211AMNW	ESTs {IMAGE:573365}	573365 IMAGE Consortium	AA118680
8847 H	6 0211AMNW	IMAGE EST {IMAGE:574725}	574725 IMAGE Consortium	AA120657
8848 H	8 0211AMNW	ESTs {IMAGE:577409}	577409 IMAGE Consortium	AA116618
8849 H	10 0211AMNW	IMAGE EST {IMAGE:597035}	597035 IMAGE Consortium	AA145508
8850 H	12 0211AMNW	ESTs, Weakly similar to C15H9.5 [C.elegans] {IMAGE:583925}	583925 IMAGE Consortium	AA154527
8851 B	2 0218AMNX	IMAGE EST {IMAGE:598128}	598128 IMAGE Consortium	AA265439
8852 B	4 0218AMNX	ESTs, Weakly similar to D2085.4 [C.elegans] {IMAGE:618327}	618327 IMAGE Consortium	AA175034
8853 B	6 0218AMNX	ESTs {IMAGE:620311}	620311 IMAGE Consortium	AA177968
8854 B	8 0218AMNX	ESTs {IMAGE:619745}	619745 IMAGE Consortium	AA172771
8855 B	10 0218AMNX	IMAGE EST {IMAGE:634588}	634588 IMAGE Consortium	AA185939
8856 B	12 0218AMNX	ESTs {IMAGE:621883}	621883 IMAGE Consortium	AA178388
8857 D	2 0218AMNX	ESTs {IMAGE:596879}	596879 IMAGE Consortium	AA138580
8858 D	4 0218AMNX	IMAGE EST {IMAGE:618306}	618306 IMAGE Consortium	AA170599
8859 D	6 0218AMNX	IMAGE EST {IMAGE:620101}	620101 IMAGE Consortium	AA176037
8860 D	8 0218AMNX	ESTs {IMAGE:621316}	621316 IMAGE Consortium	AA178549
8861 D	10 0218AMNX	IMAGE EST {IMAGE:621630}	621630 IMAGE Consortium	AA183684
8862 D	12 0218AMNX	ESTs {IMAGE:621960}	621960 IMAGE Consortium	AA183747
8863 F	2 0218AMNX	IMAGE EST {IMAGE:595951}	595951 IMAGE Consortium	AA140542
8864 F	4 0218AMNX	ESTs {IMAGE:617360}	617360 IMAGE Consortium	AA174512
8865 F	6 0218AMNX	ESTs {IMAGE:619430}	619430 IMAGE Consortium	AA175625

8866 F	8 0218AMNX	IMAGE EST {IMAGE:622028}	622028 IMAGE Consortium	AA177505
8867 F	10 0218AMNX	ESTs, Moderately similar to Toll-like receptor 1 [H.sapiens] {IMAGE:622702}	622702 IMAGE Consortium	AA177549
8868 F	12 0218AMNX	ESTs {IMAGE:634728}	634728 IMAGE Consortium	AA163094
8869 H	2 0218AMNX			
8870 H	4 0218AMNX	ESTs, Weakly similar to CMRF35 ANTIGEN PRECURSOR [H.sapiens] {IMAGE:618824}	618824 IMAGE Consortium	AA175290
8871 H	6 0218AMNX	ESTs {IMAGE:620547}	620547 IMAGE Consortium	AA178286
8872 H	8 0218AMNX	ESTs {IMAGE:622542}	622542 IMAGE Consortium	AA183587
8873 H	10 0218AMNX	Immunoglobulin heavy chain 8 (heavy chain of IgG3) {IMAGE:635556}	635556 IMAGE Consortium	AA166427
8874 H	12 0218AMNX	ESTs, Moderately similar to JAK3 [M.musculus] {IMAGE:635916}	635916 IMAGE Consortium	AA183273
8875 B	2 021FAMNY	ESTs {IMAGE:621991}	621991 IMAGE Consortium	AA183803
8876 B	4 021FAMNY	ESTs {IMAGE:640096}	640096 IMAGE Consortium	AA197454
8877 B	6 021FAMNY	ESTs {IMAGE:639994}	639994 IMAGE Consortium	AA197601
8878 B	8 021FAMNY	ESTs {IMAGE:642836}	642836 IMAGE Consortium	AA185271
8879 B	10 021FAMNY	IMAGE EST {IMAGE:653189}	653189 IMAGE Consortium	AA217174
8880 B	12 021FAMNY	disintegrin metalloprotease (decysin)	635258 IMAGE Consortium	AA189495
8881 D	2 021FAMNY	Interleukin 16 {IMAGE:634936}	634936 IMAGE Consortium	AA155344
8882 D	4 021FAMNY			
8883 D	6 021FAMNY	IMAGE EST {IMAGE:639902}	639902 IMAGE Consortium	AA198514
8884 D	8 021FAMNY	ESTs, Highly similar to PUTATIVE UBIQUITIN CARBOXYL-TERMINAL HYDROLASE C13A11.04C [Schizosaccharomyces pombe] {IMAGE:642148}	642148 IMAGE Consortium	AA212162
8885 D	10 021FAMNY	protein tyrosine phosphatase, receptor type, E	635150 IMAGE Consortium	AI322925
8886 D	12 021FAMNY	ESTs, Highly similar to RL21_MOUSE 60S RIBOSOMAL PROTEIN L21□ [M.musculus]	635282 IMAGE Consortium	AA189503
8887 F	2 021FAMNY	Zinc finger protein X-linked {IMAGE:634652}	634652 IMAGE Consortium	AA155377
8888 F	4 021FAMNY	Dlk1-like homolog (Drosophila) {IMAGE:636875}	636875 IMAGE Consortium	AA189431
8889 F	6 021FAMNY	Mus musculus interferon regulatory factor 5 (mirf5) mRNA, complete cds {IMAGE:637338}	637338 IMAGE Consortium	AA185025
8890 F	8 021FAMNY	ESTs {IMAGE:652666}	652666 IMAGE Consortium	AA216849
8891 F	10 021FAMNY	ESTs	635010 IMAGE Consortium	AA155493
8892 F	12 021FAMNY	ESTs	636035 IMAGE Consortium	AI645132
8893 H	2 021FAMNY	Mus musculus mRNA for N-glycan alpha 2,8-sialyltransferase {IMAGE:638406}	638406 IMAGE Consortium	AA217008
8894 H	4 021FAMNY	ESTs {IMAGE:638217}	638217 IMAGE Consortium	AA200330
8895 H	6 021FAMNY	HpalI tiny fragments locus 9c {IMAGE:641594}	641594 IMAGE Consortium	AA209823
8896 H	8 021FAMNY	IMAGE EST {IMAGE:657952}	657952 IMAGE Consortium	AA198145
8897 H	10 021FAMNY	ESTs	635028 IMAGE Consortium	AI592792
8898 H	12 021FAMNY	ESTs	635317 IMAGE Consortium	AI592823
8899 B	2 021MAMNZ	ESTs	634841 IMAGE Consortium	AI592763

8900 B	4 021MAMNZ	ESTs, Weakly similar to XY40 protein [R.norvegicus]	635728 IMAGE Consortium	AA163479
8901 B	6 021MAMNZ	Fc receptor, IgE, high affinity I, gamma polypeptide	316763 IMAGE Consortium	W11251
8902 B	8 021MAMNZ	gap junction membrane channel protein alpha 4	332069 IMAGE Consortium	AI390909
8903 B	10 021MAMNZ	cytochrome c oxidase, subunit VIIIb	334467 IMAGE Consortium	W36632
8904 B	12 021MAMNZ	synaptosomal-associated protein, 91 kDa	387263 IMAGE Consortium	AI893941
8905 D	2 021MAMNZ	ESTs	634892 IMAGE Consortium	AA154777
8906 D	4 021MAMNZ	ESTs	635760 IMAGE Consortium	AI509284
8907 D	6 021MAMNZ	yolk sac gene 2	317643 IMAGE Consortium	W11483
8908 D	8 021MAMNZ	intercellular adhesion molecule	332372 IMAGE Consortium	W08586
8909 D	10 021MAMNZ	leukotriene C4 synthase	348993 IMAGE Consortium	W29221
8910 D	12 021MAMNZ	TAP binding protein	391467 IMAGE Consortium	AI430210
8911 F	2 021MAMNZ	ESTs	635603 IMAGE Consortium	AI509257
8912 F	4 021MAMNZ	ESTs	635786 IMAGE Consortium	AI509288
8913 F	6 021MAMNZ	prostaglandin I2 (prostacyclin) synthase	317686 IMAGE Consortium	W13004
8914 F	8 021MAMNZ	secretin	332800 IMAGE Consortium	AI325342
8915 F	10 021MAMNZ	sine oculis-related homeobox 1 homolog (Drosophila)	350912 IMAGE Consortium	AI893912
8916 F	12 021MAMNZ	regulator of G protein signaling 7	400795 IMAGE Consortium	W76764
8917 H	2 021MAMNZ	ESTs, Moderately similar to multidrug resistance protein [M.musculus]	635617 IMAGE Consortium	AA165775
8918 H	4 021MAMNZ	ESTs	635961 IMAGE Consortium	AI645119
8919 H	6 021MAMNZ	tenascin C	317871 IMAGE Consortium	W12942
8920 H	8 021MAMNZ	plasmin inhibitor alpha 2	333193 IMAGE Consortium	W33918
8921 H	10 021MAMNZ	aggrecan, structural proteoglycan of cartilage	353693 IMAGE Consortium	W41795
8922 H	12 021MAMNZ	alpha internexin neuronal intermediate filament protein	401796 IMAGE Consortium	W78548
8923 B	2 021MAMO0	Bcl2 homologous antagonist/killer	400326 IMAGE Consortium	W77319
8924 B	4 021MAMO0	sparc/osteonectin, cwcv and kazal-like domains proteoglycan 1	456265 IMAGE Consortium	AI894071
8925 B	6 021MAMO0	trefoil factor 2 (spasmolytic protein 1)	438574 IMAGE Consortium	AI894032
8926 B	8 021MAMO0	polymerase, gamma 2	581961 IMAGE Consortium	AA154131
8927 B	10 021MAMO0	zinc finger homeobox 1a	620244 IMAGE Consortium	AA178068
8928 B	12 021MAMO0	CD4 antigen	641964 IMAGE Consortium	AA210044
8929 D	2 021MAMO0	ectonucleoside triphosphate diphosphohydrolase 1	402959 IMAGE Consortium	W80311
8930 D	4 021MAMO0	Mus musculus major histocompatibility locus class III region:butyrophilin-like protein gene, partial cds; Notch4, PBX2, RAGE, lysophatidic acid acyl transferase-alpha, palmitoyl-protein thioesterase 2 (PPT2), CREB-RP, and tenascin	475474 IMAGE Consortium	AA050074
8931 D	6 021MAMO0	protein C receptor, endothelial	444429 IMAGE Consortium	AA016910
8932 D	8 021MAMO0	ESTs, Moderately similar to DNA NUCLEOTIDYLEXOTRANSFERASE [Bos taurus]	583353 IMAGE Consortium	AI326745
8933 D	10 021MAMO0	tissue inhibitor of metalloproteinase	622732 IMAGE Consortium	AA184223
8934 D	12 021MAMO0	breast cancer 1	643280 IMAGE Consortium	AA189699
8935 F	2 021MAMO0	manic fringe homolog (Drosophila)	408533 IMAGE Consortium	AA052104



8936 F	4 021MAM00	proprotein convertase subtilisin/kexin type 2	477515 IMAGE Consortium	AA048432
8937 F	6 021MAM00	sema domain, transmembrane domain (TM), and cytoplasmic domain, (semaphorin) 6A	456781 IMAGE Consortium	AA023538
8938 F	8 021MAM00	BMX non-receptor tyrosine kinase	596729 IMAGE Consortium	AA138327
8939 F	10 021MAM00	kinesin family member C2	634870 IMAGE Consortium	AA154770
8940 F	12 021MAM00	cytoplasmic tyrosine kinase, Dscr28C related (Drosophila)	653787 IMAGE Consortium	AA218066
8941 H	2 021MAM00	ral guanine nucleotide dissociation stimulator,-like 1	422579 IMAGE Consortium	AI894000
8942 H	4 021MAM00	testis specific protein kinase 1	477798 IMAGE Consortium	AA048413
8943 H	6 021MAM00	ESTs	463011 IMAGE Consortium	AI325350
8944 H	8 021MAM00	ESTs, Highly similar to MOX2_MOUSE HOMEBOX PROTEIN MOX-2 [M.musculus]	597262 IMAGE Consortium	AI324355
8945 H	10 021MAM00	chemokine (C-C) receptor 2	635981 IMAGE Consortium	AA183002
8946 H	12 021MAM00	forkhead box A1	659086 IMAGE Consortium	AA217937
8947 B	2 021TAM01	CEA-related cell adhesion molecule 1	680715 IMAGE Consortium	AA242406
8948 B	4 021TAM01	cytochrome c oxidase, subunit VIIa 1	334181 IMAGE Consortium	W16266
8949 B	6 021TAM01	retinol dehydrogenase type 5	692779 IMAGE Consortium	AA239932
8950 B	8 021TAM01	immunoglobulin mu binding protein 2	764717 IMAGE Consortium	AA274243
8951 B	10 021TAM01	NK2 transcription factor related, locus 2 (Drosophila)	406655 IMAGE Consortium	W83713
8952 B	12 021TAM01	COMPLEMENT FACTOR H PRECURSOR	681810 IMAGE Consortium	AA244704
8953 D	2 021TAM01	complement component 4 binding protein	681482 IMAGE Consortium	AA238308
8954 D	4 021TAM01	solute carrier family 28 (sodium-coupled nucleoside transporter), member 2	349446 IMAGE Consortium	AI391105
8955 D	6 021TAM01	cytochrome P450, 1b1, benz[a]anthracene inducible	720892 IMAGE Consortium	AA267178
8956 D	8 021TAM01	FBJ osteosarcoma oncogene B	765039 IMAGE Consortium	AA273581
8957 D	10 021TAM01	Rhesus blood group-like	439864 IMAGE Consortium	AA008551
8958 D	12 021TAM01	CD38 antigen	750428 IMAGE Consortium	AA472876
8959 F	2 021TAM01	glucagon receptor	681657 IMAGE Consortium	AA237855
8960 F	4 021TAM01	glycoprotein 9 (platelet)	355508 IMAGE Consortium	W48187
8961 F	6 021TAM01	glycophorin A	734015 IMAGE Consortium	AA272078
8962 F	8 021TAM01	lymphocyte antigen 9	777346 IMAGE Consortium	AA276920
8963 F	10 021TAM01	vesicle transport through interaction with t-SNAREs 1 homolog	444735 IMAGE Consortium	AA016934
8964 F	12 021TAM01	neurofilament, heavy polypeptide	639382 IMAGE Consortium	AA200799
8965 H	2 021TAM01	solute carrier family 1, member 2	693294 IMAGE Consortium	AA397107
8966 H	4 021TAM01	recombination activating gene 1 gene activation	390128 IMAGE Consortium	W71636
8967 H	6 021TAM01	zinc finger protein 46	735790 IMAGE Consortium	AA269904
8968 H	8 021TAM01	amine oxidase, copper containing 3	820177 IMAGE Consortium	AA423217
8969 H	10 021TAM01	glucokinase activity	468297 IMAGE Consortium	AI325384
8970 H	12 021TAM01	ESTs	638216 IMAGE Consortium	AI646670
8971 B	2 0210AM02	ESTs, Weakly similar to S57968 Ran-binding protein 2 - mouse [M.musculus]	639263 IMAGE Consortium	AI645951
8972 B	4 0210AM02	ESTs	639781 IMAGE Consortium	AA197547
8973 B	6 0210AM02	ESTs	640707 IMAGE Consortium	AA203872

8974 B	8 0210AMO2	ESTs	636252 IMAGE Consortium	AI645156
8975 B	10 0210AMO2	ESTs	636454 IMAGE Consortium	AI645182
8976 B	12 0210AMO2	ESTs	636642 IMAGE Consortium	AI646620
8977 D	2 0210AMO2	ESTs	637814 IMAGE Consortium	AI509349
8978 D	4 0210AMO2	Mus musculus K20D4 (K20D4) mRNA, complete cds	639797 IMAGE Consortium	AA197557
8979 D	6 0210AMO2	ESTs	640733 IMAGE Consortium	AA203881
8980 D	8 0210AMO2	ESTs	637017 IMAGE Consortium	AI645230
8981 D	10 0210AMO2	ESTs	637208 IMAGE Consortium	AA189544
8982 D	12 0210AMO2	ESTs	636657 IMAGE Consortium	AI646624
8983 F	2 0210AMO2	ESTs	637832 IMAGE Consortium	AA189752
8984 F	4 0210AMO2	ESTs	640069 IMAGE Consortium	AA197355
8985 F	6 0210AMO2	ESTs	636738 IMAGE Consortium	AA182342
8986 F	8 0210AMO2	etoile	637059 IMAGE Consortium	AA189288
8987 F	10 0210AMO2	ESTs	637214 IMAGE Consortium	AI645262
8988 F	12 0210AMO2	ESTs	636674 IMAGE Consortium	AI646627
8989 H	2 0210AMO2	ESTs	637864 IMAGE Consortium	AA189571
8990 H	4 0210AMO2	lunatic fringe gene homolog (Drosophila)	640133 IMAGE Consortium	AA197529
8991 H	6 0210AMO2	ESTs	636838 IMAGE Consortium	AI645202
8992 H	8 0210AMO2	ESTs	637072 IMAGE Consortium	AI645238
8993 H	10 0210AMO2	RAB4A, member RAS oncogene family	636512 IMAGE Consortium	AA182236
8994 H	12 0210AMO2	ESTs	640725 IMAGE Consortium	AA204409
8995 B	2 0217AMO3	Mus musculus mRNA for beta chimaerin (bch gene)	642415 IMAGE Consortium	AI606014
8996 B	4 0217AMO3	ESTs	642612 IMAGE Consortium	AA184890
8997 B	6 0217AMO3	ESTs, Highly similar to PYRUVATE DEHYDROGENASE E1 COMPONENT, BETA SUBUNIT PRECURSOR [Rattus norvegicus]	347856 IMAGE Consortium	AI391020
8998 B	8 0217AMO3	ESTs	643155 IMAGE Consortium	AI627090
8999 B	10 0217AMO3			
9000 B	12 0217AMO3	ESTs, Moderately similar to AF151838_1 CGI-80 protein [H.sapiens]	644301 IMAGE Consortium	AI450449
9001 D	2 0217AMO3	ESTs	642456 IMAGE Consortium	AI606018
9002 D	4 0217AMO3	ESTs	642682 IMAGE Consortium	AA184930
9003 D	6 0217AMO3	f-box only protein 14	336022 IMAGE Consortium	W36379
9004 D	8 0217AMO3	ESTs	643167 IMAGE Consortium	AI627091
9005 D	10 0217AMO3	ESTs, Weakly similar to XY40 protein [R.norvegicus]	643455 IMAGE Consortium	AA185989
9006 D	12 0217AMO3	ESTs	643611 IMAGE Consortium	AI390462
9007 F	2 0217AMO3	ESTs	642488 IMAGE Consortium	AI606021
9008 F	4 0217AMO3	ESTs	336672 IMAGE Consortium	W36776
9009 F	6 0217AMO3	ESTs	336150 IMAGE Consortium	AI390168
9010 F	8 0217AMO3	ESTs	643248 IMAGE Consortium	AI627100
9011 F	10 0217AMO3	ESTs	642750 IMAGE Consortium	AA184976
9012 F	12 0217AMO3	ESTs	644350 IMAGE Consortium	AA204513

9013 H	2	0217AMO3	EST	642494	IMAGE Consortium	AI451550
9014 H	4	0217AMO3	EST	337426	IMAGE Consortium	AI415099
9015 H	6	0217AMO3	ARS component B gene	337123	IMAGE Consortium	W20775
9016 H	8	0217AMO3	ESTs	643306	IMAGE Consortium	AI627109
9017 H	10	0217AMO3	ESTs	642775	IMAGE Consortium	AI627053
9018 H	12	0217AMO3	ESTs	643649	IMAGE Consortium	AA200316
9019 B	2	021EAMO4	ESTs	643719	IMAGE Consortium	AI390474
9020 B	4	021EAMO4	ESTs	659313	IMAGE Consortium	AA218015
9021 B	6	021EAMO4	ESTs	659710	IMAGE Consortium	AA220093
9022 B	8	021EAMO4	ESTs	669810	IMAGE Consortium	AA230635
9023 B	10	021EAMO4	ESTs	653475	IMAGE Consortium	AI390732
9024 B	12	021EAMO4	ESTs	658320	IMAGE Consortium	AA198486
9025 D	2	021EAMO4	ESTs	643824	IMAGE Consortium	AI450413
9026 D	4	021EAMO4	ESTs, Highly similar to WNT-5B PROTEIN PRECURSOR [Mus musculus]	659366	IMAGE Consortium	AA218013
9027 D	6	021EAMO4	ESTs	659726	IMAGE Consortium	AA220036
9028 D	8	021EAMO4	retinoic acid receptor, gamma	669879	IMAGE Consortium	AA230517
9029 D	10	021EAMO4	ESTs	652771	IMAGE Consortium	AI390653
9030 D	12	021EAMO4	ESTs	653037	IMAGE Consortium	AI390682
9031 F	2	021EAMO4	ESTs	643855	IMAGE Consortium	AA204265
9032 F	4	021EAMO4	ESTs, Weakly similar to IER5 [M.musculus]	658686	IMAGE Consortium	AI390417
9033 F	6	021EAMO4	calcium/calmodulin-dependent serine protein kinase	659937	IMAGE Consortium	AI390207
9034 F	8	021EAMO4	ESTs	660540	IMAGE Consortium	AI390228
9035 F	10	021EAMO4	ESTs	652807	IMAGE Consortium	AA217469
9036 F	12	021EAMO4	ESTs	653056	IMAGE Consortium	AI390685
9037 H	2	021EAMO4	ESTs, Moderately similar to I37275 death-associated protein kinase [H.sapiens]	643889	IMAGE Consortium	AA203937
9038 H	4	021EAMO4	ESTs	659527	IMAGE Consortium	AI644835
9039 H	6	021EAMO4	ESTs	660062	IMAGE Consortium	AI390212
9040 H	8	021EAMO4	ESTs	669986	IMAGE Consortium	AA253844
9041 H	10	021EAMO4	ESTs	652861	IMAGE Consortium	AI390664
9042 H	12	021EAMO4	ESTs, Moderately similar to unnamed protein product [H.sapiens]	653170	IMAGE Consortium	AA217311
9043 B	2	021LAMO5	ESTs	652630	IMAGE Consortium	AI390588
9044 B	4	021LAMO5	EST X83306	680030	IMAGE Consortium	AA242091
9045 B	6	021LAMO5	ESTs	671739	IMAGE Consortium	AI391347
9046 B	8	021LAMO5	ESTs, Weakly similar to A48998 nucleolar protein p120 - mouse [M.musculus]	679473	IMAGE Consortium	AA244646
9047 B	10	021LAMO5	bone morphogenetic protein 5	681111	IMAGE Consortium	AA242542
9048 B	12	021LAMO5	ESTs	681616	IMAGE Consortium	AA237978
9049 D	2	021LAMO5	ESTs	652655	IMAGE Consortium	AI390593
9050 D	4	021LAMO5	mab-21-like 2 (C. elegans)	672060	IMAGE Consortium	AA245183
9051 D	6	021LAMO5	ESTs, Weakly similar to AF177041_1 aldo-keto reductase	677021	IMAGE Consortium	AI390246

		a [M.musculus]		
9052 D	8 021LAMO5	ESTs	681306 IMAGE Consortium	AA242421
9053 D	10 021LAMO5	ESTs	681270 IMAGE Consortium	AA244951
9054 D	12 021LAMO5	ESTs	681644 IMAGE Consortium	AA237987
9055 F	2 021LAMO5	ESTs	652684 IMAGE Consortium	AI390598
9056 F	4 021LAMO5	ESTs, Weakly similar to zinc finger protein 95 [M.musculus]	680242 IMAGE Consortium	AA237521
9057 F	6 021LAMO5	ESTs, Weakly similar to I48668 zinc finger protein 51 - mouse [M.musculus]	677214 IMAGE Consortium	AA213131
9058 F	8 021LAMO5	villin	697307 IMAGE Consortium	AA242102
9059 F	10 021LAMO5	CD52 antigen	680584 IMAGE Consortium	AA238062
9060 F	12 021LAMO5	ESTs	680928 IMAGE Consortium	AI451864
9061 H	2 021LAMO5	ESTs, Highly similar to retinoblastoma-interacting protein [H.sapiens]	653426 IMAGE Consortium	AI390725
9062 H	4 021LAMO5	sema domain, immunoglobulin domain (Ig), transmembrane domain (TM) and short cytoplasmic domain, (semaphorin) 4G	680284 IMAGE Consortium	AA238294
9063 H	6 021LAMO5	ESTs	676623 IMAGE Consortium	AI391425
9064 H	8 021LAMO5	Socs-5	697031 IMAGE Consortium	AI425448
9065 H	10 021LAMO5	ESTs	681328 IMAGE Consortium	AA242433
9066 H	12 021LAMO5	Bcl2-associated athanogene 3	681707 IMAGE Consortium	AA245702
9067 B	2 021SAMO6	alcohol dehydrogenase 4 (class II), pi polypeptide	692690 IMAGE Consortium	AI390641
9068 B	4 021SAMO6	ESTs	719175 IMAGE Consortium	AA267319
9069 B	6 021SAMO6	SH2 domain protein 1A	719479 IMAGE Consortium	AI464772
9070 B	8 021SAMO6	S100 calcium-binding protein A9 (calgranulin B)	720190 IMAGE Consortium	AA255025
9071 B	10 021SAMO6	DNA segment, Chr 1, ERATO Doi 622, expressed	718373 IMAGE Consortium	AA267725
9072 B	12 021SAMO6	ESTs	718611 IMAGE Consortium	AI464630
9073 D	2 021SAMO6	ESTs	693498 IMAGE Consortium	AA239380
9074 D	4 021SAMO6	ESTs	719210 IMAGE Consortium	AA254541
9075 D	6 021SAMO6	ESTs	719490 IMAGE Consortium	AI464771
9076 D	8 021SAMO6	ESTs	720277 IMAGE Consortium	AI464812
9077 D	10 021SAMO6	ESTs, Weakly similar to KIAA0946 protein [H.sapiens]	717648 IMAGE Consortium	AA250335
9078 D	12 021SAMO6	ESTs	718648 IMAGE Consortium	AI464634
9079 F	2 021SAMO6	vanin 3	692993 IMAGE Consortium	AA238409
9080 F	4 021SAMO6	ESTs	719254 IMAGE Consortium	AI464736
9081 F	6 021SAMO6	ESTs	719591 IMAGE Consortium	AI464778
9082 F	8 021SAMO6	ESTs, Weakly similar to unnamed protein product [M.musculus]	720388 IMAGE Consortium	AI464818
9083 F	10 021SAMO6	ESTs	718399 IMAGE Consortium	AA267733
9084 F	12 021SAMO6	ESTs	718671 IMAGE Consortium	AA265639
9085 H	2 021SAMO6	ESTs	693736 IMAGE Consortium	AA265096
9086 H	4 021SAMO6	ESTs	719296 IMAGE Consortium	AI464745
9087 H	6 021SAMO6	ESTs	718878 IMAGE Consortium	AA265821

9088 H	8 021SAMO6	ESTs	720591 IMAGE Consortium	AA261404
9089 H	10 021SAMO6	ESTs	718431 IMAGE Consortium	AI464603
9090 H	12 021SAMO6	ESTs	718711 IMAGE Consortium	AI464652
9091 B	2 021ZAMO7	ESTs	717127 IMAGE Consortium	AI466871
9092 B	4 021ZAMO7	ESTs	721080 IMAGE Consortium	AI464897
9093 B	6 021ZAMO7	ESTs	721383 IMAGE Consortium	AA267259
9094 B	8 021ZAMO7	ESTs	721539 IMAGE Consortium	AI464956
9095 B	10 021ZAMO7	ESTs	720983 IMAGE Consortium	AA266871
9096 B	12 021ZAMO7	ESTs	733870 IMAGE Consortium	AI529431
9097 D	2 021ZAMO7	ESTs	717171 IMAGE Consortium	AI466873
9098 D	4 021ZAMO7	ESTs	721120 IMAGE Consortium	AI464901
9099 D	6 021ZAMO7	ESTs	722104 IMAGE Consortium	AA268576
9100 D	8 021ZAMO7	ESTs	721577 IMAGE Consortium	AA267673
9101 D	10 021ZAMO7	ESTs	720995 IMAGE Consortium	AI464881
9102 D	12 021ZAMO7	intestine N-acetylglucosamine 6-O-sulfotransferase	734674 IMAGE Consortium	AI529474
9103 F	2 021ZAMO7	ESTs	718013 IMAGE Consortium	AA261547
9104 F	4 021ZAMO7	ESTs	721170 IMAGE Consortium	AI464903
9105 F	6 021ZAMO7	dopa/tyrosine sulfotransferase	721406 IMAGE Consortium	AI464939
9106 F	8 021ZAMO7	ESTs	720858 IMAGE Consortium	AI464856
9107 F	10 021ZAMO7	ESTs	721726 IMAGE Consortium	AI595294
9108 F	12 021ZAMO7	ESTs	733965 IMAGE Consortium	AI529444
9109 H	2 021ZAMO7	ESTs	718051 IMAGE Consortium	AA261194
9110 H	4 021ZAMO7	N-methylpurine-DNA glycosylase	721243 IMAGE Consortium	AI464913
9111 H	6 021ZAMO7	ESTs	721433 IMAGE Consortium	AI464942
9112 H	8 021ZAMO7	vimentin	720884 IMAGE Consortium	AI464862
9113 H	10 021ZAMO7	ESTs	721744 IMAGE Consortium	AI595297
9114 H	12 021ZAMO7	LIM homeobox protein 2	734707 IMAGE Consortium	AA259281
9115 B	2 0216AMO8	ESTs	722346 IMAGE Consortium	AI608085
9116 B	4 0216AMO8	ESTs	723270 IMAGE Consortium	AI463762
9117 B	6 0216AMO8	ESTs	722876 IMAGE Consortium	AI608143
9118 B	8 0216AMO8	ESTs	749035 IMAGE Consortium	AA288152
9119 B	10 0216AMO8	ESTs, Weakly similar to AF151830_1 CGI-72 protein [H.sapiens]	749402 IMAGE Consortium	AA289555
9120 B	12 0216AMO8	ESTs	749784 IMAGE Consortium	AA396292
9121 D	2 0216AMO8	ESTs	722356 IMAGE Consortium	AA254255
9122 D	4 0216AMO8	EST	723279 IMAGE Consortium	AI661398
9123 D	6 0216AMO8	ESTs, Weakly similar to sex-determination protein homolog Fem1a [M.musculus]	722935 IMAGE Consortium	AI608162
9124 D	8 0216AMO8	ESTs	749100 IMAGE Consortium	AI644772
9125 D	10 0216AMO8	epithelial V-like antigen	748712 IMAGE Consortium	AA288427
9126 D	12 0216AMO8	ESTs	749882 IMAGE Consortium	AA388662
9127 F	2 0216AMO8	ESTs	722366 IMAGE Consortium	AI608089

9128 F	4 0216AMO8	ESTs	723293 IMAGE Consortium	AA267824
9129 F	6 0216AMO8	ESTs	734149 IMAGE Consortium	AA268607
9130 F	8 0216AMO8	ESTs, Highly similar to AF161493_1 HSPC144 [H.sapiens]	749136 IMAGE Consortium	AA290555
9131 F	10 0216AMO8	ESTs	749542 IMAGE Consortium	AA288278
9132 F	12 0216AMO8	ESTs	749925 IMAGE Consortium	AA389281
9133 H	2 0216AMO8	ESTs	722408 IMAGE Consortium	AI608093
9134 H	4 0216AMO8	ESTs	723318 IMAGE Consortium	AA267871
9135 H	6 0216AMO8	ESTs	734205 IMAGE Consortium	AA268634
9136 H	8 0216AMO8	ESTs	749260 IMAGE Consortium	AI644788
9137 H	10 0216AMO8	ESTs	749564 IMAGE Consortium	AA289741
9138 H	12 0216AMO8	ESTs, Moderately similar to hT41 [H.sapiens]	749994 IMAGE Consortium	AA388951
9139 B	2 021DAMO9	fatty acid binding protein 7, brain	334095 IMAGE Consortium	W16203
9140 B	4 021DAMO9	ESTs, Weakly similar to GLYCOGEN DEBRANCHING ENZYME [Homo sapiens]	335260 IMAGE Consortium	AI322281
9141 B	6 021DAMO9	ESTs, Weakly similar to T00068 hypothetical protein KIAA0443 - human [H.sapiens]	736432 IMAGE Consortium	AI463878
9142 B	8 021DAMO9	ESTs	746549 IMAGE Consortium	AI607675
9143 B	10 021DAMO9	ESTs, Weakly similar to EPS8_MOUSE EPIDERMAL GROWTH FACTOR RECEPTOR KINASE SUBSTRATE EPS8 [M.musculus]	751279 IMAGE Consortium	AA290368
9144 B	12 021DAMO9	ESTs	751505 IMAGE Consortium	AI508727
9145 D	2 021DAMO9	ESTs	334238 IMAGE Consortium	AI391288
9146 D	4 021DAMO9	crystallin, gamma C	334649 IMAGE Consortium	W19024
9147 D	6 021DAMO9	growth factor independent 1B	735830 IMAGE Consortium	AI463842
9148 D	8 021DAMO9	ESTs, Moderately similar to similar to C.elegans protein [H.sapiens]	746670 IMAGE Consortium	AI607681
9149 D	10 021DAMO9	Mus musculus (strain C57Bl/6) mRNA sequence	750569 IMAGE Consortium	AA472976
9150 D	12 021DAMO9	ESTs, Weakly similar to anti-Fas-induced apoptosis [H.sapiens]	750847 IMAGE Consortium	AI508502
9151 F	2 021DAMO9	ESTs	335076 IMAGE Consortium	AI390099
9152 F	4 021DAMO9	ESTs	334680 IMAGE Consortium	W19015
9153 F	6 021DAMO9	ESTs	736632 IMAGE Consortium	AI463891
9154 F	8 021DAMO9	ESTs, Weakly similar to hypothetical protein [H.sapiens]	746875 IMAGE Consortium	AA274891
9155 F	10 021DAMO9	ESTs	750590 IMAGE Consortium	AA472950
9156 F	12 021DAMO9	ESTs, Moderately similar to CYP2C40 [M.musculus]	751595 IMAGE Consortium	AA396053
9157 H	2 021DAMO9	ESTs, Highly similar to unknown [H.sapiens]	335078 IMAGE Consortium	AI390100
9158 H	4 021DAMO9	ESTs, Weakly similar to podocalyxin-like protein 1 [M.musculus]	334816 IMAGE Consortium	W36438
9159 H	6 021DAMO9	ESTs	738204 IMAGE Consortium	AA270832
9160 H	8 021DAMO9	DNA polymerase alpha 1, 180 kDa	751125 IMAGE Consortium	AA288474
9161 H	10 021DAMO9	ESTs	751316 IMAGE Consortium	AI508703
9162 H	12 021DAMO9	ESTs, Moderately similar to unnamed protein product	750896 IMAGE Consortium	AA290276

		[H.sapiens]		
9163 B	2 021XAMOA	ESTs	751663 IMAGE Consortium	AA396100
9164 B	4 021XAMOA	ESTs	750285 IMAGE Consortium	AI606539
9165 B	6 021XAMOA	ESTs, Highly similar to DRNG_MOUSE DEOXYRIBONUCLEASE GAMMA PRECURSOR	752217 IMAGE Consortium	AA289054
		[M.musculus]		
9166 B	8 021XAMOA	ESTs	762234 IMAGE Consortium	AI509003
9167 B	10 021XAMOA	ESTs	764113 IMAGE Consortium	AA285993
9168 B	12 021XAMOA	ESTs	764395 IMAGE Consortium	AA277178
9169 D	2 021XAMOA	ESTs	751668 IMAGE Consortium	AA396089
9170 D	4 021XAMOA	ESTs	751012 IMAGE Consortium	AI508523
9171 D	6 021XAMOA	ESTs	752244 IMAGE Consortium	AI508570
9172 D	8 021XAMOA	ESTs	762309 IMAGE Consortium	AI595030
9173 D	10 021XAMOA	ESTs	764151 IMAGE Consortium	AA285835
9174 D	12 021XAMOA	ESTs	762993 IMAGE Consortium	AI595068
9175 F	2 021XAMOA	cholesterol 25-hydroxylase	750976 IMAGE Consortium	AA289153
9176 F	4 021XAMOA	Mus musculus germline J and C regions of immuoglobulin lambda-2 and germline J and C regions of the immuoglobulin lambda-4 pseudo-gene	751026 IMAGE Consortium	AA289453
9177 F	6 021XAMOA	ESTs	752309 IMAGE Consortium	AI508578
9178 F	8 021XAMOA	ESTs	762365 IMAGE Consortium	AI595033
9179 F	10 021XAMOA	ESTs, Weakly similar to AF113615_1 FH1/FH2 domain- containing protein FHOS [H.sapiens]	764169 IMAGE Consortium	AI606558
9180 F	12 021XAMOA	ESTs	764472 IMAGE Consortium	AA274113
9181 H	2 021XAMOA	ESTs	751784 IMAGE Consortium	AA396195
9182 H	4 021XAMOA	ESTs	750332 IMAGE Consortium	AI606544
9183 H	6 021XAMOA	RAB geranylgeranyl transferase, b subunit	752360 IMAGE Consortium	AA290003
9184 H	8 021XAMOA	ESTs	763119 IMAGE Consortium	AA271256
9185 H	10 021XAMOA	ESTs	764182 IMAGE Consortium	AI606556
9186 H	12 021XAMOA	ESTs	778290 IMAGE Consortium	AA413858
9187 B	2 0214AMOB	M.musculus mRNA for transferrin receptor	778391 IMAGE Consortium	AA413490
9188 B	4 0214AMOB	ligase III, DNA, ATP-dependent	778550 IMAGE Consortium	AA413595
9189 B	6 0214AMOB	mannosidase 2, alpha 1	778857 IMAGE Consortium	AA414150
9190 B	8 0214AMOB	ESTs	777405 IMAGE Consortium	AA276580
9191 B	10 0214AMOB	ESTs	765105 IMAGE Consortium	AI510250
9192 B	12 0214AMOB	ESTs	764826 IMAGE Consortium	AA274074
9193 D	2 0214AMOB	DNA segment, Chr 10, ERATO Doi 802, expressed	779164 IMAGE Consortium	AA467574
9194 D	4 0214AMOB	cyclin G2	778625 IMAGE Consortium	AA413576
9195 D	6 0214AMOB	ESTs, Highly similar to AC004858_3 U1 small ribonucleoprotein 1SNRP homolog [H.sapiens]	778928 IMAGE Consortium	AA414279
9196 D	8 0214AMOB	ESTs	777458 IMAGE Consortium	AA276616
9197 D	10 0214AMOB	ESTs	765244 IMAGE Consortium	AA273297
9198 D	12 0214AMOB	ESTs	765796 IMAGE Consortium	AA275356

9199 F	2 0214AMOB	ESTs, Highly similar to CYCLIN-DEPENDENT KINASES REGULATORY SUBUNIT 2 [Homo sapiens]	778476 IMAGE Consortium	AA413740
9200 F	4 0214AMOB	ESTs, Weakly similar to FLI-LRR associated protein-1 [M.musculus]	778633 IMAGE Consortium	AA413640
9201 F	6 0214AMOB	ESTs	778952 IMAGE Consortium	AA414311
9202 F	8 0214AMOB	ESTs	777536 IMAGE Consortium	AA276747
9203 F	10 0214AMOB	ESTs	765290 IMAGE Consortium	AA275143
9204 F	12 0214AMOB	ESTs	765928 IMAGE Consortium	AA277899
9205 H	2 0214AMOB	replication protein A2	778482 IMAGE Consortium	AA413512
9206 H	4 0214AMOB	hypothetical protein	778680 IMAGE Consortium	AA413600
9207 H	6 0214AMOB	zinc finger protein, subfamily 1A, 1 (Ikaros)	765132 IMAGE Consortium	AA273924
9208 H	8 0214AMOB	leptin receptor	777547 IMAGE Consortium	AA286541
9209 H	10 0214AMOB	ESTs	765309 IMAGE Consortium	AA275241
9210 H	12 0214AMOB	ESTs	765956 IMAGE Consortium	AA277951
9211 B	2 021BAMOC	ESTs	779033 IMAGE Consortium	AA413766
9212 B	4 021BAMOC	immunophilin-like protein NG7	779254 IMAGE Consortium	AA445853
9213 B	6 021BAMOC	peroxisomal delta3, delta2-enoyl-Coenzyme A isomerase	779426 IMAGE Consortium	AA414193
9214 B	8 021BAMOC	ESTs	779658 IMAGE Consortium	AA466936
9215 B	10 021BAMOC	ESTs, Weakly similar to TRANSLATION INITIATION FACTOR EIF-3 P39 SUBUNIT [Saccharomyces cerevisiae]	779995 IMAGE Consortium	AA414888
9216 B	12 021BAMOC	glutamate oxaloacetate transaminase 1, soluble	791106 IMAGE Consortium	AA415254
9217 D	2 021BAMOC	ESTs	779038 IMAGE Consortium	AA413765
9218 D	4 021BAMOC	ESTs, Highly similar to CCHL_HUMAN CYTOCHROME C-TYPE HEME LYASE [H.sapiens]	779297 IMAGE Consortium	AA445869
9219 D	6 021BAMOC	ESTs, Weakly similar to PRP3_MOUSE PROLINE-RICH PROTEIN MP-3 [M.musculus]	779429 IMAGE Consortium	AA414200
9220 D	8 021BAMOC	ESTs, Weakly similar to LA_RAT LUPUS LA PROTEIN HOMOLOG [R.norvegicus]	779672 IMAGE Consortium	AA413852
9221 D	10 021BAMOC	ESTs	780075 IMAGE Consortium	AA414696
9222 D	12 021BAMOC	EST AI325948	791117 IMAGE Consortium	AA415265
9223 F	2 021BAMOC	ESTs	779057 IMAGE Consortium	AA413794
9224 F	4 021BAMOC	zinc finger protein 148	779320 IMAGE Consortium	AA414802
9225 F	6 021BAMOC	upstream binding protein 1	779457 IMAGE Consortium	AA414213
9226 F	8 021BAMOC	ESTs	779705 IMAGE Consortium	AA414629
9227 F	10 021BAMOC	leukotriene A4 hydrolase	780114 IMAGE Consortium	AA414790
9228 F	12 021BAMOC	superoxide dismutase 2, mitochondrial	791140 IMAGE Consortium	AA415267
9229 H	2 021BAMOC	ESTs, Weakly similar to S37583 RING finger protein rfp - mouse [M.musculus]	779073 IMAGE Consortium	AA414078
9230 H	4 021BAMOC	ESTs, Highly similar to HYPOTHETICAL 167.8 KD PROTEIN CCE1-CAP1 INTERGENIC REGION [Saccharomyces cerevisiae]	779345 IMAGE Consortium	AA414812
9231 H	6 021BAMOC	protein phosphatase 2, regulatory subunit B (B56),	779461 IMAGE Consortium	AA414215



9232 H	8 021BAMOC	gamma isoform ESTs	779493 IMAGE Consortium	AA414491
9233 H	10 021BAMOC	ethanol induced 1	791008 IMAGE Consortium	AA473948
9234 H	12 021BAMOC	adenomatosis polyposis coli binding protein Eb1	791155 IMAGE Consortium	AA415514
9235 B	2 021IAMOD	GA repeat binding protein, alpha	791281 IMAGE Consortium	AA420357
9236 B	4 021IAMOD	ESTs, Highly similar to AF151078_1 HSPC244 [H.sapiens]	790952 IMAGE Consortium	AA415323
9237 B	6 021IAMOD	ESTs, Weakly similar to CRTC_MOUSE CALRETICULIN PRECURSOR [M.musculus]	791900 IMAGE Consortium	AA414421
9238 B	8 021IAMOD	ESTs, Highly similar to TIP120 [R.norvegicus]	792889 IMAGE Consortium	AA415523
9239 B	10 021IAMOD	Mus musculus mRNA for SSeCKS, complete cds	793039 IMAGE Consortium	AA444232
9240 B	12 021IAMOD	steroid dehydrogenase	791955 IMAGE Consortium	AA414292
9241 D	2 021IAMOD	G protein-coupled receptor 37	790595 IMAGE Consortium	AA289047
9242 D	4 021IAMOD	ESTs	790977 IMAGE Consortium	AA415361
9243 D	6 021IAMOD	Mus musculus E-cadherin binding protein E7 mRNA, complete cds	792644 IMAGE Consortium	AA416395
9244 D	8 021IAMOD	ESTs, Moderately similar to AF151064_1 HSPC230 [H.sapiens]	792914 IMAGE Consortium	AA415528
9245 D	10 021IAMOD	ESTs, Weakly similar to dJ126A5.2.1 [H.sapiens]	793106 IMAGE Consortium	AA437864
9246 D	12 021IAMOD	protein phosphatase 1B, magnesium dependent, beta isoform	791985 IMAGE Consortium	AA414499
9247 F	2 021IAMOD	phosphatidylinositol glycan, class N	790626 IMAGE Consortium	AA289117
9248 F	4 021IAMOD	ESTs, Weakly similar to unnamed protein product [M.musculus]	791827 IMAGE Consortium	AA414381
9249 F	6 021IAMOD	Mus musculus 15 kDa selenoprotein (Sep15) mRNA, complete cds	791920 IMAGE Consortium	AA414427
9250 F	8 021IAMOD	serum response factor	792937 IMAGE Consortium	AA415341
9251 F	10 021IAMOD	ESTs, Highly similar to UDP-N-acetylglucosamine pyrophosphorylase [H.sapiens]	793128 IMAGE Consortium	AA437972
9252 F	12 021IAMOD	DNA segment, Chr 15, ERATO Doi 806, expressed	791323 IMAGE Consortium	AA473957
9253 H	2 021IAMOD	ESTs, Weakly similar to splicing factor SC35 [M.musculus]	790765 IMAGE Consortium	AA388448
9254 H	4 021IAMOD	fibulin 1	792563 IMAGE Consortium	AA416352
9255 H	6 021IAMOD	protein phosphatase 1B, magnesium dependent, beta isoform	792656 IMAGE Consortium	AA416338
9256 H	8 021IAMOD	WSB-2 protein	792939 IMAGE Consortium	AA415342
9257 H	10 021IAMOD	baculoviral IAP repeat-containing 4	793152 IMAGE Consortium	AA437994
9258 H	12 021IAMOD	poly(rC)-binding protein 1	791346 IMAGE Consortium	AA473961
9259 B	2 021PAMOE	TATA box binding protein (Tbp)-associated factor, RNA polymerase I, C	791417 IMAGE Consortium	AA416401
9260 B	4 021PAMOE	Mus musculus B-cell activating factor (Baff) mRNA, complete cds	791670 IMAGE Consortium	AA437602
9261 B	6 021PAMOE	ESTs	792855 IMAGE Consortium	AA415560
9262 B	8 021PAMOE	ESTs, Highly similar to RIBOSOMAL PROTEIN S6	793754 IMAGE Consortium	AA415545

		KINASE [Homo sapiens]		
9263 B	10 021PAMOE	serine/threonine kinase 18	793053 IMAGE Consortium	AA444238
9264 B	12 021PAMOE	protein phosphatase 1, catalytic subunit, beta isoform	793195 IMAGE Consortium	AA438025
9265 D	2 021PAMOE	ESTs	791446 IMAGE Consortium	AA416413
9266 D	4 021PAMOE	ESTs, Weakly similar to BACR37P7.g [D.melanogaster]	791687 IMAGE Consortium	AA474020
9267 D	6 021PAMOE	ESTs, Weakly similar to PRAJA1 [M.musculus]	792875 IMAGE Consortium	AA415565
9268 D	8 021PAMOE	ESTs, Highly similar to CBP_MOUSE CREB-BINDING PROTEIN□ [M.musculus]	793853 IMAGE Consortium	AA422662
9269 D	10 021PAMOE	ornithine decarboxylase, structural	793067 IMAGE Consortium	AA444254
9270 D	12 021PAMOE	uncoupling protein, mitochondrial	352000 IMAGE Consortium	W41372
9271 F	2 021PAMOE	ESTs	791502 IMAGE Consortium	AA437424
9272 F	4 021PAMOE	mitogen activated protein kinase kinase 4	791764 IMAGE Consortium	AA414670
9273 F	6 021PAMOE	ESTs	793607 IMAGE Consortium	AA415800
9274 F	8 021PAMOE	bone morphogenetic protein 15	793859 IMAGE Consortium	AA422665
9275 F	10 021PAMOE	cyclin E	793114 IMAGE Consortium	AA437868
9276 F	12 021PAMOE	wingless related MMTV integration site 2b	353765 IMAGE Consortium	W44010
9277 H	2 021PAMOE	antigenic determinant of rec-A protein	791528 IMAGE Consortium	AA437439
9278 H	4 021PAMOE	cDNA sequence AB017026	791784 IMAGE Consortium	AA414046
9279 H	6 021PAMOE	ESTs, Highly similar to hypothetical protein [H.sapiens]	792925 IMAGE Consortium	AA415581
9280 H	8 021PAMOE	activated leukocyte cell adhesion molecule	793950 IMAGE Consortium	AA422593
9281 H	10 021PAMOE	Rho-associated coiled-coil forming kinase 1	793119 IMAGE Consortium	AA437878
9282 H	12 021PAMOE	ESTs, Highly similar to LACTASE-PHLORIZIN HYDROLASE PRECURSOR [Homo sapiens]	351960 IMAGE Consortium	AI322356
9283 B	2 021WAMOF	ESTs, Highly similar to Smad6 [M.musculus]	354487 IMAGE Consortium	AI427265
9284 B	4 021WAMOF	ESTs	350145 IMAGE Consortium	AI391165
9285 B	6 021WAMOF	ESTs	351313 IMAGE Consortium	AI450530
9286 B	8 021WAMOF	ESTs	808029 IMAGE Consortium	AA432428
9287 B	10 021WAMOF			
9288 B	12 021WAMOF	ESTs	804168 IMAGE Consortium	AA474311
9289 D	2 021WAMOF	ESTs	355218 IMAGE Consortium	W47917
9290 D	4 021WAMOF	ESTs	350183 IMAGE Consortium	AI391167
9291 D	6 021WAMOF	vesicle-associated membrane protein 1	351383 IMAGE Consortium	AI450533
9292 D	8 021WAMOF			
9293 D	10 021WAMOF	ESTs	793671 IMAGE Consortium	AA415684
9294 D	12 021WAMOF	ESTs, Moderately similar to KIAA0879 protein [H.sapiens]	804191 IMAGE Consortium	AA474449
9295 F	2 021WAMOF	retinoic acid induced 1	355284 IMAGE Consortium	AI893920
9296 F	4 021WAMOF	interleukin 1 receptor antagonist homolog 1	350220 IMAGE Consortium	AI391169
9297 F	6 021WAMOF	Mus musculus mRNA for Reprimo, complete cds	352107 IMAGE Consortium	W33504
9298 F	8 021WAMOF			
9299 F	10 021WAMOF	ESTs	809168 IMAGE Consortium	AA434680
9300 F	12 021WAMOF	synaptonemal complex protein 3	804247 IMAGE Consortium	AA474336
9301 H	2 021WAMOF	ESTs	355309 IMAGE Consortium	W48326

9302 H	4 021WAMOF	ESTs, Weakly similar to MUP4_MOUSE MAJOR URINARY PROTEIN 4 PRECURSOR [M.musculus]	350266 IMAGE Consortium	AI391171
9303 H	6 021WAMOF	ESTs	350785 IMAGE Consortium	W41129
9304 H	8 021WAMOF	ESTs	807477 IMAGE Consortium	AA412978
9305 H	10 021WAMOF	SRY-box containing gene 2	793768 IMAGE Consortium	AA415862
9306 H	12 021WAMOF	Bet3 homolog (S. cerevisiae)	804268 IMAGE Consortium	AA474650
9307 B	2 0213AMOG	Mouse testis abundant mRNA sequence	803673 IMAGE Consortium	AA422834
9308 B	4 0213AMOG	ESTs, Weakly similar to similar to acyl-CoA dehydrogenases and epoxide hydrolases [C.elegans]	807019 IMAGE Consortium	AA404087
9309 B	6 0213AMOG	ESTs	832768 IMAGE Consortium	AI549830
9310 B	8 0213AMOG	ESTs, Weakly similar to Very similar and perhaps identical to Hs-CUL-4B. [H.sapiens]	835510 IMAGE Consortium	AA422439
9311 B	10 0213AMOG	retinol binding protein 7, cellular	831668 IMAGE Consortium	AA466092
9312 B	12 0213AMOG	ESTs	818409 IMAGE Consortium	AA434859
9313 D	2 0213AMOG	hypothetical protein MNCb-2146	803724 IMAGE Consortium	AA422886
9314 D	4 0213AMOG	ESTs	833188 IMAGE Consortium	AA467595
9315 D	6 0213AMOG	ESTs, Moderately similar to snRNA activating protein complex 19kDa subunit [H.sapiens]	833529 IMAGE Consortium	AA465997
9316 D	8 0213AMOG	ESTs, Highly similar to KIAA1108 protein [H.sapiens]	835805 IMAGE Consortium	AA422613
9317 D	10 0213AMOG	ESTs	831680 IMAGE Consortium	AA467022
9318 D	12 0213AMOG	ESTs, Weakly similar to CG11095 gene product [D.melanogaster]	818486 IMAGE Consortium	AA434932
9319 F	2 0213AMOG	polycystic kidney disease 2-like 2	803776 IMAGE Consortium	AA427304
9320 F	4 0213AMOG	hypothetical protein MNCb-2032	833225 IMAGE Consortium	AA432994
9321 F	6 0213AMOG	vascular endothelial growth factor	832825 IMAGE Consortium	AA427234
9322 F	8 0213AMOG	ESTs	835821 IMAGE Consortium	AA422617
9323 F	10 0213AMOG	ESTs, Highly similar to AF039942_1 HCF-binding transcription factor Zhangfei [H.sapiens]	831791 IMAGE Consortium	AA444626
9324 F	12 0213AMOG	ESTs	818605 IMAGE Consortium	AA444318
9325 H	2 0213AMOG	ESTs	803811 IMAGE Consortium	AA422971
9326 H	4 0213AMOG	osmotic stress protein 94 kDa	833248 IMAGE Consortium	AA432997
9327 H	6 0213AMOG	ESTs	832952 IMAGE Consortium	AA423051
9328 H	8 0213AMOG	growth differentiation factor 9	835857 IMAGE Consortium	AA422195
9329 H	10 0213AMOG	ESTs, Weakly similar to unnamed protein product [H.sapiens]	831853 IMAGE Consortium	AA444683
9330 H	12 0213AMOG	nectin-3 beta	818672 IMAGE Consortium	AA466550
9331 B	2 021AAMOH	mitogen activated protein kinase kinase 7	821017 IMAGE Consortium	AA647161
9332 B	4 021AAMOH	ESTs, Weakly similar to Fish protein [M.musculus]	836296 IMAGE Consortium	AA426833
9333 B	6 021AAMOH	ESTs	846306 IMAGE Consortium	AI553039
9334 B	8 021AAMOH	ESTs	845892 IMAGE Consortium	AA416143
9335 B	10 021AAMOH	ESTs	846924 IMAGE Consortium	AI553074
9336 B	12 021AAMOH	ESTs	848695 IMAGE Consortium	AA445438
9337 D	2 021AAMOH	histocompatibility 2, T region locus 17	820348 IMAGE Consortium	AA423123

9338 D	4 021AAMOH	ESTs	836321 IMAGE Consortium	AA426919
9339 D	6 021AAMOH	ESTs	847076 IMAGE Consortium	AA419833
9340 D	8 021AAMOH	ESTs	847332 IMAGE Consortium	AI552154
9341 D	10 021AAMOH	ESTs	846976 IMAGE Consortium	AA433721
9342 D	12 021AAMOH	ESTs	849400 IMAGE Consortium	AI552257
9343 F	2 021AAMOH	ESTs	821121 IMAGE Consortium	AI552344
9344 F	4 021AAMOH			
9345 F	6 021AAMOH	ATP-binding cassette, sub-family C (CFTR/MRP), member 8	846353 IMAGE Consortium	AI553043
9346 F	8 021AAMOH	ESTs	846613 IMAGE Consortium	AA422260
9347 F	10 021AAMOH	membrane-spanning 4-domains, subfamily A, member 8	849103 IMAGE Consortium	AI552228
9348 F	12 021AAMOH	ESTs	849450 IMAGE Consortium	AI552266
9349 H	2 021AAMOH	ESTs	820441 IMAGE Consortium	AA423358
9350 H	4 021AAMOH	ESTs, Weakly similar to 573K1.5 [M.musculus]	836411 IMAGE Consortium	AA426930
9351 H	6 021AAMOH	ESTs	846412 IMAGE Consortium	AI553048
9352 H	8 021AAMOH	m6a methyltransferase	845994 IMAGE Consortium	AI553023
9353 H	10 021AAMOH	ESTs	848434 IMAGE Consortium	AI552204
9354 H	12 021AAMOH	ESTs	849492 IMAGE Consortium	AI552275
9355 B	2 021HAMOI	cholecystokinin	849926 IMAGE Consortium	AA450795
9356 B	4 021HAMOI	ESTs, Weakly similar to DHAP_MOUSE ALDEHYDE DEHYDROGENASE, DIMERIC NADP-PREFERRING [M.musculus]	850493 IMAGE Consortium	AA451349
9357 B	6 021HAMOI	ESTs	860268 IMAGE Consortium	AA500903
9358 B	8 021HAMOI	ESTs	860662 IMAGE Consortium	AI613804
9359 B	10 021HAMOI	ESTs	875296 IMAGE Consortium	AA475858
9360 B	12 021HAMOI	ESTs	874820 IMAGE Consortium	AA472749
9361 D	2 021HAMOI	ESTs, Weakly similar to phospholipase C-beta-1b [M.musculus]	850046 IMAGE Consortium	AA444565
9362 D	4 021HAMOI			
9363 D	6 021HAMOI	ESTs, Moderately similar to Y391_HUMAN HYPOTHETICAL PROTEIN KIAA0391 [H.sapiens]	851559 IMAGE Consortium	AI613595
9364 D	8 021HAMOI	selectin, platelet	860885 IMAGE Consortium	AA500260
9365 D	10 021HAMOI	ESTs, Weakly similar to LEM3_MOUSE P-SELECTIN PRECURSOR [M.musculus]	876010 IMAGE Consortium	AA475278
9366 D	12 021HAMOI	ESTs	876339 IMAGE Consortium	AA476196
9367 F	2 021HAMOI	ESTs	850172 IMAGE Consortium	AA451389
9368 F	4 021HAMOI	ESTs	851279 IMAGE Consortium	AA463075
9369 F	6 021HAMOI	ESTs	851609 IMAGE Consortium	AI481758
9370 F	8 021HAMOI	ESTs	874189 IMAGE Consortium	AA472307
9371 F	10 021HAMOI	homeo box A10	875342 IMAGE Consortium	AA475215
9372 F	12 021HAMOI	DNA segment, Chr 9, ERATO Doi 402, expressed	875702 IMAGE Consortium	AA476097
9373 H	2 021HAMOI	ESTs	850217 IMAGE Consortium	AA451444
9374 H	4 021HAMOI	ESTs	851319 IMAGE Consortium	AA461645

9375 H	6 021HAMOI	Mus musculus calcium/calmodulin-dependent protein kinase II delta mRNA, partial cds	850980 IMAGE Consortium	AA462389
9376 H	8 021HAMOI	ESTs	874274 IMAGE Consortium	AA472523
9377 H	10 021HAMOI	ESTs	875355 IMAGE Consortium	AA475241
9378 H	12 021HAMOI	ESTs	875738 IMAGE Consortium	AA476131
9379 B	2 021OAMOJ	receptor (calcitonin) activity modifying protein 2	871603 IMAGE Consortium	AA462231
9380 B	4 021OAMOJ	like-glycosyltransferase	871082 IMAGE Consortium	AI552901
9381 B	6 021OAMOJ	ESTs, Weakly similar to AF220187_1 uncharacterized hypothalamus protein HT013 [H.sapiens]	891170 IMAGE Consortium	AI614502
9382 B	8 021OAMOJ	SEC22 vesicle trafficking protein-like 1 (S. cerevisiae)	890818 IMAGE Consortium	AA510654
9383 B	10 021OAMOJ	ESTs, Highly similar to CRP-ductin-alpha [M.musculus]	888508 IMAGE Consortium	AA497618
9384 B	12 021OAMOJ	ESTs, Weakly similar to B39066 proline-rich protein 15 - rat [R.norvegicus]	903578 IMAGE Consortium	AA623060
9385 D	2 021OAMOJ	ESTs, Weakly similar to T2_MOUSE OCTAPEPTIDE-REPEAT PROTEIN T2 [M.musculus]	871629 IMAGE Consortium	AA462255
9386 D	4 021OAMOJ	prostaglandin D2 synthase (21 kDa, brain)	871122 IMAGE Consortium	AA462585
9387 D	6 021OAMOJ	plasminogen activator, urokinase	891219 IMAGE Consortium	AA510298
9388 D	8 021OAMOJ	ESTs, Moderately similar to T08762 hypothetical protein DKFZp586P2219.1 - human [H.sapiens]	890831 IMAGE Consortium	AA510671
9389 D	10 021OAMOJ	bone morphogenetic protein 4	888656 IMAGE Consortium	AA498724
9390 D	12 021OAMOJ	ESTs	904375 IMAGE Consortium	AA616977
9391 F	2 021OAMOJ	ESTs	871695 IMAGE Consortium	AI552925
9392 F	4 021OAMOJ	ESTs	870828 IMAGE Consortium	AA462012
9393 F	6 021OAMOJ	ESTs	890589 IMAGE Consortium	AA509748
9394 F	8 021OAMOJ	endomucin	876330 IMAGE Consortium	AA475776
9395 F	10 021OAMOJ	syntaxin binding protein 1	888693 IMAGE Consortium	AA498760
9396 F	12 021OAMOJ	osteoglycin	902951 IMAGE Consortium	AA518187
9397 H	2 021OAMOJ	ESTs	871780 IMAGE Consortium	AI614438
9398 H	4 021OAMOJ	ESTs	870889 IMAGE Consortium	AA462133
9399 H	6 021OAMOJ	ESTs	891371 IMAGE Consortium	AA510722
9400 H	8 021OAMOJ	ESTs	890977 IMAGE Consortium	AA511234
9401 H	10 021OAMOJ	death-associated kinase 2	902193 IMAGE Consortium	AA516800
9402 H	12 021OAMOJ	solute carrier family 7 (cationic amino acid transporter, y+ system), member 5	902964 IMAGE Consortium	AA518295
9403 B	2 021VAMOK	ESTs	904896 IMAGE Consortium	AA519025
9404 B	4 021VAMOK	ESTs	904062 IMAGE Consortium	AA518560
9405 B	6 021VAMOK	ESTs, Highly similar to KIAA0336 [H.sapiens]	949441 IMAGE Consortium	AI596789
9406 B	8 021VAMOK	ESTs, Weakly similar to F-box protein FBX13 [M.musculus]		
9407 B	10 021VAMOK	ESTs, Moderately similar to Edp1 protein [M.musculus]	959586 IMAGE Consortium	AA546545
9408 B	12 021VAMOK	ESTs, Weakly similar to delta7-sterol reductase [M.musculus]	959783 IMAGE Consortium	AA546940
9409 D	2 021VAMOK	carboxypeptidase D	905307 IMAGE Consortium	AI553302

9410 D	4 021VAMOK	ESTs	904905 IMAGE Consortium	AA519037
9411 D	6 021VAMOK	sema domain, immunoglobulin domain (Ig), short basic domain, secreted, (semaphorin) 3B	902634 IMAGE Consortium	AI553114
9412 D	8 021VAMOK	ESTs	949517 IMAGE Consortium	AI596800
9413 D	10 021VAMOK	ESTs	959604 IMAGE Consortium	AA546625
9414 D	12 021VAMOK	torsin family 1, member B	959808 IMAGE Consortium	AA546959
9415 F	2 021VAMOK	ESTs, Moderately similar to SRE2_HUMAN STEROL REGULATORY ELEMENT BINDING PROTEIN-2 [H.sapiens]	904632 IMAGE Consortium	AI553249
9416 F	4 021VAMOK	ESTs	905677 IMAGE Consortium	AA529248
9417 F	6 021VAMOK	nestin	903381 IMAGE Consortium	AA518370
9418 F	8 021VAMOK	ESTs	949556 IMAGE Consortium	AI607582
9419 F	10 021VAMOK	ESTs	959632 IMAGE Consortium	AA546647
9420 F	12 021VAMOK	ESTs	959820 IMAGE Consortium	AA546964
9421 H	2 021VAMOK	DNA segment, Chr 15, Wayne State University 122, expressed	903982 IMAGE Consortium	AA522079
9422 H	4 021VAMOK	ESTs	904983 IMAGE Consortium	AI553279
9423 H	6 021VAMOK	ESTs	901938 IMAGE Consortium	AA517795
9424 H	8 021VAMOK	ESTs	948907 IMAGE Consortium	AA544962
9425 H	10 021VAMOK	ESTs	959673 IMAGE Consortium	AA546753
9426 H	12 021VAMOK	ESTs	959862 IMAGE Consortium	AA547034
9427 B	2 0212AMOL	pentaxin related gene	355881 IMAGE Consortium	AI893922
9428 B	4 0212AMOL	sarcoglycan, beta (43kD dystrophin-associated glycoprotein)	368542 IMAGE Consortium	W61688
9429 B	6 0212AMOL	ESTs	921646 IMAGE Consortium	AA529879
9430 B	8 0212AMOL	ESTs	944304 IMAGE Consortium	AA545496
9431 B	10 0212AMOL	ESTs	944739 IMAGE Consortium	AA545305
9432 B	12 0212AMOL	baculoviral IAP repeat-containing 6	959926 IMAGE Consortium	AA547134
9433 D	2 0212AMOL	Scgn10 like-protein	368000 IMAGE Consortium	W53771
9434 D	4 0212AMOL	even skipped homeotic gene 2 homolog	367878 IMAGE Consortium	W53902
9435 D	6 0212AMOL	ESTs	921690 IMAGE Consortium	AA529898
9436 D	8 0212AMOL	ESTs	944331 IMAGE Consortium	AA545529
9437 D	10 0212AMOL	serine/threonine kinase 13 (aurora/IPL-like)	944766 IMAGE Consortium	AA545317
9438 D	12 0212AMOL	U1 small nuclear ribonucleoprotein 1C	959949 IMAGE Consortium	AA547160
9439 F	2 0212AMOL	respiratory neuron homeobox	367322 IMAGE Consortium	W53935
9440 F	4 0212AMOL	calcium channel beta 3 subunit	367924 IMAGE Consortium	W54062
9441 F	6 0212AMOL	ESTs	921775 IMAGE Consortium	AA530026
9442 F	8 0212AMOL	ESTs	935627 IMAGE Consortium	AA543690
9443 F	10 0212AMOL	ESTs	944788 IMAGE Consortium	AA545395
9444 F	12 0212AMOL	ESTs, Weakly similar to Keap1 [M.musculus]	960022 IMAGE Consortium	AA547245
9445 H	2 0212AMOL	ESTs	368180 IMAGE Consortium	AI427414
9446 H	4 0212AMOL	ESTs, Highly similar to HOMEBOX PROTEIN HOX-B2 [Homo sapiens]	367979 IMAGE Consortium	W53769

9447 H	6 0212AMOL	ESTs	921872 IMAGE Consortium	AA512099
9448 H	8 0212AMOL	interleukin 1 receptor, type I	935643 IMAGE Consortium	AA543982
9449 H	10 0212AMOL	ESTs	944798 IMAGE Consortium	AA545399
9450 H	12 0212AMOL	ESTs	960057 IMAGE Consortium	AA547487
9451 B	2 0219AMOM	ESTs	960276 IMAGE Consortium	AA549705
9452 B	4 0219AMOM	ESTs	960466 IMAGE Consortium	AA549305
9453 B	6 0219AMOM	histone acetyltransferase	960722 IMAGE Consortium	AA549040
9454 B	8 0219AMOM	ESTs	961887 IMAGE Consortium	AA672319
9455 B	10 0219AMOM	ESTs	961363 IMAGE Consortium	AA547664
9456 B	12 0219AMOM	ESTs	1003019 IMAGE Consortium	AA684464
9457 D	2 0219AMOM	ESTs, Moderately similar to HYPOTHETICAL PROTEIN KIAA0310 [H.sapiens]	960289 IMAGE Consortium	AA549732
9458 D	4 0219AMOM	phosphoinositide kinase, fyve-containing	961229 IMAGE Consortium	AA547524
9459 D	6 0219AMOM	DNA segment, Chr 2, ERATO Doi 694, expressed	960748 IMAGE Consortium	AA549134
9460 D	8 0219AMOM	superoxide dismutase 1, soluble	963308 IMAGE Consortium	AA671734
9461 D	10 0219AMOM	ESTs	961390 IMAGE Consortium	AA547674
9462 D	12 0219AMOM	ESTs, Weakly similar to Nbr1 [M.musculus]	1003044 IMAGE Consortium	AA684474
9463 F	2 0219AMOM	ESTs, Moderately similar to putative [C.elegans]	961023 IMAGE Consortium	AA549662
9464 F	4 0219AMOM	ESTs	961272 IMAGE Consortium	AA547551
9465 F	6 0219AMOM	ESTs, Highly similar to hypothetical protein [H.sapiens]	960757 IMAGE Consortium	AA549146
9466 F	8 0219AMOM	ESTs, Highly similar to CD2 cytoplasmic domain binding protein [H.sapiens]	961900 IMAGE Consortium	AA672170
9467 F	10 0219AMOM	ESTs	961409 IMAGE Consortium	AA553306
9468 F	12 0219AMOM	differentially expressed in B16F10 1	1003047 IMAGE Consortium	AA684483
9469 H	2 0219AMOM	ESTs, Highly similar to unnamed protein product [H.sapiens]	961030 IMAGE Consortium	AA549656
9470 H	4 0219AMOM	ESTs	960585 IMAGE Consortium	AA549524
9471 H	6 0219AMOM	ESTs, Highly similar to AF223467_1 NPD008 protein [H.sapiens]	960834 IMAGE Consortium	AA549260
9472 H	8 0219AMOM	ESTs	963385 IMAGE Consortium	AA670581
9473 H	10 0219AMOM	ESTs	961429 IMAGE Consortium	AA549181
9474 H	12 0219AMOM	ESTs	1003050 IMAGE Consortium	AA684477
9475 B	2 021GAMON	nibrin	1003235 IMAGE Consortium	AA684114
9476 B	4 021GAMON	fused toes	1003440 IMAGE Consortium	AA684254
9477 B	6 021GAMON	protein kinase, cAMP dependent, catalytic, beta	1003630 IMAGE Consortium	AA683704
9478 B	8 021GAMON	transcription factor AP-2, gamma	976458 IMAGE Consortium	AA619859
9479 B	10 021GAMON	ESTs	976585 IMAGE Consortium	AA619725
9480 B	12 021GAMON	ornithine decarboxylase antizyme 2	976684 IMAGE Consortium	AA619805
9481 D	2 021GAMON	Mus musculus mRNA for AWP1 protein (Awp1 gene)	1003332 IMAGE Consortium	AA666810
9482 D	4 021GAMON	protein tyrosine phosphatase 4a1	1003466 IMAGE Consortium	AA684319
9483 D	6 021GAMON	ESTs, Moderately similar to DTD_MOUSE SULFATE TRANSPORTER [M.musculus]	1002926 IMAGE Consortium	AA792032
9484 D	8 021GAMON	ESTs, Weakly similar to HYPOTHETICAL 38.5 KD	976466 IMAGE Consortium	AA619874

PROTEIN IN SUI2-TDH2 INTERGENIC REGION [Saccharomyces cerevisiae]				
9485 D	10 021GAMON	eukaryotic translation initiation factor 3, subunit 4 (delta, 44 kDa)	976592 IMAGE Consortium	AA619722
9486 D	12 021GAMON	coatamer protein complex, subunit beta 2 (beta prime)	976688 IMAGE Consortium	AA619806
9487 F	2 021GAMON	ESTs	1003369 IMAGE Consortium	AA684166
9488 F	4 021GAMON	protein phosphatase 1A, magnesium dependent, alpha isoform	1003470 IMAGE Consortium	AA684321
9489 F	6 021GAMON	ESTs, Highly similar to A57284 spermatid perinuclear RNA-binding protein Spnr - mouse [M.musculus]	1002999 IMAGE Consortium	AA684448
9490 F	8 021GAMON	sortilin-related receptor, LDLR class A repeats-containing	976482 IMAGE Consortium	AA619881
9491 F	10 021GAMON	T-cell lymphoma breakpoint 1	976610 IMAGE Consortium	AA619733
9492 F	12 021GAMON	Mus musculus mRNA for beta-catenin-interacting protein ICAT, complete cds	976694 IMAGE Consortium	AA619808
9493 H	2 021GAMON	ESTs	1003377 IMAGE Consortium	AA684170
9494 H	4 021GAMON	ESTs, Moderately similar to This gene is novel. [H.sapiens]	1003481 IMAGE Consortium	AA684334
9495 H	6 021GAMON	ESTs	976397 IMAGE Consortium	AA619782
9496 H	8 021GAMON	cyclin I	976489 IMAGE Consortium	AA619950
9497 H	10 021GAMON	ring finger protein 2	976633 IMAGE Consortium	AA619751
9498 H	12 021GAMON	ESTs, Weakly similar to RB3A_MOUSE RAS-RELATED PROTEIN RAB-3A [M.musculus]	976704 IMAGE Consortium	AA619816
9499 B	2 021NAMOO	diaphanous homolog 1 (Drosophila)	976736 IMAGE Consortium	AA619832
9500 B	4 021NAMOO	ESTs	1026548 IMAGE Consortium	AA672908
9501 B	6 021NAMOO	ESTs, Weakly similar to CG14414 gene product [D.melanogaster]	1003916 IMAGE Consortium	AA674573
9502 B	8 021NAMOO	transcription factor 17	1026103 IMAGE Consortium	AA616580
9503 B	10 021NAMOO	cholinergic receptor, nicotinic, alpha polypeptide 1 (muscle)	1050557 IMAGE Consortium	AA596914
9504 B	12 021NAMOO	ESTs	1052385 IMAGE Consortium	AI593925
9505 D	2 021NAMOO	ESTs, Highly similar to AF091457_1 zinc finger protein RIN ZF [R.norvegicus]	976740 IMAGE Consortium	AA619834
9506 D	4 021NAMOO	Rac GTPase-activating protein 1	1004004 IMAGE Consortium	AA674774
9507 D	6 021NAMOO	ESTs	1003942 IMAGE Consortium	AA674592
9508 D	8 021NAMOO	cyclic AMP phosphoprotein, 19kD	1026153 IMAGE Consortium	AI642230
9509 D	10 021NAMOO	ESTs	1052005 IMAGE Consortium	AA592637
9510 D	12 021NAMOO	ESTs	1050974 IMAGE Consortium	AA592797
9511 F	2 021NAMOO	Mus musculus nuclear ATP/GTP-binding protein (Nna1) mRNA, complete cds	976758 IMAGE Consortium	AA619843
9512 F	4 021NAMOO	guanylate cyclase activator 1a (retina)	1004019 IMAGE Consortium	AA674797
9513 F	6 021NAMOO			
9514 F	8 021NAMOO	ESTs	1025464 IMAGE Consortium	AA590694
9515 F	10 021NAMOO	ESTs	1052047 IMAGE Consortium	AI594442
9516 F	12 021NAMOO	ESTs	1053118 IMAGE Consortium	AA606747



9517 H	2 021NAMOO	H3 histone, family 3B	976761 IMAGE Consortium	AA619848
9518 H	4 021NAMOO	receptor-like tyrosine kinase	1003828 IMAGE Consortium	AA675371
9519 H	6 021NAMOO	cyclin-dependent kinase inhibitor 2D (p19, inhibits CDK4)	1003993 IMAGE Consortium	AA674780
9520 H	8 021NAMOO	ESTs	1024791 IMAGE Consortium	AA793372
9521 H	10 021NAMOO	ESTs	1051396 IMAGE Consortium	AI429031
9522 H	12 021NAMOO	ESTs	1052414 IMAGE Consortium	AA839333
9523 B	2 021UAMOP	ESTs	1051949 IMAGE Consortium	AA597263
9524 B	4 021UAMOP	ESTs, Moderately similar to dJ967N21.4 [H.sapiens]	1053755 IMAGE Consortium	AA638578
9525 B	6 021UAMOP	sialyltransferase 4A (beta-galactosidase alpha-2,3-sialyltransferase)	1068701 IMAGE Consortium	AA796822
9526 B	8 021UAMOP	ESTs	1078592 IMAGE Consortium	AA727914
9527 B	10 021UAMOP	ESTs	1066958 IMAGE Consortium	AA616122
9528 B	12 021UAMOP	ESTs	1064618 IMAGE Consortium	AA615712
9529 D	2 021UAMOP	ESTs	1052737 IMAGE Consortium	AA607553
9530 D	4 021UAMOP	ESTs	1053853 IMAGE Consortium	AI594104
9531 D	6 021UAMOP	synaptogamin V	1068724 IMAGE Consortium	AA796832
9532 D	8 021UAMOP	kelch-like (D. melanogaster)	1077898 IMAGE Consortium	AA796762
9533 D	10 021UAMOP	ESTs, Moderately similar to unnamed protein product [H.sapiens]	1066986 IMAGE Consortium	AA616180
9534 D	12 021UAMOP	ESTs, Highly similar to YSPL-1 form 1 □ [M.musculus]	1065366 IMAGE Consortium	AA615692
9535 F	2 021UAMOP	ESTs	1052753 IMAGE Consortium	AI593952
9536 F	4 021UAMOP	ESTs	1053944 IMAGE Consortium	AI594111
9537 F	6 021UAMOP	ESTs	1068763 IMAGE Consortium	AA796658
9538 F	8 021UAMOP	kinesin family member 3c	1078706 IMAGE Consortium	AA823329
9539 F	10 021UAMOP	ESTs	1064337 IMAGE Consortium	AI603990
9540 F	12 021UAMOP	ESTs	1065389 IMAGE Consortium	AA615785
9541 H	2 021UAMOP	ESTs	1052758 IMAGE Consortium	AI593950
9542 H	4 021UAMOP	ESTs	1067728 IMAGE Consortium	AA764531
9543 H	6 021UAMOP	ESTs	1078215 IMAGE Consortium	AI593576
9544 H	8 021UAMOP	ESTs	1078904 IMAGE Consortium	AI593604
9545 H	10 021UAMOP	ESTs	1064364 IMAGE Consortium	AA615373
9546 H	12 021UAMOP	ESTs	1066103 IMAGE Consortium	AA674000
9547 B	2 0211AMOQ	mesothelin	1065669 IMAGE Consortium	AA673869
9548 B	4 0211AMOQ	ESTs	1065922 IMAGE Consortium	AI428996
9549 B	6 0211AMOQ	ESTs	1093696 IMAGE Consortium	AA869499
9550 B	8 0211AMOQ	ESTs	1096012 IMAGE Consortium	AA871251
9551 B	10 0211AMOQ	defensin related cryptdin, related sequence 2	1096595 IMAGE Consortium	AA871641
9552 B	12 0211AMOQ	ESTs, Weakly similar to LIS1_MOUSE PLATELET-ACTIVATING FACTOR ACETYLHYDROLASE IB ALPHA SUBUNIT [M.musculus]	400361 IMAGE Consortium	AI425696
9553 D	2 0211AMOQ	ESTs	1066442 IMAGE Consortium	AI594391
9554 D	4 0211AMOQ	DNA segment, Chr 2, ERATO Doi 210, expressed	1065951 IMAGE Consortium	AA611788
9555 D	6 0211AMOQ	ESTs	1095148 IMAGE Consortium	AI591658

9556 D	8 0211AMQ	bone morphogenetic protein 8a	1096039 IMAGE Consortium	AA871280
9557 D	10 0211AMQ	defensin related cryptdin, related sequence 7	1096615 IMAGE Consortium	AA871663
9558 D	12 0211AMQ	paired-like homeodomain transcription factor 2	400449 IMAGE Consortium	AI893277
9559 F	2 0211AMQ	ESTs	1066450 IMAGE Consortium	AA615889
9560 F	4 0211AMQ	ephrin A2	1066697 IMAGE Consortium	AA611809
9561 F	6 0211AMQ	ESTs	1079127 IMAGE Consortium	AA823451
9562 F	8 0211AMQ	ESTs	1096050 IMAGE Consortium	AA871274
9563 F	10 0211AMQ	nuclear factor, erythroid derived 2, like 3	1096647 IMAGE Consortium	AA871687
9564 F	12 0211AMQ	neuropeptide nociceptin 1	400481 IMAGE Consortium	W77365
9565 H	2 0211AMQ	ESTs	1066483 IMAGE Consortium	AA615928
9566 H	4 0211AMQ	ESTs	1066704 IMAGE Consortium	AA611812
9567 H	6 0211AMQ	ESTs	1079133 IMAGE Consortium	AA823452
9568 H	8 0211AMQ	ESTs	1096104 IMAGE Consortium	AA871326
9569 H	10 0211AMQ	ESTs, Weakly similar to Hrs [M.musculus]	1096744 IMAGE Consortium	AA871749
9570 H	12 0211AMQ	ESTs	400531 IMAGE Consortium	W76731
9571 B	2 0218AMOR	ESTs, Moderately similar to unnamed protein product [H.sapiens]	400713 IMAGE Consortium	W75868
9572 B	4 0218AMOR	ESTs, Weakly similar to CG17883 gene product [D.melanogaster]	389909 IMAGE Consortium	W70701
9573 B	6 0218AMOR	ESTs, Weakly similar to AF137367_1 VPS10 domain receptor protein SORCS [M.musculus]	400093 IMAGE Consortium	W76988
9574 B	8 0218AMOR	ESTs, Highly similar to RasGAP-related protein [H.sapiens]	1107602 IMAGE Consortium	AA623836
9575 B	10 0218AMOR	ESTs	1107733 IMAGE Consortium	AA623991
9576 B	12 0218AMOR	ESTs, Highly similar to dJ347H13.5 [H.sapiens]	1125013 IMAGE Consortium	AA692962
9577 D	2 0218AMOR	ESTs	400767 IMAGE Consortium	AI425727
9578 D	4 0218AMOR	glypican 4	402621 IMAGE Consortium	AI893963
9579 D	6 0218AMOR	Ras-GTPase-activating protein SH3-domain binding protein	1108013 IMAGE Consortium	AA623896
9580 D	8 0218AMOR	ESTs, Moderately similar to HAT1_HUMAN HISTONE ACETYLTRANSFERASE TYPE B CATALYTIC SUBUNIT [H.sapiens]	1107685 IMAGE Consortium	AA623936
9581 D	10 0218AMOR	adaptor protein complex AP-2, mu1	1107755 IMAGE Consortium	AA624010
9582 D	12 0218AMOR	alkaline phosphatase 5	1125142 IMAGE Consortium	AA693081
9583 F	2 0218AMOR	ESTs	389563 IMAGE Consortium	W71566
9584 F	4 0218AMOR	oligodendrocyte transcription factor 2	390022 IMAGE Consortium	AI430166
9585 F	6 0218AMOR	hypothetical protein	1107314 IMAGE Consortium	AA645457
9586 F	8 0218AMOR	pleckstrin homology, Sec7 and coiled/coil domains 3	1107751 IMAGE Consortium	AA624008
9587 F	10 0218AMOR	ESTs	1107849 IMAGE Consortium	AA623676
9588 F	12 0218AMOR	ESTs	1125282 IMAGE Consortium	AA693302
9589 H	2 0218AMOR	transmembrane tryptase	402278 IMAGE Consortium	AI425768
9590 H	4 0218AMOR	FK506 binding protein 1b (12.6 kDa)	390134 IMAGE Consortium	AI385895
9591 H	6 0218AMOR	golgi autoantigen, golgin subfamily a, 4	1107386 IMAGE Consortium	AA647230

9592 H	8 0218AMOR	protein tyrosine kinase 9 related protein	1107846 IMAGE Consortium	AA623663
9593 H	10 0218AMOR	ESTs	1107874 IMAGE Consortium	AA623706
9594 H	12 0218AMOR	ESTs, Highly similar to AF158186_1 pEachy [R.norvegicus]	1126031 IMAGE Consortium	AA691050
9595 B	2 021FAMOS	ESTs, Weakly similar to NTC3_MOUSE NEUROGENIC LOCUS NOTCH 3 PROTEIN [M.musculus]	1125569 IMAGE Consortium	AA675294
9596 B	4 021FAMOS	DNA segment, Chr 8, ERATO Doi 580, expressed	1134492 IMAGE Consortium	AA681114
9597 B	6 021FAMOS	ESTs, Moderately similar to ribonuclease P protein subunit p14 [H.sapiens]	1135437 IMAGE Consortium	AA823904
9598 B	8 021FAMOS	ESTs, Weakly similar to cDNA EST EMBL:D70762 comes from this gene [C.elegans]	1152659 IMAGE Consortium	AA645304
9599 B	10 021FAMOS	ESTs	1136838 IMAGE Consortium	AA688811
9600 B	12 021FAMOS	ESTs	1139347 IMAGE Consortium	AA656612
9601 D	2 021FAMOS	ESTs	1125584 IMAGE Consortium	AA824068
9602 D	4 021FAMOS	ESTs, Weakly similar to TFSL_MOUSE TRANSCRIPTION FACTOR S-II-RELATED PROTEIN 3 [M.musculus]	1125748 IMAGE Consortium	AA691124
9603 D	6 021FAMOS	heparan sulfate 2-O-sulfotransferase 1	1153047 IMAGE Consortium	AI593151
9604 D	8 021FAMOS	ESTs	1135957 IMAGE Consortium	AA734656
9605 D	10 021FAMOS	serine protease inhibitor, Kazal type 4	1138342 IMAGE Consortium	AA821396
9606 D	12 021FAMOS	ESTs	1137940 IMAGE Consortium	AA691435
9607 F	2 021FAMOS	ESTs, Moderately similar to unknown [M.musculus]	1134304 IMAGE Consortium	AA675537
9608 F	4 021FAMOS	mitogen activated protein kinase kinase 1	1134560 IMAGE Consortium	AA681161
9609 F	6 021FAMOS	ESTs	1137054 IMAGE Consortium	AA689070
9610 F	8 021FAMOS	DNA segment, Chr 16, Wayne State University 119, expressed	1135983 IMAGE Consortium	AA734673
9611 F	10 021FAMOS	ESTs	1138375 IMAGE Consortium	AA691557
9612 F	12 021FAMOS	ESTs	1137996 IMAGE Consortium	AA691514
9613 H	2 021FAMOS	dynein, axon, light chain 4	1134310 IMAGE Consortium	AA675540
9614 H	4 021FAMOS	RNA polymerase 1-2 (128 kDa subunit)	1134569 IMAGE Consortium	AA681171
9615 H	6 021FAMOS	ESTs	1136409 IMAGE Consortium	AA683926
9616 H	8 021FAMOS	calcium/calmodulin-dependent protein kinase II, beta	1138142 IMAGE Consortium	AA691743
9617 H	10 021FAMOS	ESTs	1137748 IMAGE Consortium	AA689631
9618 H	12 021FAMOS	ESTs	1138711 IMAGE Consortium	AA734168
9619 B	2 021MAMOT	ESTs	1164708 IMAGE Consortium	AA690434
9620 B	4 021MAMOT	ESTs, Highly similar to putative G protein-coupled receptor, P40GPRT [M.musculus]	1154112 IMAGE Consortium	AA796946
9621 B	6 021MAMOT	runt related transcription factor 2	1162652 IMAGE Consortium	AA771675
9622 B	8 021MAMOT	ESTs, Highly similar to T00093 hypothetical protein KIAA0467 - human [H.sapiens]	1162985 IMAGE Consortium	AI592483
9623 B	10 021MAMOT	ESTs	1195078 IMAGE Consortium	AA690500
9624 B	12 021MAMOT	sterol-C5-desaturase (fungal ERG3, delta-5-desaturase) homolog (S. cerevisae)	1195500 IMAGE Consortium	AA690751

9625 D	2	021MAMOT	ESTs	1166200	IMAGE Consortium	AA710588
9626 D	4	021MAMOT	cholinergic receptor, nicotinic, delta polypeptide	1162269	IMAGE Consortium	AA691939
9627 D	6	021MAMOT	alcohol dehydrogenase family 3, subfamily A1	1162684	IMAGE Consortium	AA770709
9628 D	8	021MAMOT	RAB17, member RAS oncogene family	1165130	IMAGE Consortium	AA709984
9629 D	10	021MAMOT	ESTs, Weakly similar to eIF4E-like protein 4E-LP [M.musculus]	1195132	IMAGE Consortium	AA690645
9630 D	12	021MAMOT	ESTs	1195582	IMAGE Consortium	AA711684
9631 F	2	021MAMOT	ESTs	1167049	IMAGE Consortium	AA711115
9632 F	4	021MAMOT	syntaxin 4A (placental)	1162298	IMAGE Consortium	AA691960
9633 F	6	021MAMOT	ESTs	1164110	IMAGE Consortium	AA690711
9634 F	8	021MAMOT	ESTs, Weakly similar to unnamed protein product [H.sapiens]	1164502	IMAGE Consortium	AA929194
9635 F	10	021MAMOT	ESTs	1195244	IMAGE Consortium	AA718108
9636 F	12	021MAMOT	ESTs	1195607	IMAGE Consortium	AA711713
9637 H	2	021MAMOT	ESTs	1166415	IMAGE Consortium	AA710200
9638 H	4	021MAMOT	angiopoietin	1163088	IMAGE Consortium	AA691900
9639 H	6	021MAMOT	lymphoid blast crisis-like 1	1162716	IMAGE Consortium	AA770840
9640 H	8	021MAMOT	ESTs	1163837	IMAGE Consortium	AA692621
9641 H	10	021MAMOT	ESTs	1195251	IMAGE Consortium	AA718166
9642 H	12	021MAMOT	ESTs	1195654	IMAGE Consortium	AA711779
9643 B	2	021TAMOU	hepatic nuclear factor 4	386299	IMAGE Consortium	AI893936
9644 B	4	021TAMOU	ESTs	373295	IMAGE Consortium	W57281
9645 B	6	021TAMOU	ESTs	374665	IMAGE Consortium	AI430462
9646 B	8	021TAMOU	ESTs	387162	IMAGE Consortium	AI430622
9647 B	10	021TAMOU	transmembrane protein 6	1178215	IMAGE Consortium	AA718842
9648 B	12	021TAMOU	ESTs	1225038	IMAGE Consortium	AA756334
9649 D	2	021TAMOU	granin-like neuroendocrine peptide precursor	387474	IMAGE Consortium	W66762
9650 D	4	021TAMOU	ESTs, Highly similar to RPB17 [R.norvegicus]	374167	IMAGE Consortium	W59105
9651 D	6	021TAMOU	ESTs	373943	IMAGE Consortium	AI430090
9652 D	8	021TAMOU	ESTs	387191	IMAGE Consortium	W65543
9653 D	10	021TAMOU	ESTs	1176879	IMAGE Consortium	AA733381
9654 D	12	021TAMOU	cathepsin F	1225054	IMAGE Consortium	AA756341
9655 F	2	021TAMOU	v-maf musculoaponeurotic fibrosarcoma oncogene family, protein F (avian)	387565	IMAGE Consortium	W70403
9656 F	4	021TAMOU	ESTs, Moderately similar to PB39 [H.sapiens]	374260	IMAGE Consortium	W59204
9657 F	6	021TAMOU	Public domain EST	374889	IMAGE Consortium	
9658 F	8	021TAMOU	ESTs, Highly similar to CYTIDINE DEAMINASE [Homo sapiens]	386489	IMAGE Consortium	AI430581
9659 F	10	021TAMOU	bone morphogenetic protein 6	1177726	IMAGE Consortium	AA718365
9660 F	12	021TAMOU	ESTs	1225084	IMAGE Consortium	AA756379
9661 H	2	021TAMOU	ESTs	387744	IMAGE Consortium	AI430641
9662 H	4	021TAMOU	ribosomal protein L7a	374336	IMAGE Consortium	AI430447
9663 H	6	021TAMOU	ESTs	374997	IMAGE Consortium	W62781

9664 H	8 021TAMOU	ESTs	388017 IMAGE Consortium	W70441
9665 H	10 021TAMOU	ESTs, Highly similar to AF155196_1 mindin precursor [R.norvegicus]	1177751 IMAGE Consortium	AA718392
9666 H	12 021TAMOU	ESTs, Highly similar to RRB_MOUSE RETINOIC ACID RECEPTOR BETA [M.musculus]	1225132 IMAGE Consortium	AA762870
9667 B	2 0210AMOV			
9668 B	4 0210AMOV			
9669 B	6 0210AMOV	ESTs, Moderately similar to molybdopterin synthase sulfurylase [H.sapiens]	1225852 IMAGE Consortium	AA739189
9670 B	8 0210AMOV	growth factor independent 1	1195959 IMAGE Consortium	AA716844
9671 B	10 0210AMOV	coagulation factor III	1230698 IMAGE Consortium	AA879919
9672 B	12 0210AMOV	ZW10 homolog (Drosophila), centromere/kinetochore protein	1226392 IMAGE Consortium	AA940442
9673 D	2 0210AMOV	ESTs, Moderately similar to BRDT [H.sapiens]	1226138 IMAGE Consortium	AA739436
9674 D	4 0210AMOV	phospholipid scramblase 1	1225724 IMAGE Consortium	AA764163
9675 D	6 0210AMOV	crystallin, alpha binding protein 1	1225883 IMAGE Consortium	AA739343
9676 D	8 0210AMOV	ESTs	1230263 IMAGE Consortium	AA879516
9677 D	10 0210AMOV	ESTs, Moderately similar to TDT_MOUSE DNA NUCLEOTIDYLEXOTRANSFERASE [M.musculus]	1230732 IMAGE Consortium	AA879946
9678 D	12 0210AMOV	ESTs, Weakly similar to cDNA EST CEMSA26F comes from this gene [C.elegans]	1226487 IMAGE Consortium	AI466137
9679 F	2 0210AMOV	ESTs	1226169 IMAGE Consortium	AI430246
9680 F	4 0210AMOV	cardiotrophin 1	1225744 IMAGE Consortium	AA764184
9681 F	6 0210AMOV	MYLE protein	1195838 IMAGE Consortium	AA712093
9682 F	8 0210AMOV	ESTs	1230309 IMAGE Consortium	AA880168
9683 F	10 0210AMOV	metallocarboxypeptidase CPX-1	1230769 IMAGE Consortium	AA880365
9684 F	12 0210AMOV	ESTs	1226533 IMAGE Consortium	AA756602
9685 H	2 0210AMOV	ESTs	1225511 IMAGE Consortium	AA762168
9686 H	4 0210AMOV	ESTs	1225772 IMAGE Consortium	AA764252
9687 H	6 0210AMOV	tumor necrosis factor receptor superfamily, member 7	1195887 IMAGE Consortium	AA717022
9688 H	8 0210AMOV	ESTs	1230487 IMAGE Consortium	AA880254
9689 H	10 0210AMOV	ESTs, Weakly similar to AC004858_3 U1 small ribonucleoprotein 1SNRP homolog [H.sapiens]	1230800 IMAGE Consortium	AA879989
9690 H	12 0210AMOV	Mus musculus clone IMAGE:349147 unknown protein mRNA, partial cds	1226559 IMAGE Consortium	AA756707
9691 B	2 0217AMOW	ESTs	1226752 IMAGE Consortium	AA738627
9692 B	4 0217AMOW	ESTs	1263009 IMAGE Consortium	AA856261
9693 B	6 0217AMOW	ESTs	1248131 IMAGE Consortium	AA762383
9694 B	8 0217AMOW	transcription factor 20	1262899 IMAGE Consortium	AA839615
9695 B	10 0217AMOW	pre-B lymphocyte gene 3	1246296 IMAGE Consortium	AA823083
9696 B	12 0217AMOW	ESTs, Moderately similar to hypothetical protein 384D8_7 [H.sapiens]	1246510 IMAGE Consortium	AI465805
9697 D	2 0217AMOW	plasminogen activator inhibitor, type II	1226790 IMAGE Consortium	AA738996

9698 D	4 0217AMOW	CD7 antigen	1263105 IMAGE Consortium	AA856336
9699 D	6 0217AMOW	small inducible cytokine subfamily A17	1263461 IMAGE Consortium	AA863527
9700 D	8 0217AMOW	ESTs	1247634 IMAGE Consortium	AA763437
9701 D	10 0217AMOW	CD8 antigen, alpha chain	1247019 IMAGE Consortium	AA839138
9702 D	12 0217AMOW	ESTs	1246535 IMAGE Consortium	AI465811
9703 F	2 0217AMOW	src associated in mitosis, 68 kDa	1226871 IMAGE Consortium	AA738714
9704 F	4 0217AMOW	ESTs	1263158 IMAGE Consortium	AA856374
9705 F	6 0217AMOW	ESTs	1248193 IMAGE Consortium	AA763026
9706 F	8 0217AMOW	ESTs	1248386 IMAGE Consortium	AA789647
9707 F	10 0217AMOW	ESTs	1247093 IMAGE Consortium	AI465828
9708 F	12 0217AMOW	ESTs	1245824 IMAGE Consortium	AA822817
9709 H	2 0217AMOW	ESTs, Weakly similar to nuclear localization signal binding protein [M.musculus]	1243612 IMAGE Consortium	AA789810
9710 H	4 0217AMOW	phosphodiesterase 1C	1263202 IMAGE Consortium	AA856411
9711 H	6 0217AMOW	ESTs	1248231 IMAGE Consortium	AA762267
9712 H	8 0217AMOW	ESTs	1248592 IMAGE Consortium	AA759412
9713 H	10 0217AMOW	phosphorylase kinase gamma	1245679 IMAGE Consortium	AA798637
9714 H	12 0217AMOW	ESTs	1246587 IMAGE Consortium	AA832598
9715 B	2 021EAMOX	mitogen-activated protein kinase 12	1247477 IMAGE Consortium	AA855742
9716 B	4 021EAMOX	ESTs	1244663 IMAGE Consortium	AA797268
9717 B	6 021EAMOX	ESTs	1264282 IMAGE Consortium	AA881315
9718 B	8 021EAMOX	ESTs	1265152 IMAGE Consortium	AA880133
9719 B	10 021EAMOX	src-like adapter protein	1265347 IMAGE Consortium	AA867150
9720 B	12 021EAMOX	Public domain EST	1265532 IMAGE Consortium	
9721 D	2 021EAMOX	H5AR gene	1246969 IMAGE Consortium	AA839098
9722 D	4 021EAMOX	ESTs	1245461 IMAGE Consortium	AA797868
9723 D	6 021EAMOX	ESTs, Moderately similar to unnamed protein product [H.sapiens]	1264298 IMAGE Consortium	AI509580
9724 D	8 021EAMOX	ESTs	1265186 IMAGE Consortium	AA867873
9725 D	10 021EAMOX	ESTs, Weakly similar to ANKYRIN, BRAIN VARIANT 2 [Homo sapiens]	1265364 IMAGE Consortium	AA867157
9726 D	12 021EAMOX	ESTs	1264851 IMAGE Consortium	AI509658
9727 F	2 021EAMOX	ESTs	1244275 IMAGE Consortium	AA790810
9728 F	4 021EAMOX	apoptosis-associated tyrosine kinase	1245479 IMAGE Consortium	AA797877
9729 F	6 021EAMOX	ESTs	1264303 IMAGE Consortium	AI509586
9730 F	8 021EAMOX	ESTs	1264482 IMAGE Consortium	AI509615
9731 F	10 021EAMOX	ESTs	1265371 IMAGE Consortium	AI509758
9732 F	12 021EAMOX	ESTs	1264889 IMAGE Consortium	AI509667
9733 H	2 021EAMOX	ESTs	1245190 IMAGE Consortium	AI465992
9734 H	4 021EAMOX	adaptor-related protein complex AP-3, beta 2 subunit	1263332 IMAGE Consortium	AI509574
9735 H	6 021EAMOX	solute carrier family 25 (mitochondrial carrier, brain) member 14	1265034 IMAGE Consortium	AA881591
9736 H	8 021EAMOX	ESTs	1264487 IMAGE Consortium	AA881377

9737 H	10 021EAMOX	ESTs	1264659 IMAGE Consortium	AA881388
9738 H	12 021EAMOX	ESTs	1264930 IMAGE Consortium	AA881492
9739 B	2 021LAMOY	ESTs, Highly similar to 5-LIPOXYGENASE ACTIVATING PROTEIN [Mus musculus]	1329349 IMAGE Consortium	AA930477
9740 B	4 021LAMOY	lymphocyte antigen 55 complex, locus B	1332385 IMAGE Consortium	AI877001
9741 B	6 021LAMOY	E2F transcription factor 3	1378891 IMAGE Consortium	AI020268
9742 B	8 021LAMOY	ESTs, Weakly similar to CG14648 gene product [D.melanogaster]	1366033 IMAGE Consortium	AI020795
9743 B	10 021LAMOY	UDP-Gal:betaGlcNAc beta 1,3-galactosyltransferase, polypeptide 2	1365902 IMAGE Consortium	AI021204
9744 B	12 021LAMOY	ESTs	1363041 IMAGE Consortium	AI006445
9745 D	2 021LAMOY	ESTs	1281324 IMAGE Consortium	AA874612
9746 D	4 021LAMOY	ESTs, Moderately similar to AF133123_1 transcription factor IIIC102 [H.sapiens]	1379052 IMAGE Consortium	AI050248
9747 D	6 021LAMOY	myosin Vb	1378935 IMAGE Consortium	AI020338
9748 D	8 021LAMOY	RAS, dexamethasone-induced 1	1366165 IMAGE Consortium	AI020897
9749 D	10 021LAMOY	N-myc downstream regulated 3	1345003 IMAGE Consortium	AA939860
9750 D	12 021LAMOY	transcription factor 7, T-cell specific	1344933 IMAGE Consortium	AA958693
9751 F	2 021LAMOY	ESTs, Weakly similar to AF159159_1 cyclin ania-6a [M.musculus]	1281358 IMAGE Consortium	AA874156
9752 F	4 021LAMOY	ESTs, Weakly similar to KIAA1374 protein [H.sapiens]	1379223 IMAGE Consortium	AI050373
9753 F	6 021LAMOY	ESTs	1379846 IMAGE Consortium	AI060394
9754 F	8 021LAMOY	secreted frizzled-related sequence protein 4	1364770 IMAGE Consortium	AI019575
9755 F	10 021LAMOY	C-terminal binding protein 1	1345215 IMAGE Consortium	AA940130
9756 F	12 021LAMOY	ESTs	1346479 IMAGE Consortium	AA958426
9757 H	2 021LAMOY	Mus musculus NOTCH4-like protein (Zneu1) mRNA, complete cds	1281405 IMAGE Consortium	AA874240
9758 H	4 021LAMOY	monocyte macrophage 19	1378509 IMAGE Consortium	AI019319
9759 H	6 021LAMOY	Mus musculus mRNA for ATP-dependent interferon response protein 1 (Adir gene)	1379900 IMAGE Consortium	AI049439
9760 H	8 021LAMOY	ESTs	1365506 IMAGE Consortium	AI020979
9761 H	10 021LAMOY	ESTs	1361416 IMAGE Consortium	AA989995
9762 H	12 021LAMOY	ESTs, Highly similar to AF252290_1 PAR6A [M.musculus]	1345804 IMAGE Consortium	AA967374
9763 B	2 021SAMOZ	mut S homolog 5 (E. coli)	1349053 IMAGE Consortium	AA966993
9764 B	4 021SAMOZ	ESTs, Highly similar to HETEROGENEOUS NUCLEAR RIBONUCLEOPROTEIN K [Homo sapiens; Rattus norvegicus]	1382116 IMAGE Consortium	AA797976
9765 B	6 021SAMOZ	zinc fingers and homeoboxes protein 1	1383175 IMAGE Consortium	AI120674
9766 B	8 021SAMOZ	ESTs, Highly similar to UDP-GalNAc:polypeptide N-acetylgalactosaminyltransferase T6 [R.norvegicus]	1382697 IMAGE Consortium	AI036097
9767 B	10 021SAMOZ	ESTs, Weakly similar to AF233582_1 GTPase Rab37 [M.musculus]	403386 IMAGE Consortium	AI425830
9768 B	12 021SAMOZ	hepatoma-derived growth factor, related protein 3	404454 IMAGE Consortium	W83156
9769 D	2 021SAMOZ	gamma-aminobutyric acid (GABA-A) receptor, subunit rho	1348747 IMAGE Consortium	AA981457

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9770 D	4	021SAMOZ	ESTs	1382141 IMAGE Consortium	AA798210
9771 D	6	021SAMOZ	ESTs	1382469 IMAGE Consortium	AI036150
9772 D	8	021SAMOZ	ESTs	1382734 IMAGE Consortium	AI036116
9773 D	10	021SAMOZ	ESTs	403464 IMAGE Consortium	AI425834
9774 D	12	021SAMOZ	LIM only 2	404586 IMAGE Consortium	W82892
9775 F	2	021SAMOZ	interleukin 11 receptor, alpha chain 1	1349690 IMAGE Consortium	AA980775
9776 F	4	021SAMOZ	J domain protein 1	1382224 IMAGE Consortium	AA798311
9777 F	6	021SAMOZ	ESTs	1383240 IMAGE Consortium	AI120705
9778 F	8	021SAMOZ	ESTs	1382782 IMAGE Consortium	AI120186
9779 F	10	021SAMOZ			
9780 F	12	021SAMOZ	calsenilin/DREAM	406125 IMAGE Consortium	W83313
9781 H	2	021SAMOZ	ESTs	1349832 IMAGE Consortium	AA980898
9782 H	4	021SAMOZ	ESTs	1382235 IMAGE Consortium	AA798371
9783 H	6	021SAMOZ	ESTs, Moderately similar to RBB1_HUMAN RETINOBLASTOMA BINDING PROTEIN 1 [H.sapiens]	1382543 IMAGE Consortium	AI036486
9784 H	8	021SAMOZ	ESTs, Moderately similar to myosin phosphatase target subunit 1 [H.sapiens]	1382856 IMAGE Consortium	AI120239
9785 H	10	021SAMOZ	ESTs	403609 IMAGE Consortium	AI425844
9786 H	12	021SAMOZ	dermis expressed 1	406299 IMAGE Consortium	W83374
9787 B	2	021SAMP0	ESTs	405783 IMAGE Consortium	AI425333
9788 B	4	021SAMP0	ESTs, Weakly similar to SMD1_HUMAN SMALL NUCLEAR RIBONUCLEOPROTEIN SM D1 [M.musculus]	407559 IMAGE Consortium	W88137
9789 B	6	021SAMP0	casein alpha	1383863 IMAGE Consortium	AI181722
9790 B	8	021SAMP0	ESTs	1383115 IMAGE Consortium	AI120472
9791 B	10	021SAMP0	syntrophin, basic 1	1383625 IMAGE Consortium	AI180768
9792 B	12	021SAMP0	macrophage galactose N-acetyl-galactosamine specific lectin	1433164 IMAGE Consortium	AA832918
9793 D	2	021SAMP0	UDP-Gal:betaGlcNAc beta 1,4- galactosyltransferase, polypeptide 2	406506 IMAGE Consortium	W83570
9794 D	4	021SAMP0	ESTs	406903 IMAGE Consortium	W83611
9795 D	6	021SAMP0	ESTs	1383872 IMAGE Consortium	AI461743
9796 D	8	021SAMP0	SEC22 vesicle trafficking protein-like 1 (S. cerevisiae)	1383163 IMAGE Consortium	AI120657
9797 D	10	021SAMP0	ESTs	1383633 IMAGE Consortium	AI180772
9798 D	12	021SAMP0	ESTs	1433414 IMAGE Consortium	AA833037
9799 F	2	021SAMP0	ESTs	405828 IMAGE Consortium	AI425334
9800 F	4	021SAMP0	SRY-box containing gene 18	407667 IMAGE Consortium	AI893332
9801 F	6	021SAMP0	ESTs, Weakly similar to Y281_HUMAN HYPOTHETICAL PROTEIN KIAA0281 [H.sapiens]	1395288 IMAGE Consortium	AI117357
9802 F	8	021SAMP0	ATP-binding cassette, sub-family G (WHITE), member 1	1383276 IMAGE Consortium	AI120733
9803 F	10	021SAMP0	ESTs	1383654 IMAGE Consortium	AI180782
9804 F	12	021SAMP0	gamma-glutamyltransferase-like activity 1	1434243 IMAGE Consortium	AA833290
9805 H	2	021SAMP0	ESTs, Moderately similar to IdlBp [M.musculus]	406629 IMAGE Consortium	W83706



9806 H	4 021SAMP0	ESTs	408629 IMAGE Consortium	AI430293
9807 H	6 021SAMP0	ESTs, Highly similar to dJ50O24.2.1 [H.sapiens]	1397046 IMAGE Consortium	AI098438
9808 H	8 021SAMP0	CD5 antigen	1384126 IMAGE Consortium	AI117259
9809 H	10 021SAMP0	ESTs, Weakly similar to AF116865_1 hedgehog-interacting protein [M.musculus]	1383720 IMAGE Consortium	AI180891
9810 H	12 021SAMP0	histocompatibility-2 complex class 1-like sequence	1432976 IMAGE Consortium	AA833460
9811 B	2 0215AMSI	ESTs	1434695 IMAGE Consortium	AI156999
9812 B	4 0215AMSI	granzyme A	1430039 IMAGE Consortium	AI122257
9813 B	6 0215AMSI	ESTs	1400496 IMAGE Consortium	AI117913
9814 B	8 0215AMSI	casein gamma	1400955 IMAGE Consortium	AI121961
9815 B	10 0215AMSI	melanocortin 2 receptor	1449443 IMAGE Consortium	AI153955
9816 B	12 0215AMSI	ESTs	1449937 IMAGE Consortium	AI196970
9817 D	2 0215AMSI	ESTs	1434829 IMAGE Consortium	AI157060
9818 D	4 0215AMSI	interferon regulatory factor 7	1430219 IMAGE Consortium	AI122517
9819 D	6 0215AMSI	thymidylate kinase homologue	1400538 IMAGE Consortium	AI117832
9820 D	8 0215AMSI	ESTs	1400986 IMAGE Consortium	AI121731
9821 D	10 0215AMSI	RAB23, member RAS oncogene family	1448784 IMAGE Consortium	AI573485
9822 D	12 0215AMSI	REV3-like, catalytic subunit of DNA polymerase zeta RAD54 like (S. cerevisiae)	1450026 IMAGE Consortium	AI197110
9823 F	2 0215AMSI	EST AI428435	1446025 IMAGE Consortium	AI157093
9824 F	4 0215AMSI	ESTs, Moderately similar to KIAA0290 [H.sapiens]	1430379 IMAGE Consortium	AI182149
9825 F	6 0215AMSI	ESTs	1400590 IMAGE Consortium	AI117878
9826 F	8 0215AMSI	ESTs	1399236 IMAGE Consortium	AI120109
9827 F	10 0215AMSI	pale ear	1449566 IMAGE Consortium	AI157856
9828 F	12 0215AMSI	ESTs	1450134 IMAGE Consortium	AI154756
9829 H	2 0215AMSI	ESTs	1446080 IMAGE Consortium	AI157126
9830 H	4 0215AMSI	signal-induced proliferation associated gene 1	1430397 IMAGE Consortium	AI182158
9831 H	6 0215AMSI	butyrophilin	1400638 IMAGE Consortium	AI117962
9832 H	8 0215AMSI	ESTs	1429452 IMAGE Consortium	AI183233
9833 H	10 0215AMSI			
9834 H	12 0215AMSI	ESTs, Weakly similar to mosaic serine protease epitheliasin [M.musculus]	1450203 IMAGE Consortium	AI155132
9835 B	2 0216AMP2			
9836 B	4 0216AMP2	small inducible cytokine A25	1446756 IMAGE Consortium	AI158138
9837 B	6 0216AMP2	ESTs, Highly similar to PLCA_MOUSE 1-ACYL-SN- GLYCEROL-3-PHOSPHATE ACYLTRANSFERASE ALPHA [M.musculus]	1448573 IMAGE Consortium	AI528891
9838 B	8 0216AMP2			
9839 B	10 0216AMP2	ESTs	1477883 IMAGE Consortium	AI152317
9840 B	12 0216AMP2	ESTs	1479915 IMAGE Consortium	AI156363
9841 D	2 0216AMP2	fibroblast growth factor 2	1447251 IMAGE Consortium	AI158649
9842 D	4 0216AMP2	calcitonin	1448200 IMAGE Consortium	AI158820
9843 D	6 0216AMP2	topoisomerase 3 alpha	1449321 IMAGE Consortium	AI153894

9844 D	8 0216AMP2	proliferin	1478339 IMAGE Consortium	AI152562
9845 D	10 0216AMP2	ESTs	1494672 IMAGE Consortium	AI197142
9846 D	12 0216AMP2	ESTs	1479941 IMAGE Consortium	AI156378
9847 F	2 0216AMP2	guanine nucleotide binding protein (G protein), gamma 10	1447284 IMAGE Consortium	AI158565
9848 F	4 0216AMP2	ESTs	1448215 IMAGE Consortium	AI173066
9849 F	6 0216AMP2	hydroxysteroid 17-beta dehydrogenase 7	1450079 IMAGE Consortium	AI154030
9850 F	8 0216AMP2	ESTs	1478377 IMAGE Consortium	AI152637
9851 F	10 0216AMP2	ESTs	1494687 IMAGE Consortium	AI197159
9852 F	12 0216AMP2			
9853 H	2 0216AMP2	carnitine deficiency-associated gene expressed in ventricle 1	1448001 IMAGE Consortium	AI159416
9854 H	4 0216AMP2			
9855 H	6 0216AMP2	homeo box A11	1450222 IMAGE Consortium	AI155130
9856 H	8 0216AMP2	Public domain EST	1477679 IMAGE Consortium	
9857 H	10 0216AMP2	DNA segment, Chr 4, ERATO Doi 27, expressed	1478681 IMAGE Consortium	AI197337
9858 H	12 0216AMP2	fibroblast growth factor receptor 4	1480142 IMAGE Consortium	AI156257
9859 B	2 021DAMP3	placenta specific homeobox 1	1749860 IMAGE Consortium	AI047063
9860 B	4 021DAMP3	ESTs, Highly similar to DAF2_MOUSE COMPLEMENT DECAY-ACCELERATING FACTOR, TRANSMEMBRANE PRECURSOR [M.musculus]	1746318 IMAGE Consortium	AI037616
9861 B	6 021DAMP3	ESTs	1745956 IMAGE Consortium	AI036202
9862 B	8 021DAMP3	ESTs	1510989 IMAGE Consortium	AI225717
9863 B	10 021DAMP3			
9864 B	12 021DAMP3	prolactin-like protein B	1497865 IMAGE Consortium	AI226131
9865 D	2 021DAMP3	ESTs, Weakly similar to The KIAA0135 gene is related to pim-1 oncogene. [H.sapiens]	1746011 IMAGE Consortium	AI036899
9866 D	4 021DAMP3	insulin-like 3	1746336 IMAGE Consortium	AI037624
9867 D	6 021DAMP3	ESTs	1764149 IMAGE Consortium	AI120125
9868 D	8 021DAMP3	ESTs, Highly similar to TSHR_MOUSE THYROTROPIN RECEPTOR PRECURSOR [M.musculus]	1497189 IMAGE Consortium	AI194528
9869 D	10 021DAMP3	ESTs	1497580 IMAGE Consortium	AI664114
9870 D	12 021DAMP3	ESTs	1511278 IMAGE Consortium	AI527654
9871 F	2 021DAMP3	ESTs	1746048 IMAGE Consortium	AI036784
9872 F	4 021DAMP3			
9873 F	6 021DAMP3	ESTs, Moderately similar to AF164349_1 Na,K-ATPase alpha-4 subunit [M.musculus]	1762756 IMAGE Consortium	AI098372
9874 F	8 021DAMP3	pregnancy specific glycoprotein pseudogene 1	1498035 IMAGE Consortium	AI225368
9875 F	10 021DAMP3	ESTs	1617252 IMAGE Consortium	AA982422
9876 F	12 021DAMP3	ESTs, Highly similar to Similar to S.cerevisiae EMP70 protein precursor [H.sapiens]	1617665 IMAGE Consortium	AA968131
9877 H	2 021DAMP3	ESTs	1746075 IMAGE Consortium	AI036800
9878 H	4 021DAMP3	ESTs	1762441 IMAGE Consortium	AI181619
9879 H	6 021DAMP3	ESTs	1749110 IMAGE Consortium	AI046955

9880 H	8 021DAMP3	ESTs	1617046 IMAGE Consortium	AA967800
9881 H	10 021DAMP3	ESTs	1497623 IMAGE Consortium	AI664145
9882 H	12 021DAMP3			
9883 B	2 021KAMP4	glutamate receptor, ionotropic, NMDA2C (epsilon 3)	1852361 IMAGE Consortium	AI256808
9884 B	4 021KAMP4	ESTs, Moderately similar to MHC-TLA-T13a□ [M.musculus]	1765401 IMAGE Consortium	AI173114
9885 B	6 021KAMP4	solute carrier family 19 (sodium/hydrogen exchanger), member 1	537142 IMAGE Consortium	AA117918
9886 B	8 021KAMP4			
9887 B	10 021KAMP4	ATPase, H <sup>+</sup> transporting, lysosomal I	536913 IMAGE Consortium	AA114648
9888 B	12 021KAMP4	solute carrier family 9 (sodium/hydrogen exchanger), isoform 3 regulator 1	538100 IMAGE Consortium	AA114621
9889 D	2 021KAMP4	single-minded 1	1764188 IMAGE Consortium	AI047689
9890 D	4 021KAMP4	Public domain EST	1764729 IMAGE Consortium	
9891 D	6 021KAMP4	ESTs, Weakly similar to AF140691_1 melusin [M.musculus]	537177 IMAGE Consortium	AI550095
9892 D	8 021KAMP4	ESTs	537444 IMAGE Consortium	AA118478
9893 D	10 021KAMP4			
9894 D	12 021KAMP4	ESTs, Highly similar to unnamed protein product [H.sapiens]	538103 IMAGE Consortium	AA114625
9895 F	2 021KAMP4	ESTs	1764215 IMAGE Consortium	AI047720
9896 F	4 021KAMP4			
9897 F	6 021KAMP4	midkine	537195 IMAGE Consortium	AI325764
9898 F	8 021KAMP4	ESTs, Highly similar to JE0239 lin-10 protein - rat□ [R.norvegicus]	537466 IMAGE Consortium	AI550110
9899 F	10 021KAMP4			
9900 F	12 021KAMP4	ESTs	538115 IMAGE Consortium	AA120302
9901 H	2 021KAMP4			
9902 H	4 021KAMP4	ESTs, Weakly similar to breast cancer suppressor candidate 1 [H.sapiens]	537027 IMAGE Consortium	AA114721
9903 H	6 021KAMP4			
9904 H	8 021KAMP4	ubiquitin specific protease 5 (isopeptidase T)	537499 IMAGE Consortium	AA118127
9905 H	10 021KAMP4			
9906 H	12 021KAMP4	DNA segment, Chr 5, ERATO Doi 606, expressed	538143 IMAGE Consortium	AA120307
9907 B	2 021RAMP5	protease (prosome, macropain) 26S subunit, ATPase 5	538322 IMAGE Consortium	AA116811
9908 B	4 021RAMP5	chaperonin subunit 5 (epsilon)	538586 IMAGE Consortium	AI325795
9909 B	6 021RAMP5	guanine nucleotide binding protein, beta 1	538757 IMAGE Consortium	AI325800
9910 B	8 021RAMP5			
9911 B	10 021RAMP5	platelet derived growth factor, alpha	404427 IMAGE Consortium	AI893970
9912 B	12 021RAMP5			
9913 D	2 021RAMP5	ESTs	538335 IMAGE Consortium	AI550166
9914 D	4 021RAMP5	origin of replication 3 homolog (S. cerevisiae)	537862 IMAGE Consortium	AA118183
9915 D	6 021RAMP5	ESTs, Weakly similar to AF099018_1 hepatocyte growth	538825 IMAGE Consortium	AI550199

		factor activator inhibitor type 1 [M.musculus]		
9916 D	8 021RAMP5			
9917 D	10 021RAMP5	inhibitor of DNA binding 2	403789 IMAGE Consortium	AI322367
9918 D	12 021RAMP5	Retinol binding protein 1, cellular {IMAGE:406897}	406897 IMAGE Consortium	W83609
9919 F	2 021RAMP5			
9920 F	4 021RAMP5	gene trap ROSA 26 antisense, Philippe Soriano	537872 IMAGE Consortium	AA119806
9921 F	6 021RAMP5	ESTs, Highly similar to AF257319_1 SH3-containing protein SH3GLB2 [H.sapiens]	538849 IMAGE Consortium	AA123176
9922 F	8 021RAMP5	Mouse mRNA for small GTP-binding protein S10, complete cds {IMAGE:406177}	406177 IMAGE Consortium	W83866
9923 F	10 021RAMP5	putative phosphatase	406066 IMAGE Consortium	W83266
9924 F	12 021RAMP5	Mus musculus Rho-GDI3 mRNA, complete cds {IMAGE:407050}	407050 IMAGE Consortium	W83964
9925 H	2 021RAMP5	ESTs, Weakly similar to KIAA0294 [H.sapiens]	538420 IMAGE Consortium	AA117005
9926 H	4 021RAMP5	ESTs	538623 IMAGE Consortium	AI550184
9927 H	6 021RAMP5	X-linked nuclear protein	989178 IMAGE Consortium	AA571771
9928 H	8 021RAMP5	selenium binding protein 1	400720 IMAGE Consortium	W77429
9929 H	10 021RAMP5	annexin A3	403935 IMAGE Consortium	W82228
9930 H	12 021RAMP5	M.musculus p16K gene for 16 kDa protein {IMAGE:407552}	407552 IMAGE Consortium	W88130
9931 B	2 021YAMP6	coronin, actin binding protein 1C	407653 IMAGE Consortium	AI894359
9932 B	4 021YAMP6	IMAGE EST {IMAGE:424610}	424610 IMAGE Consortium	AA000495
9933 B	6 021YAMP6	CD151 antigen	427184 IMAGE Consortium	AA003005
9934 B	8 021YAMP6	IMAGE EST {IMAGE:439033}	439033 IMAGE Consortium	AA008240
9935 B	10 021YAMP6	delta-aminolevulinate dehydratase	439481 IMAGE Consortium	AA008712
9936 B	12 021YAMP6	Mus musculus mRNA for heme-binding protein, complete cds {IMAGE:457198}	457198 IMAGE Consortium	AA022130
9937 D	2 021YAMP6	early B-cell factor 2	409129 IMAGE Consortium	AA052141
9938 D	4 021YAMP6	ESTs, Highly similar to PROBABLE UBIQUITIN CARBOXYL-TERMINAL HYDROLASE [Mus musculus] {IMAGE:425458}	425458 IMAGE Consortium	AA000038
9939 D	6 021YAMP6	adaptor protein complex AP-1, gamma 1 subunit	426530 IMAGE Consortium	AA002389
9940 D	8 021YAMP6	IMAGE EST {IMAGE:439829}	439829 IMAGE Consortium	AA008529
9941 D	10 021YAMP6	adaptor-related protein complex AP-4, sigma 1	442436 IMAGE Consortium	AA015545
9942 D	12 021YAMP6	Insulin-like growth factor 2 {IMAGE:463135}	463135 IMAGE Consortium	AA030603
9943 F	2 021YAMP6	ribonucleotide reductase M2	407733 IMAGE Consortium	W85674
9944 F	4 021YAMP6	RAB24, member RAS oncogene family {IMAGE:438755}	438755 IMAGE Consortium	AA051083
9945 F	6 021YAMP6	ATPase, H+ transporting, lysosomal (vacuolar proton pump), 42 kDa	422990 IMAGE Consortium	AI325883
9946 F	8 021YAMP6	Mus musculus FAN protein mRNA, complete cds {IMAGE:442332}	442332 IMAGE Consortium	AA013700
9947 F	10 021YAMP6	plectin 1	442473 IMAGE Consortium	AA015107
9948 F	12 021YAMP6	ESTs, Moderately similar to MULTIDRUG RESISTANCE-	459410 IMAGE Consortium	AA027420

		ASSOCIATED PROTEIN 1 [Homo sapiens] {IMAGE:459410}		
9949 H	2 021YAMP6	staufen (RNA-binding protein) homolog 1 (Drosophila)	406367 IMAGE Consortium	W83453
9950 H	4 021YAMP6	Mus musculus thrombospondin-4 mRNA, complete cds {IMAGE:437213}	437213 IMAGE Consortium	AA003452
9951 H	6 021YAMP6	phosphodiesterase 6D, cGMP-specific, rod, delta	438236 IMAGE Consortium	AA008383
9952 H	8 021YAMP6	Amyloid beta (A4) precursor-like protein 2 {IMAGE:444846}	444846 IMAGE Consortium	AA016732
9953 H	10 021YAMP6	dishevelled 2, dsh homolog (Drosophila)	445346 IMAGE Consortium	AA017823
9954 H	12 021YAMP6	Erythrocyte protein band 4.1 {IMAGE:463625}	463625 IMAGE Consortium	AA027522
9955 B	2 0215AMP7	mast cell protease 4	463795 IMAGE Consortium	AA028376
9956 B	4 0215AMP7	Keratin complex 2, basic, gene 1 {IMAGE:476999}	476999 IMAGE Consortium	AA047991
9957 B	6 0215AMP7	EST {IMAGE:480859}	480859 IMAGE Consortium	AA060540
9958 B	8 0215AMP7	signal transducer and activator of transcription 3	479013 IMAGE Consortium	AA049185
9959 B	10 0215AMP7	Inter-alpha trypsin inhibitor, heavy chain 3 {IMAGE:482895}	482895 IMAGE Consortium	AA062129
9960 B	12 0215AMP7	fibromodulin	482824 IMAGE Consortium	AA062196
9961 D	2 0215AMP7	kinesin light chain 2	464393 IMAGE Consortium	AA030661
9962 D	4 0215AMP7	proteasome (prosome, macropain) subunit, beta type 10	476405 IMAGE Consortium	AA050168
9963 D	6 0215AMP7	Mus musculus mRNA for high-sulfur keratin protein, partial cds	480917 IMAGE Consortium	AA064187
9964 D	8 0215AMP7	ATX1 (antioxidant protein 1) homolog 1 (yeast)	479066 IMAGE Consortium	AA049031
9965 D	10 0215AMP7	M.musculus mRNA for vacuolar adenosine triphosphatase subunit B {IMAGE:481410}	481410 IMAGE Consortium	AA060559
9966 D	12 0215AMP7	protein kinase, DNA activated, catalytic polypeptide interacting protein	483576 IMAGE Consortium	AA061743
9967 F	2 0215AMP7	aquarius	466656 IMAGE Consortium	AA031006
9968 F	4 0215AMP7	actinin alpha 3	473018 IMAGE Consortium	AA038082
9969 F	6 0215AMP7	cellular retinoic acid binding protein II	478667 IMAGE Consortium	AA051397
9970 F	8 0215AMP7	tuftelin-interacting protein 33	479809 IMAGE Consortium	AA051492
9971 F	10 0215AMP7	ATPase, Na <sup>+</sup> /K <sup>+</sup> transporting, beta 1 polypeptide {IMAGE:493488}	493488 IMAGE Consortium	AA087098
9972 F	12 0215AMP7	Mus musculus interleukin enhancer binding factor 3 (Ilf3) mRNA, complete cds {IMAGE:571805}	571805 IMAGE Consortium	AA108959
9973 H	2 0215AMP7	sex-lethal interactor homolog (Drosophila)	445990 IMAGE Consortium	AA020052
9974 H	4 0215AMP7	laminin, beta 2	476963 IMAGE Consortium	AA048030
9975 H	6 0215AMP7	adaptor protein complex AP-1, sigma 1	479438 IMAGE Consortium	AA049247
9976 H	8 0215AMP7			
9977 H	10 0215AMP7	keratin complex 1, gene 3	482803 IMAGE Consortium	AA062193
9978 H	12 0215AMP7	COPROPORPHYRINOGEN III OXIDASE PRECURSOR {IMAGE:570602}	570602 IMAGE Consortium	AA108600
9979 B	2 021CAMP8	dipeptidase 1 (renal)	570539 IMAGE Consortium	AI323162
9980 B	4 021CAMP8	tubulointerstitial nephritis antigen	579632 IMAGE Consortium	AA116770

9981 B	6 021CAMP8	kidney-derived aspartic protease-like protein	596420 IMAGE Consortium	AI386398
9982 B	8 021CAMP8	sialyltransferase 7 ((alpha-N-acetylneuraminy) 2,3-betagalactosyl-1,3)-N-acetyl galactosaminide alpha-2,6-sialyltransferase) C	619209 IMAGE Consortium	AA175746
9983 B	10 021CAMP8	vaccinia related kinase 1	659173 IMAGE Consortium	AI644821
9984 B	12 021CAMP8	lymphocyte antigen 6 complex, locus H	670629 IMAGE Consortium	AA221937
9985 D	2 021CAMP8	FXYD domain-containing ion transport regulator 2	570682 IMAGE Consortium	AA108723
9986 D	4 021CAMP8	nucleotide binding protein 1	581981 IMAGE Consortium	AA152999
9987 D	6 021CAMP8	nuclear protein 220	621331 IMAGE Consortium	AA177324
9988 D	8 021CAMP8	histocompatibility 2, class II antigen A, alpha	639586 IMAGE Consortium	AA199011
9989 D	10 021CAMP8	zuotin related factor 2	660560 IMAGE Consortium	AA222952
9990 D	12 021CAMP8	homer, neuronal immediate early gene, 3	670721 IMAGE Consortium	AI466720
9991 F	2 021CAMP8	proline dehydrogenase	574536 IMAGE Consortium	AA124857
9992 F	4 021CAMP8	DNA segment, Chr 3, Nuffield Department of Surgery 3	303581 IMAGE Consortium	W09198
9993 F	6 021CAMP8	hepsin	634773 IMAGE Consortium	AA154674
9994 F	8 021CAMP8	Kruppel-like factor 2 (lung)	642661 IMAGE Consortium	AI606037
9995 F	10 021CAMP8	Rho guanine nucleotide exchange factor (GEF) 1	670987 IMAGE Consortium	AA241401
9996 F	12 021CAMP8	cortactin	660636 IMAGE Consortium	AA222953
9997 H	2 021CAMP8	casein kinase II, alpha 2, polypeptide	575323 IMAGE Consortium	AI386077
9998 H	4 021CAMP8	protein C	314450 IMAGE Consortium	W10700
9999 H	6 021CAMP8	reticulocalbin 2	634095 IMAGE Consortium	AA184659
10000 H	8 021CAMP8	integrin-associated protein	643537 IMAGE Consortium	AA204140
10001 H	10 021CAMP8	coagulation factor II	657895 IMAGE Consortium	AA199269
10002 H	12 021CAMP8	IMAGE EST {IMAGE:676921}	676921 IMAGE Consortium	AA212990
10003 B	2 021JAMP9	peroxisomal sarcosine oxidase	680723 IMAGE Consortium	AA242349
10004 B	4 021JAMP9	Amino acid transporter, cationic 2 (low affinity) {IMAGE:693244}	693244 IMAGE Consortium	AA250170
10005 B	6 021JAMP9	coagulation factor XIII, beta subunit	692419 IMAGE Consortium	AA237845
10006 B	8 021JAMP9	killer cell lectin-like receptor, subfamily D, member 1	721336 IMAGE Consortium	AA267100
10007 B	10 021JAMP9	alcohol dehydrogenase family 1, subfamily A7	734272 IMAGE Consortium	AA268920
10008 B	12 021JAMP9	ESTs	750453 IMAGE Consortium	AA472886
10009 D	2 021JAMP9	glycerol-3-phosphate acyltransferase, mitochondrial	677460 IMAGE Consortium	AA213107
10010 D	4 021JAMP9	Cytochrome P450, 2f2 {IMAGE:695667}	695667 IMAGE Consortium	AA220582
10011 D	6 021JAMP9	aldo-keto reductase AKR1C12	693222 IMAGE Consortium	AA250679
10012 D	8 021JAMP9	suppressor of Lec15 homolog (C.griseus)	720108 IMAGE Consortium	AA254930
10013 D	10 021JAMP9	ESTs {IMAGE:747208}	747208 IMAGE Consortium	AA260490
10014 D	12 021JAMP9	C-reactive protein, petaxin related	738016 IMAGE Consortium	AA271526
10015 F	2 021JAMP9	p100 co-activator	681213 IMAGE Consortium	AA244816
10016 F	4 021JAMP9	M.musculus mRNA for Pr22 protein {IMAGE:716630}	716630 IMAGE Consortium	AA265396
10017 F	6 021JAMP9	lysosomal membrane glycoprotein 2	681744 IMAGE Consortium	AA245754
10018 F	8 021JAMP9	granulin	720182 IMAGE Consortium	AA255191
10019 F	10 021JAMP9	Mus musculus mRNA encoding lysine-ketoglutarate reductase/saccharopine dehydrogenase {IMAGE:738285}	738285 IMAGE Consortium	AA271121

10020 F	12 021JAMP9	B-cell translocation gene 3	751930 IMAGE Consortium	AA396277
10021 H	2 021JAMP9	urate oxidase	679112 IMAGE Consortium	AI551358
10022 H	4 021JAMP9	Cytochrome P450, 2e1, ethanol inducible {IMAGE:696374}	696374 IMAGE Consortium	AA239467
10023 H	6 021JAMP9	Fibroblast growth factor 1 {IMAGE:720400}	720400 IMAGE Consortium	AA261582
10024 H	8 021JAMP9	ubiquitin specific protease 9, X chromosome	720999 IMAGE Consortium	AA266753
10025 H	10 021JAMP9	E2F transcription factor 1 {IMAGE:751755}	751755 IMAGE Consortium	AA396123
10026 H	12 021JAMP9			
10027 B	2 0213AMPA	ubiquitin specific protease 18	751642 IMAGE Consortium	AA396056
10028 B	4 0213AMPA	solute carrier family 22 (organic cation transporter), member 5	790431 IMAGE Consortium	AI608375
10029 B	6 0213AMPA	TRANSCRIPTION FACTOR MTF-1 {IMAGE:849197}	849197 IMAGE Consortium	AA469826
10030 B	8 0213AMPA	RAD51 associated protein 1	849369 IMAGE Consortium	AI552251
10031 B	10 0213AMPA	Mus musculus mRNA for TOM1 protein {IMAGE:873940}	873940 IMAGE Consortium	AA475177
10032 B	12 0213AMPA	nicotinamide nucleotide transhydrogenase	871047 IMAGE Consortium	AI552897
10033 D	2 0213AMPA	myosin X	737186 IMAGE Consortium	AA272581
10034 D	4 0213AMPA	ATP-binding cassette, sub-family G (WHITE), member 3	764392 IMAGE Consortium	AA277174
10035 D	6 0213AMPA	Public domain EST	818229 IMAGE Consortium	
10036 D	8 0213AMPA	macrophage migration inhibitory factor	315447 IMAGE Consortium	W14053
10037 D	10 0213AMPA	Xanthine dehydrogenase {IMAGE:873928}	873928 IMAGE Consortium	AA472074
10038 D	12 0213AMPA	cytochrome P450, steroid inducible 3a13	875431 IMAGE Consortium	AA475351
10039 F	2 0213AMPA	Mus musculus mRNA for Guanine Nucleotide Regulatory Protein, partial cds {IMAGE:776403}	776403 IMAGE Consortium	AA285484
10040 F	4 0213AMPA	ribosomal protein S4, X-linked	776714 IMAGE Consortium	AA273270
10041 F	6 0213AMPA	ESTs, Highly similar to PIAS3 [M.musculus]	849780 IMAGE Consortium	AA450397
10042 F	8 0213AMPA	proteasome (prosome, macropain) 28 subunit, 3	316192 IMAGE Consortium	W12636
10043 F	10 0213AMPA	carnitine acetyltransferase	872058 IMAGE Consortium	AA473526
10044 F	12 0213AMPA	four and a half LIM domains 2	860266 IMAGE Consortium	AA500902
10045 H	2 0213AMPA	CD81 ANTIGEN {IMAGE:775893}	775893 IMAGE Consortium	AA276336
10046 H	4 0213AMPA	phosphoglycerate mutase muscle-specific subunit	807512 IMAGE Consortium	AA412795
10047 H	6 0213AMPA	LIM and cysteine-rich domains 1	807658 IMAGE Consortium	AA470244
10048 H	8 0213AMPA	ribosomal protein S15	316465 IMAGE Consortium	W11185
10049 H	10 0213AMPA	ESTs, Highly similar to JE0378 DNA [R.norvegicus]	851859 IMAGE Consortium	AA499666
10050 H	12 0213AMPA	cytochrome b-245, alpha polypeptide	871947 IMAGE Consortium	AA473417
10051 B	2 021AAMPB	creatine kinase, brain	902005 IMAGE Consortium	AA517889
10052 B	4 021AAMPB	endothelial precursor protein B9	890786 IMAGE Consortium	AA510502
10053 B	6 021AAMPB	Bce-1 protein	949746 IMAGE Consortium	AA537451
10054 B	8 021AAMPB	procollagen-lysine, 2-oxoglutarate 5-dioxygenase 1	1122946 IMAGE Consortium	AA681596
10055 B	10 021AAMPB	LanC (bacterial lantibiotic synthetase component C)-like	1195279 IMAGE Consortium	AA754883
10056 B	12 021AAMPB	myosin heavy chain, cardiac muscle, fetal	1162830 IMAGE Consortium	AA770785
10057 D	2 021AAMPB	ESTs, Highly similar to CASL_MOUSE ENHANCER OF FILMENTATION 1 [M.musculus]	902894 IMAGE Consortium	AA518954
10058 D	4 021AAMPB	interferon gamma inducible protein, 47 kDa	890843 IMAGE Consortium	AA510689

10059 D	6 021AAMPB	ribosomal protein L36	1106464 IMAGE Consortium	AA674987
10060 D	8 021AAMPB	aldo-keto reductase family 1, member B1 (aldose reductase)	1123007 IMAGE Consortium	AA794074
10061 D	10 021AAMPB	neutrophil cytosolic factor 4	1177854 IMAGE Consortium	AA718504
10062 D	12 021AAMPB	PR domain containing 1, with ZNF domain	1165721 IMAGE Consortium	AA710344
10063 F	2 021AAMPB	Bcl2-associated athanogene 1	902140 IMAGE Consortium	AA516684
10064 F	4 021AAMPB	A kinase anchor protein 2	1053953 IMAGE Consortium	AA638739
10065 F	6 021AAMPB	nuclear factor of kappa light polypeptide gene enhancer in B-cells inhibitor-like 1	1139159 IMAGE Consortium	AA655300
10066 F	8 021AAMPB	aldehyde dehydrogenase 2, mitochondrial	1123015 IMAGE Consortium	AA794082
10067 F	10 021AAMPB	exostoses (multiple) 1	1166257 IMAGE Consortium	AA710704
10068 F	12 021AAMPB	tyrosine kinase receptor 1	1196238 IMAGE Consortium	AA794239
10069 H	2 021AAMPB	ATP-BINDING CASSETTE TRANSPORTER 2	903120 IMAGE Consortium	AA521698
10070 H	4 021AAMPB	DNA segment, Chr 9, ERATO Doi 818, expressed	1025625 IMAGE Consortium	AA590873
10071 H	6 021AAMPB	DNA segment, Chr 16, Wayne State University 103, expressed	1106517 IMAGE Consortium	AA930084
10072 H	8 021AAMPB	protein tyrosine phosphatase, receptor-type, F interacting protein, binding protein 2	1066891 IMAGE Consortium	AA616002
10073 H	10 021AAMPB	heat shock protein, 70 kDa 4	1179327 IMAGE Consortium	AA717993
10074 H	12 021AAMPB	transaldolase 1	1195576 IMAGE Consortium	AA711681
10075 B	2 021HAMPC	Mus musculus SH2-containing inositol 5-phosphatase 2 mRNA, complete cds	1181853 IMAGE Consortium	AA711654
10076 B	4 021HAMPC	lymphocyte antigen 6 complex, locus C	1248254 IMAGE Consortium	AA762277
10077 B	6 021HAMPC	cyclin C	1265335 IMAGE Consortium	AA867145
10078 B	8 021HAMPC	solute carrier family 11 (proton-coupled divalent metal ion transporters), member 1	1313921 IMAGE Consortium	AA914844
10079 B	10 021HAMPC	attractin	1346975 IMAGE Consortium	AA958714
10080 B	12 021HAMPC	seven in absentia 1A	1382061 IMAGE Consortium	AA982860
10081 D	2 021HAMPC	S3-12 protein	1195723 IMAGE Consortium	AA711925
10082 D	4 021HAMPC	death-associated kinase 3	1247540 IMAGE Consortium	AA763351
10083 D	6 021HAMPC	nucleoprotein 50	1264626 IMAGE Consortium	AI509632
10084 D	8 021HAMPC	Mouse 19.5 mRNA	1332871 IMAGE Consortium	AI159179
10085 D	10 021HAMPC	T-cell specific GTPase	1380433 IMAGE Consortium	AI060720
10086 D	12 021HAMPC	CD3 antigen, delta polypeptide	1366111 IMAGE Consortium	AI020855
10087 F	2 021HAMPC	Mus musculus calmodulin III (Calm3) mRNA, 3' untranslated region	1181217 IMAGE Consortium	AA754689
10088 F	4 021HAMPC	retinal short-chain dehydrogenase/reductase 1	1265881 IMAGE Consortium	AA881922
10089 F	6 021HAMPC	tumor necrosis factor receptor superfamily, member 10b	1345564 IMAGE Consortium	AI154278
10090 F	8 021HAMPC	integrin beta 2 (Cd18)	1327790 IMAGE Consortium	AA915313
10091 F	10 021HAMPC	ninjurin 1	1382604 IMAGE Consortium	AI036381
10092 F	12 021HAMPC	procollagen, type I, alpha 2	1382197 IMAGE Consortium	AA798297
10093 H	2 021HAMPC	insulin-like 5	1167437 IMAGE Consortium	AA692587
10094 H	4 021HAMPC	ESTs, Highly similar to PROTEIN TRANSPORT	1265180 IMAGE Consortium	AA867862



Accession	Gene ID	Gene Name	Source	Accession
		PROTEIN SEC23 [ <i>Saccharomyces cerevisiae</i> ]		
10095 H	6 021HAMPC	retinoblastoma binding protein 6	1265469 IMAGE Consortium	AA867250
10096 H	8 021HAMPC	inactive X specific transcripts	1332154 IMAGE Consortium	AI877091
10097 H	10 021HAMPC	<i>M. musculus</i> mRNA for glutamyl-tRNA synthetase	1382622 IMAGE Consortium	AI036400
10098 H	12 021HAMPC	small inducible cytokine subfamily B, member 15	1348708 IMAGE Consortium	AA983108
10099 B	2 021OAMPD	geranylgeranyl diphosphate synthase 1	1397751 IMAGE Consortium	AI118639
10100 B	4 021OAMPD	interferon (alpha and beta) receptor 2	1383585 IMAGE Consortium	AI120990
10101 B	6 021OAMPD	septin 6	1429859 IMAGE Consortium	AI122472
10102 B	8 021OAMPD	glutathione S-transferase, mu 5	1430286 IMAGE Consortium	AI181560
10103 B	10 021OAMPD	FMS-like tyrosine kinase 3 ligand	1447902 IMAGE Consortium	AI159345
10104 B	12 021OAMPD	aplysia ras-related homolog G (RhoG)	330841 IMAGE Consortium	AI390848
10105 D	2 021OAMPD	ethanol decreased 2	1383289 IMAGE Consortium	AI120759
10106 D	4 021OAMPD	suppressor of initiator codon mutations, related sequence 1 ( <i>S. cerevisiae</i> )	1383601 IMAGE Consortium	AI182724
10107 D	6 021OAMPD	tumor necrosis factor (ligand) superfamily, member 12	1477875 IMAGE Consortium	AI152313
10108 D	8 021OAMPD	Mouse type I epidermal keratin mRNA, clone pkSCC-50, 3' end	1447723 IMAGE Consortium	AI154635
10109 D	10 021OAMPD	Acid beta glucosidase {IMAGE:331646}	331646 IMAGE Consortium	W08086
10110 D	12 021OAMPD	coagulation factor XII (Hageman factor)	318749 IMAGE Consortium	W34349
10111 F	2 021OAMPD	ESTs, Highly similar to sh3bgr protein [ <i>M. musculus</i> ]	1383330 IMAGE Consortium	AI120826
10112 F	4 021OAMPD	guanine nucleotide binding protein, alpha stimulating	1383669 IMAGE Consortium	AI180799
10113 F	6 021OAMPD	Public domain EST	1434326 IMAGE Consortium	
10114 F	8 021OAMPD	potassium voltage gated channel, shaker related subfamily, beta member 2	1428887 IMAGE Consortium	AI153328
10115 F	10 021OAMPD	cytochrome P450, 2j6	318342 IMAGE Consortium	W13521
10116 F	12 021OAMPD	small EDRK-rich factor 1	318264 IMAGE Consortium	W13619
10117 H	2 021OAMPD	CD24a antigen	1383367 IMAGE Consortium	AI120876
10118 H	4 021OAMPD	menage a trois 1	1383717 IMAGE Consortium	AI117392
10119 H	6 021OAMPD	ilvB (bacterial acetolactate synthase)-like	1493949 IMAGE Consortium	AI156588
10120 H	8 021OAMPD	S100 protein, beta polypeptide, neural	1434731 IMAGE Consortium	AI877462
10121 H	10 021OAMPD	guanine nucleotide binding protein, alpha inhibiting 2	318755 IMAGE Consortium	W34424
10122 H	12 021OAMPD	colony stimulating factor 2 receptor, alpha, low-affinity (granulocyte-macrophage)	2749687 IMAGE Consortium	AW496322
10123 B	2 021VAMPE	synaptotagmin 9	1616776 IMAGE Consortium	AA967661
10124 B	4 021VAMPE	Suppressor of Ty 6 ( <i>S. cerevisiae</i> ) homolog {IMAGE:333778}	333778 IMAGE Consortium	W15976
10125 B	6 021VAMPE	calmodulin 2	333646 IMAGE Consortium	W18160
10126 B	8 021VAMPE	synaptosomal-associated protein, 25 kDa, binding protein	335256 IMAGE Consortium	W17626
10127 B	10 021VAMPE			
10128 B	12 021VAMPE	alpha-2-glycoprotein 1, zinc	337041 IMAGE Consortium	W36761
10129 D	2 021VAMPE	poly A binding protein, cytoplasmic 2	1746311 IMAGE Consortium	AI037666
10130 D	4 021VAMPE	Procollagen, type VI, alpha 1 {IMAGE:334132}	334132 IMAGE Consortium	W16221
10131 D	6 021VAMPE	cathepsin H	333908 IMAGE Consortium	W17422

10132 D	8 021VAMPE	kinesin heavy chain member 1B	333857 IMAGE Consortium	W17492
10133 D	10 021VAMPE	chaperonin subunit 2 (beta)	335090 IMAGE Consortium	W17932
10134 D	12 021VAMPE			
10135 F	2 021VAMPE	close homolog of L1	1511262 IMAGE Consortium	AI527646
10136 F	4 021VAMPE			
10137 F	6 021VAMPE	solute carrier family 1, member 1	334225 IMAGE Consortium	W16288
10138 F	8 021VAMPE	leucine carboxyl methyltransferase 1	333137 IMAGE Consortium	W15851
10139 F	10 021VAMPE	histidine decarboxylase cluster	333681 IMAGE Consortium	AI322263
10140 F	12 021VAMPE	bone gamma carboxylglutamate protein 1	349238 IMAGE Consortium	W29228
10141 H	2 021VAMPE	immunoglobulin kappa chain variable 20 (V20 family)	2749406 IMAGE Consortium	AW496194
10142 H	4 021VAMPE			
10143 H	6 021VAMPE	lymphocyte antigen 84 ligand	334527 IMAGE Consortium	W36468
10144 H	8 021VAMPE	annexin A4	334798 IMAGE Consortium	W36433
10145 H	10 021VAMPE	Keratin complex 2, gene 6A {IMAGE:335736}	335736 IMAGE Consortium	W20986
10146 H	12 021VAMPE	Pak3 binding protein	349586 IMAGE Consortium	W30220
10147 B	2 0212AMPF	neural precursor cell expressed, developmentally down-regulated gene 8	335452 IMAGE Consortium	W17711
10148 B	4 0212AMPF	RAR-related orphan receptor alpha {IMAGE:350310}	350310 IMAGE Consortium	W34685
10149 B	6 0212AMPF	latent transforming growth factor beta binding protein 3	353553 IMAGE Consortium	W40918
10150 B	8 0212AMPF	homeo box C6	368453 IMAGE Consortium	W53691
10151 B	10 0212AMPF	ESTs, Highly similar to KERATIN, TYPE II CYTOSKELETAL 5 [Homo sapiens]	368199 IMAGE Consortium	W53916
10152 B	12 0212AMPF	requiem	374550 IMAGE Consortium	W64068
10153 D	2 0212AMPF	phosphatidic acid phosphatase 2a	335544 IMAGE Consortium	W18632
10154 D	4 0212AMPF	Glycerol phosphate dehydrogenase 1, mitochondrial {IMAGE:351221}	351221 IMAGE Consortium	W41175
10155 D	6 0212AMPF			
10156 D	8 0212AMPF	hemoglobin X, alpha-like embryonic chain in Hba complex	368807 IMAGE Consortium	W56995
10157 D	10 0212AMPF	nuclear protein 95	367559 IMAGE Consortium	W54459
10158 D	12 0212AMPF	open reading frame 6	373913 IMAGE Consortium	W62478
10159 F	2 0212AMPF	recombining binding protein suppressor of hairless (Drosophila)	335555 IMAGE Consortium	AI390138
10160 F	4 0212AMPF	Mus musculus Numbl like (m-nbl) mRNA, complete cds {IMAGE:352882}	352882 IMAGE Consortium	W34802
10161 F	6 0212AMPF	parotid secretory protein	349430 IMAGE Consortium	AI391093
10162 F	8 0212AMPF	milk fat globule-EGF factor 8 protein	368524 IMAGE Consortium	W53856
10163 F	10 0212AMPF	carbonyl reductase 1	368485 IMAGE Consortium	W53728
10164 F	12 0212AMPF	pro-opiomelanocortin-alpha	388641 IMAGE Consortium	W66732
10165 H	2 0212AMPF	timeless homolog (Drosophila)	335766 IMAGE Consortium	W21012
10166 H	4 0212AMPF	Mus musculus activin type IB receptor (ALK-1) mRNA, complete cds {IMAGE:355337}	355337 IMAGE Consortium	W48139
10167 H	6 0212AMPF	insulin-like growth factor binding protein 4	350397 IMAGE Consortium	AI450480
10168 H	8 0212AMPF	hydroxylacyl-Coenzyme A dehydrogenase	368571 IMAGE Consortium	W53871

10169 H	10 0212AMPF	IMAGE EST {IMAGE:385429}	385429 IMAGE Consortium	W62931
10170 H	12 0212AMPF	Bcl-associated death promoter	374295 IMAGE Consortium	W64107
10171 B	2 0211AJ7N	Internal_Control_E	Internal_Control_E	Controls
10172 B	4 0211AJ7N	Internal_Control_F	Internal_Control_F	Controls
10173 B	6 0211AJ7N	Internal_Control_G	Internal_Control_G	Controls
10174 B	8 0211AJ7N	Internal_Control_H	Internal_Control_H	Controls
10175 B	10 0211AJ7N	Internal_Control_I	Internal_Control_I	Controls
10176 B	12 0211AJ7N	Internal_Control_J	Internal_Control_J	Controls
10177 D	2 0211AJ7N	Internal_Control_K	Internal_Control_K	Controls
10178 D	4 0211AJ7N	Internal_Control_L	Internal_Control_L	Controls
10179 D	6 0211AJ7N	Internal_Control_M	Internal_Control_M	Controls
10180 D	8 0211AJ7N	Internal_Control_N	Internal_Control_N	Controls
10181 D	10 0211AJ7N	Internal_Control_O	Internal_Control_O	Controls
10182 D	12 0211AJ7N	Internal_Control_P	Internal_Control_P	Controls
10183 F	2 0211AJ7N	Internal_Control_Q	Internal_Control_Q	Controls
10184 F	4 0211AJ7N	Internal_Control_R	Internal_Control_R	Controls
10185 F	6 0211AJ7N	Internal_Control_S	Internal_Control_S	Controls
10186 F	8 0211AJ7N	Internal_Control_T	Internal_Control_T	Controls
10187 F	10 0211AJ7N	Internal_Control_U	Internal_Control_U	Controls
10188 F	12 0211AJ7N	Internal_Control_V	Internal_Control_V	Controls
10189 H	2 0211AJ7N	Internal_Control_W	Internal_Control_W	Controls
10190 H	4 0211AJ7N	Internal_Control_X	Internal_Control_X	Controls
10191 H	6 0211AJ7N	Internal_Control_Y	Internal_Control_Y	Controls
10192 H	8 0211AJ7N	Internal_Control_Z	Internal_Control_Z	Controls
10193 H	10 0211AJ7N	Internal_Control_AA	Internal_Control_AA	Controls
10194 H	12 0211AJ7N	Internal_Control_AB	Internal_Control_AB	Controls

3. **Samples:** *samples used, extract preparation and labeling:* Male rats (Wistar, Fischer 344 and Fischer x Brown Norway) were purchased from Harlan (Aged 2-6 months) or from the Rat colony at the Gerontology Research Center (GRC), Baltimore MD. Animals were housed in the GRC facility until the planned experimental age. All rats were anesthetized with sodium pentobarbital (6.5 mg/100 g body weight) and weighed. The hearts were removed and the left ventricles (including septum) were dissected free of the right ventricle and atria. Tissue samples were frozen in liquid nitrogen and stored at  $-70^{\circ}\text{C}$  until needed. Frozen tissues were homogenized (Polytron PT3000) and total RNA extracted in RNeasy lysis reagent (Qiagen, Inc). Poly(A<sup>+</sup>) RNA was prepared from total RNA by a single round of selection using Oligotex mRNA kits (Qiagen Inc, CA) according to the manufacturer's instructions. Purified mRNA was quantified using RiboGreen dye (Molecular Probes Inc) in a fluorescent assay. RiboGreen dye was diluted 1:200 (v/v final) and 1:2000 (v/v final) and mixed with known RNA concentrations (determined by absorbance at 260 nm) ranging from 1 to 1000 ng/ml. Ribosomal RNA standards (16S and 23S rRNA from *E. Coli*) were used to generate standard curves and unknown samples were diluted as necessary. Fluorescence was measured in 96-well plates with a Fluoroskan Ascent FL (Thermo Labsystems, Helsinki, Finland) fitted with 480 nm (excitation) and 520 nm (emission) filters.

According to Incyte Genomics, between 25 and 100 ng mRNA were separated on an Agilent 2100 Bioanalyzer, a high resolution electrophoresis system (Agilent Technologies, Palo Alto, CA), to examine the mRNA size distribution. Because this was a cross-species hybridization, 600 ng of purified mRNA were converted to either a Cy3- or Cy5-labeled cDNA probe using a custom labeling kit (Incyte Genomics). Each reaction contained 50 mM Tris-HCl pH 8.3, 75 mM KCl, 15 mM MgCl<sub>2</sub>, 4 mM DTT, 2 mM dNTPs (0.5 mM each), 2 μg Cy3 or Cy5 random 9mer (Trilink, San Diego, CA), 20 U RNase inhibitor (Ambion), 200 U MMLV RNase H-free reverse transcriptase (Promega, Madison, WI) and mRNA. Correspondingly labeled Cy3 and Cy5 cDNA products were combined and purified on a size exclusion column, concentrated by ethanol precipitation and resuspended in hybridization buffer

4. **Hybridizations:** *procedures and parameters:* The hybridization protocols were as described by <sup>1</sup> by Incyte Genomics. Specifically, the combined Cy3 and Cy5 labeled cDNA probes were hybridized in 20 μl of 5x SSC, 0.1% SDS, 1 mM DTT at 60°C for 6 h. Hybridization with a Cy3-labeled vector-specific oligonucleotide (5'-TTCGAGCTTGGCGTAATCATGGTCATAGCTGTTTCCTGTGTGAAATTGTTATCCGCTCA-3') (Operon Technologies) was performed at 10 ng/μl in 5x SSC, 0.1% SDS, 1 mM DTT at 60°C for 1 h. After hybridization, microarrays were washed in 1x SSC, 0.1% SDS, 1 mM DTT at 45°C for 10 min, followed by washing in 0.1x SSC, 0.2% SDS, 1 mM DTT at room temperature for 3 min. Arrays were dried by centrifugation.
5. **Measurements:** *images, quantification and specifications:* After drying, the microarrays were scanned with an Axon GenePix 4000A fluorescence reader and GenePix image acquisition software (Axon) at 535 nm for Cy3 and 625 nm for Cy5 <sup>1</sup>. The Axon scanner used by Incyte was calibrated using a primary standard and a secondary standard to account for the differences in scanner performance [laser and photomultiplier tube (PMT)] between the Cy3 and Cy5 channels. For the primary standard, the company states that hundreds of probe samples

were prepared that were fluorescently balanced in the Cy3 and Cy5 channels as determined by a Fluorolog3 fluorescence spectrophotometer (Instruments SA, Edison, NJ). These probes were hybridized to microarrays and the scanner PMTs were adjusted to give balanced fluorescence and the greatest dynamic range. Using these PMT values a fluorescent plastic slide was scanned to obtain corresponding fluorescent values, and that this secondary standard was used to calibrate scanners on a daily basis. According to Yue et al, low frequency data correction algorithms were additionally applied to compensate for systematic variations in data quality; however, in this study, we took all corrected signal intensities and subjected these values to Log<sub>2</sub> normalizations<sup>4</sup> and statistical tests. Importantly, we only used data where the plot of log Cy3 signal versus log Cy5 signal showed a signal distribution along a line with a slope of 1. In those cases where the center line of the signals did not have a slope of near 1, we assumed different rates of incorporation of the Cy3 and Cy5 dyes, and these data were excluded from subsequent analyses<sup>5</sup>.

Due to the large amount of microarray data, raw data files will be made available on a URL as required by the Journal (if requested). Secondly, image data as tiff files are currently only accessible within GEMTools software, and it is unclear how we can export this information; however, printed images can be supplied upon request.

6. **Normalization controls: types, values and specifications:** As described by the company, an image analysis algorithm (GEMTools software, Incyte Genomics) was used to quantify signal and background intensity for each target element and the ratio of the two corrected signal intensities was calculated and used as the differential expression ratio (DE) for this specific gene in the two mRNA samples. In our study, we only used the corrected signal intensities in our analyses, which involved either Log<sub>2</sub> normalization<sup>4</sup> or primary data analyses with the NIA Array Analysis Tool (<http://microarray.grc.nia.nih.gov/index.html>), a web-based tool for Analysis of Variance of gene microarray data. This was used to define significant changes in transcript abundance with age between relative lifespan groups for both strains.

In this study, all age groups from Wistar and F344 rats were reorganized into relative lifespan groups based on the predicted life expectancy of each rat strains (see text of manuscript). This reorganization allowed an estimation of survival based on the age-specific death rate (i.e., probability of dying). All data were normalized to 6-month old animals, permitting a statistical analysis of four relative lifespan groups: 4%, 8%, 30% and 80%.

## References:

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