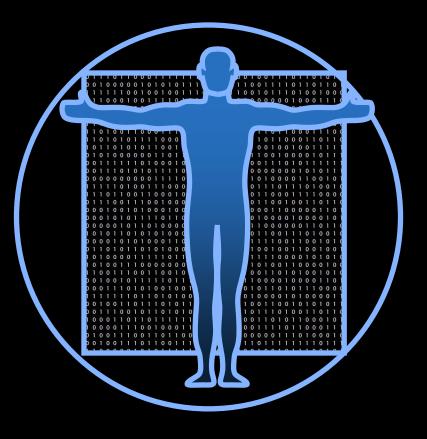
Secretary's Advisory Committee on Genetics, Health and Society

Session on Personal Genome Services July 8, 2008

Personal Genomic Information: A Consumer's Perspective

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THE EXPERIMENTAL MAN PROJECT Genes Environment Brain Body

The Center for Life Science Policy University of California at Berkeley

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Personalized Medicine

- **PAST:** Focus on the ill and the unhealthy
- FUTURE: Focus on the healthy individual -- on prevention and improving health
- I'm not sick (that I know of) and I'm reasonably healthy
- What we're doing is **predicting** an individual's future health

<u>QUESTIONS</u>

- What were your reasons for pursuing personal genome services?
- What sort of information did you anticipate receiving from these services?
- What tests did you take, and what were your results?
- Were there differences in any overlapping results or the interpretation of results from multiple services?

Questions: Expectations

- What were your reasons for pursuing personal genome services?
 - -Journalist
 - -Curiosity about technology and information
 - -Insight to my future health?
- What sort of information did you anticipate receiving from these services?
 - -Low expectations given early phase of science
 - -Confirmation that I am well

Questions: Tests and Results

- What tests did you take, and what were your results?
- Were there differences in any overlapping results or the interpretation of results from multiple services?

<u>Genetic Tests</u>

- SNPS, Insertions, Deletion: ~1.5 Million Genetic Markers
 - Illumina HumanHap 1 Million SNP/650k SNP/330K SNP
 - Affymetrix Genome-Wide Human SNP Array
 6.0
 - Sequenom Mass-Spectrometer (2001-2)

• Dozens of individual genes sequenced

- Quest Diagnostics
- Myriad
- Academic Labs
- Others

Companies, Sites and Labs

2001 (cont.) deCode Genetics Sequenom (Ancestry) Orchid

Diagnostics <u>2007-2008</u> deCode Genetics Institute deCodeme University

2007-2008

DNA Direct Family Tree

Interleukin Quest

> Myriad Coriell

Baylor

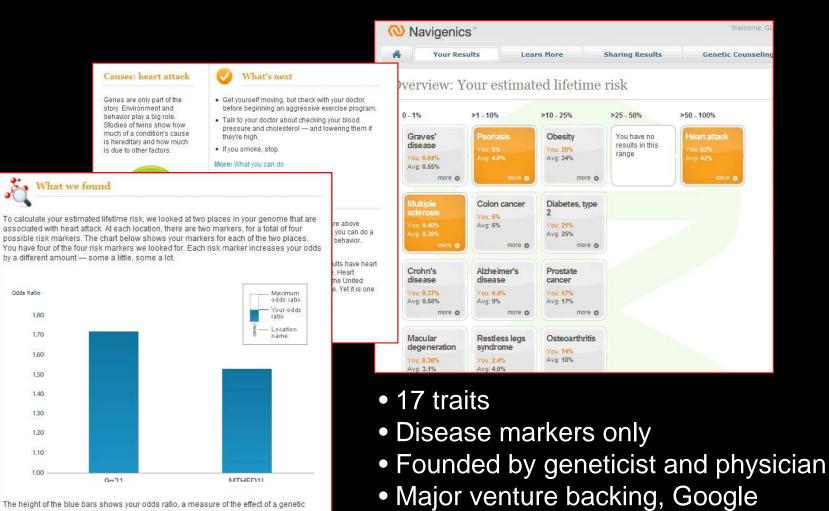
Costs (Genetic Tests)

- 3 Online Consumer Sites
 (Genome-Wide): \$8,500* (David: \$4500 Family: \$4000)
- Ancestral Testing: \$1400 (4 people)*
- DNA Direct (Myriad BRCA1-BRCA2): \$3500*
- Quest Diagnostics (15 tests): TBD (>\$2000?)*
- Other Tests: ~\$5,000*

Participants

- Mother, 75, Artist, Rockport, Maine*
- Father, 76, Architect, Rockport, Maine*
- Brother, 48, Photographer, Brunswick, Maine*
- David, 50, Journalist, San Francisco, California
- Daughter, 19, College Sophomore, St. Andrews,

Navigenics

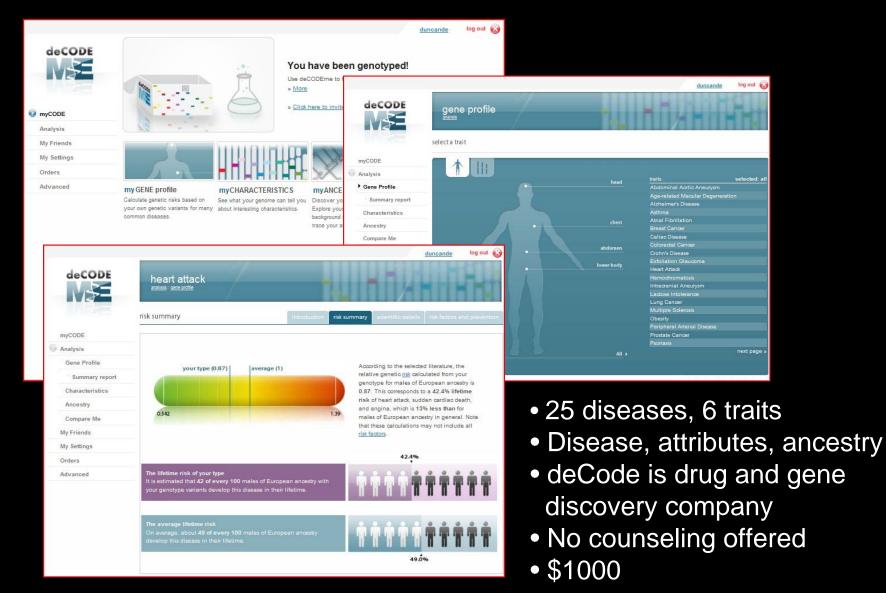


The height of the blue bars shows your odds ratio, a measure of the effect of a genetic variant on your odds of developing a condition. The clear bars represent the maximum odds ratio for each location. Roll your mouse over the chart for more information about each location.

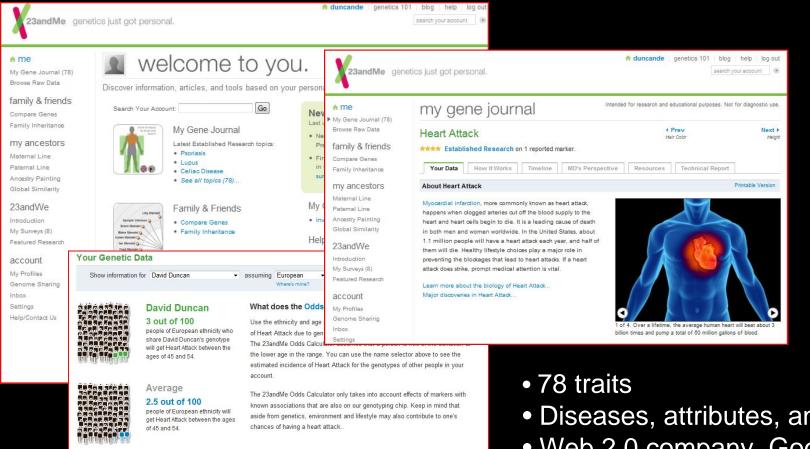
More: Your DNA

- Counseling offered
- \$2500

<u>deCodeme</u>



23andme



Genes vs. Environment



The heritability of death from a heart attack is estimated to be 38% for women and 57% for men. This means that genetic factors contribute slightly less to this condition than environmental factors in women, but contribute slightly more in men. Genetic factors that play a role in heart attack include both unknown factors and known factors such as the SNPs we describe here. Other factors that increase your risk include being older, being male, being African-American, smoking, having high blood cholesterol or high blood pressure, physical inactivity, being overweight, having diabetes. alcohol use, and stress. (sources)

- Diseases, attributes, ancestry
- Web 2.0 company, Google
- Rating system
- Counseling not offered
- \$1000

Two Other Approaches

• DNA Direct

- Online ordering and results, physician signs orders
- Offers only individual tests in common use by physicians
- Counseling before and after
- Rich information, including pros and cons of testing

• Coriell Institute

- Genome-wide data (Affymetrix)
- -15 or so diseases, website
- Nonprofit: free for 10,000-100,000 people

Sample Results

Red: risks over 1.5 times normal
Orange: risks over 1.2 time normal
Black: Average or normal risk
Yellow: Between .5 and .99 times normal
Green: Protective SNP or risk factor below .5

Age-Related Macular Degeneration

| Trait | Gene | Marker | Resul ts | Risk * | Source | Life Risk* DED |
|-------------|-------------|------------|-------------|-----------|----------------|-------------------|
| Age-Related | PLEKHA1/ARM | rs932275 | | | deCodeme | |
| Macular | S2 CFH | rs1329428 | GG G AA | 0.68 | deCodeme | 8.0% |
| Degeneratio | CFH | rs10737680 | А | 0.20 | Navigenic | 0.36% 3.1% |
| n | LOC387715 | rs10490924 | CC T | 1.0 | s Navigenic | |
| | CFB | Rs541862 | GG T | 1.0 | s Navigenic | |
| | LOC387715 | rs3750847 | 111 | 6.98 | S 23andme | 0.19% |
| | CFH | rs1061147 | CC A | 0.46 | 23andme | 1.2% |
| | | | CC | 0.34 | | |

*Sites use different methods for determining risk factors.

More Results: Comparing 3 Sites

Diabetes Type II

- 19 Different SNPs
 - 23andme: 9 Navigenics: 11 deCodeme: 10
- 15 Different Genes
- Range of SNP risk factors: 0.82 2.61
 - Lifetime Risk: 23andme: 16.8% -Navigenics: 21% - deCodeme: 18.8% -Average for U.S. Male: 25%
- 4 SNPs on all 3 sites (2 of 4 risk factors consistent)
- 4 SNPs on 2 out of 3 sites (4 out of 4 risk factors consistent)

Is Data Consistent?

- Genotyping Results (CLIA Lab): very consistent among 3 sites (GG or AG is the same)
- Risk Factor Results: mostly consistent
- Risk Factors by Disease, regardless of site: not always consistent (mix of high, med and low)

Heart Attack Gene Markers

| <u>Gene /</u> Locati | <u>SNP</u> | <u>Risk</u> Vari | <u>DED</u> <u>Resul</u> | <u>Risk</u> <u>Facto</u> | <u>Source</u> | <u>Lifetim</u> <u>e Risk*</u> |
|-------------------------|----------------|----------------------|----------------------------|-----------------------------|----------------|----------------------------------|
| CELSR2 + | rs599839 | an t G | <u>ts</u> AG | <u>r*</u> 86 | deCodeme | 42% 49% |
| 9p21 | rs101162 77 | Т | GT | 1.0 | deCodeme | |
| 9p21 | rs133304 9 | С | CC | 1.72 | Navigeni cs | 62% 49% |
| MTHFD1 L | rs692226 9 | A | AA | 1.53 | Navigeni cs | |
| 9p21 | rs238320 7 | G | GG | 1.22 | 23andme | 29.9% 17% |

*Risk factors for each site are calculated differently.

Why Different Results?

- Different SNPs/studies used
- Different methods for determining SNP risk
 - -deCodeme: Relative Risk
 - -23andme and Navigenics: odds ratios
- Different methods for determining combined SNPs risk/lifetime risk
- Reliance on correlative SNPs

End Result: head scratching, what does it mean?

Three Generation Study



| Heart Attack (| rs1075728) | Alzheimer's (rs4420638) | | | | | | |
|------------------------------|------------|-------------------------|-------------|---------|--|--|--|--|
| Father | Mother | | Father | Mother | | | | |
| AG | AG | | AG | AA | | | | |
| David | Brother | | David | Brother | | | | |
| GG | AA | | AA | AG | | | | |
| Daughter A <mark>G</mark> | | | Daugh AA | iter | | | | |

Two Brothers

(Rare Diseases vs. Common Diseases)

Disease: Osteogenesis Imperfecta (OI)

Full Sequence: COLA1A and COLA2A

Lab: Peter Byers, University of Washington

Results:

Donald Duncan, 48: Deletion in COLA1A = Positive for OI

David Duncan, 50: Normal COLA1A = No OI

Q: Should rare diseases be part of

"Recreational" and "Preliminary"

| <u>Trait</u> | <u>SNP</u> | <u>Risk</u> | DED | <u>Rati</u> | <u>Risk</u> | Source | | |
|------------------|--------------------|--------------|--------------|-------------|----------------|---------------------|--|--|
| | | | <u>Resul</u> | <u>nq</u> | <u>Factor</u> | | | |
| Ancestry | mtDNA | | Group | 5 | European | deCodeme | | |
| | | | H | | Ancestry | / | | |
| Bitter | rs71359 | G=bit | CC | 5 | No bitter | Navigeni 23andme | | |
| Taste | 8 | ter | TT | 5 | No bitter | deCodeme | | |
| Intellig | rs17268 rs36305 | T=bit ter | GG | 1 | Lower IQ | 23andme | | |
| ence Avoiding | rs18004 | A | GG | 2 | 3pts Avoids | 23andme | | |
| Errors | 97 | | | | Errors | | | |
| Back | rs20737 | G | GG | 2 | Average | 23andme | | |
| Pain Heroin | 11 rs17999 | G | AG | 1 | Substantia | 23andme | | |
| Addictio | 71 | | | | lly Higher | | | |
| n Longevit | rs25420 | С | CC | 1 | Higher | 23andme | | |
| У | 52 | | | | odds age | | | |
| Caffeine | rs76255 | А | AA | | 100 Rapid | 23andme | | |
| | 1 | | | | Metabolize | | | |
| | | | | | r | | | |

Crush of Data... Chart 24 ft. Long

| EXPERIMENTAL M | 1AN | | | | | | | | | | - | | - | | | | | |
|------------------|---------------------------|------------|---|------------|-----------|----------|-----------|------------|----------|-------|---------|-------|--------|-------|--|--------|----------|--------|
| Main Graph - Gen | e Test Results | | | | | | | | | | | | | | | | | |
| | | | for the second | 222010.000 | LOCETTE - | 1 | | 1111111 | | | | | | | | | | |
| | High Risk (over 1.5) | | Relative Ri | | | | 5 most u | | | 3 | | | | | | | | |
| | Low Risk (over 1.2) | | Odds Ratio | (no bold) | | | n Studied | and Malaka | | | | | | | | | | |
| | Average Risk (1.0) | | | | | 000 Popu | | | | | | | | | | | | |
| | Below Average Risk | (Below 1.0 |) | | | 5000 Pop | | | | | | | | | | | | |
| | Protective | | | | | | opulation | | | 19 | | | | | | | | |
| | | | | | 5: >10,0 | 00 Popul | ation | | | 5 | | | | | | | | |
| TRAIT | | | | CRITERIA | DED RES | SULTS | THREE | SENERAT | TION STU | IDY | | | - | | LOCATION, | POPULA | TIONS, S | OURCES |
| Туре | Condition | Gene | SNP/Mark | | | Risk | Father | | Mother | | Brother | Risk | Daught | Risk | Sources | Locii | | Rating |
| della maria | | | | | | | | | | | | | | | 1. | | | |
| I. ANCESTRAL | Ancestry | mDNA | NA | NA | н | * | Н | * | н | * | Н | * | U5 | * | deCodeme | mDNA | N | |
| | Ancestry | Y | NA | NA | R1B | * | R1B | * | NA | * | R1B | * | NA | * | deCodeme | Y | | |
| | %European | Auto | NA | NA | * | 95 | * | 93 | * | 93 | * | 94 | * | 95 | deCodeme | 3. | | |
| | | Auto | NA | NA | 282 | 3 | * | 5 | * | 3 | * | 4 | * | 3 | deCodeme | | | |
| | %African | Auto | NA | NA | * | 2 | * | 2 | * | 4 | * | 2 | * | 2 | deCodeme | X | | |
| | %European | х | NA | NA | * | 87 | 240 | 87 | 240 | 91 | 540 | 92 | * | 93 | deCodeme | X | | |
| | %Asian | x | NA | NA | * | 11 | * | 10 | * | 8 | * | 8 | * | 5 | deCodeme | | | |
| | %African | x | NA | NA | * | 2 | sic | 3 | | 1 | * | <1 | * | 2 | deCodeme | | | |
| | Orcadia (Scotland) | NA | NA | NA | 30 | 1 | sic | 1 | * | 1 | * | 2 | 36 | 1 | deCodeme | | | |
| | French | NA | NA | NA | * | 2 | 90 | 2 | * | 2 | * | 1 | 342 | 2 | deCodeme | | | |
| | Basque | NA | NA | NA | * | 3 | sic | 4 | * | 4 | * | 5 | 340 | 3 | deCodeme | | | |
| | | NA | NA | NA | * | 4 | 280 | 3 | 38 | 3 | * | 3 | * | 4 | deCodeme | | | |
| | Italian | NA | NA | NA | * | 5 | 940 | 6 | * | 6 | 80 | 6 | 180 | 5 | deCodeme | - 21 | | 1 |
| | Russia | NA | NA | NA | * | 6 | 94 | 5 | * | 5 | * | 4 | 36 | 6 | deCodeme | | | |
| | | | in the second | | | | | | | | 1 | | 1 | | | | | |
| II. ATTRIBUTES | Blue Eyes | SLC24A4/T | rs12913832 | G (blue) | GG | 94.17 | * | 96.62 | * | 90.48 | * | 96.62 | * | 85.37 | SNPedia | | * | |
| | Blue Eyes | OCA2 | rs1667394 | A (blue) | AA | 90 | * | | * | | 90 | | * | | deCode | 5 | | |
| | Brown Eyes | SLC24A4/T | * | * | * | 2.3 | * | 1.81 | * | 2.55 | * | 1.81 | * | 4.22 | * | | * | |
| | Green Eyes | SLC24A4/T | rs7495174 | A (blue) | AA | 3.52 | * | 1.58 | * | 6.96 | * | 1.58 | * | 10.41 | SNPedia | | * | |
| | Blond Hair | SLC24A4/K | * | * | * | 24.21 | * | 13.28 | * | 21.3 | * | 25.65 | * | 28.58 | * | | * | |
| | Brown Hair | SLC24A4/K | * | * | * | 74.92 | * | 22.85 | * | 77.83 | * | 73.48 | * | 70.55 | * | 5 | * | |
| | Red Hair | SLC24A4/K | * | * | * | 0.87 | * | 63.87 | * | 0.87 | * | 0.87 | * | 0.87 | * | | * | |
| 11 | Height | HMGA2 | rs1042725 | С | CT | .05 cm | * | | * | | * | | * | | Nature Gene | tics | * | |

- Total Markers (so Far): 1000+ (24 feet long when printed out)
- Five Sections: Ancestral, Attributes, Behavioral, Disease, Environment
- 3-Generation Study
- Risk Factors
- Rating System

Step 3: Reactions, Thoughts

- Did you alter your behavior in light of test results? If so, how?
 - One person journalist, tested on multiple sites
 - Not really... subsequent heart tests
 - convinced me to alter my diet - Breast cancer data (high risk SNPs) for

PLUSES OF DTC TESTING

- Insight into personal and societal health
- Personal empowerment
- Will push society (and health industry) to discuss guidelines, ethics, education, and funding
- Opening up new avenues for research impacting individuals and subgroups

MINUSES OF DTC TESTING

- Early days of technology
- Association studies not always applicable to individuals
- Disease and non-disease results mixed
- No standards for validity, risk factors
- Physicians not trained in genetics
- Potential to frighten
- High costs, no insurance (costs

THOUGHTS AND SUGGESTIONS

- Consumers should be free to access their information and buy services
- Encourage discussion
- Early adopters should be part of the experiment - Coriell approach, doctor's first
- Establish standards and guidelines for tests <u>and</u>

THOOGHID AND SOGGESTANS

(Cont)

- Crash program to set validation standards, refocus on preventive medicine
- Disease markers should be handled differently; counseling offered
- Physicians in companies should review disease markers, alert consumers of serious findings
- Companies should provide lists

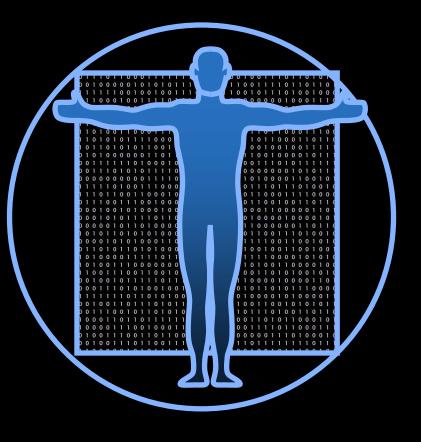
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