

Proposed NHGRI Reorganization



Eric Green, M.D., Ph.D. Director, NHGRI



- I. Background on NHGRI
- II. Proposed Reorganization
- III. Process for Implementation





Historical Context: 'The Genome Institute'

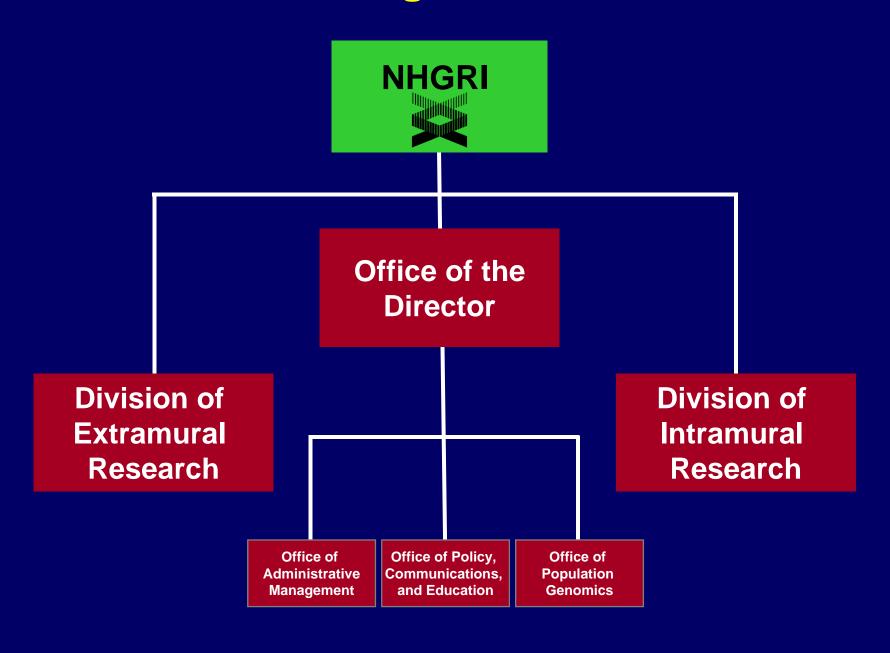


Office for Human Genome Research
1988-1989

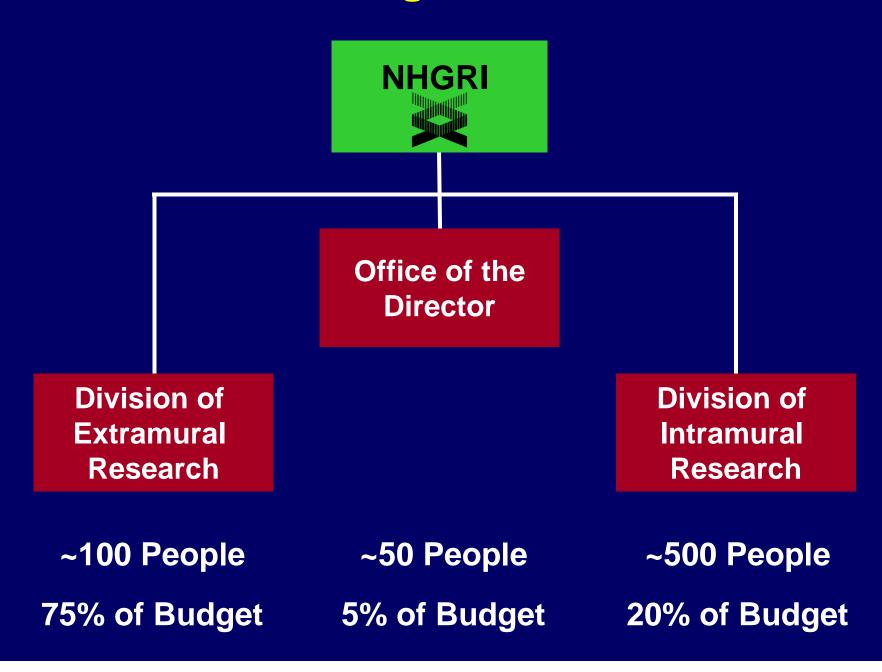
National Center for Human Genome Research 1989-1997

National Human Genome Research Institute 1997-present

Current NHGRI Organizational Structure



Current NHGRI Organizational Structure



Additional Background

- Organizational structure of the Extramural Research Program has been essentially unchanged since the Human Genome Project largely a 'flat' (non-hierarchical) structure
- The Office of the Director has grown in mission, complexity, and scale in recent years, commensurate with the Institute's expanding research portfolio

The Different Eras of NHGRI

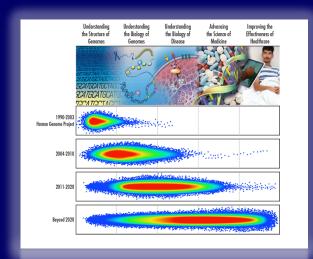
1990-2003

2003-2010

2011-??













~1 Year Ago



PERSPECTIVE

Charting a course for genomic medicine from base pairs to bedside

Eric D. Green¹, Mark S. Guyer¹ & National Human Genome Research Institute

There has been much progress in genomics in the ten years since a draft sequence of the human genome was published. Opportunities for understanding health and disease are now unprecedented, as advances in genomics are harnessed to obtain robust foundational knowledge about the structure and function of the human genome and about the genetic contributions to human health and disease. Here we articulate a 2011 vision for the future of genomics research and describe the path towards an era of genomic medicine

S ince the end of the Human Genome Project (HGP) in 2003 and the publication of a reference human genome sequence¹², genomics has become a mainstay of biomedical research. The scientific community of the scientific community range of scientific advances that the HGP has enabled, as shown in Fig. 1 (see rollfold). Optimism about the potential contributions of genomics for improving human health has been fuelled by new insights about cancer⁴⁻⁷, the molecular basis of inherited diseases (http://www.ncbi.nlm.nih.gov/ omim and http://www.genome.gov/GWAStudies) and the role of structural variation in disease⁸, some of which have already led to new therapies⁹⁻¹⁵. Other advanceshave alreadychanged medical practice (for example, micro-arrays are now used for dinical detection of genomic imbalances¹⁴ and pharmacogenomic testing is routinely performed before administration of certain medications 15). Together, these achievements (see accompanying paper¹⁶) document that genomics is contributing to a better understanding of human biology and to improving human health.

As it did eight years ago¹⁷, the National Human Genome Research Institute (NHGRI) has engaged the scientific community (http://www. genome gov/Planning) to reflect on the key attributes of genomics (Box 1) and explore future directions and challenges for the field. These discussions have led to an update dvision that focuses on understanding human biology and the diagnosis, prevention and treatment of human disease including consideration of the implications of those advances for society (but these discussions, intentionally did not address the role of genomics in agriculture, energy and other areas). Like the HGP, achieving this vision is broader than what any single organization or country can achieve realizing the full benefits of genomics will be a global effort.

This 2011 vision for genomics is organized around five domains extending from basic research to health applications (Fig. 2). It reflects the view that, over time, the most effective way to improve human health is to understand normal biology (in this case, genome biology) as a basis for understanding disease biology, which then becomes the basis for improving health. At the same time, there are other connections among these domains. Genomics offers opportunities for improving health without a thorough understanding of disease (for example, cancer therapies can be selected based on genomic profiles that identify tumour subtypes^{18,19}), and clinical discoveries can lead back to understanding disease or even basic biology.

The past decade has seen genomics contribute fundamental knowledge

about biology and its perturbation in disease. Further deepening this understanding will accelerate the transition to genomic medicine (clinical care based on genomic information). But significant change rarely comes

nity's foresight in launching this ambitious project* is evident in the broad (Fig. 2). Achieving such progress will depend not only on research, but also on new policies, practices and other developments. We have illustrated the kinds of achievements that can be anticipated with a few examples (Box 2) where a confluence of need and opportunities should lead to major accomplishments in genomic medicine in the coming decade. Similarly, we note three cross-cutting areas that are broadly relevant and fundamental across the entire spectrum of genomics and genomic medicine: bioinformatics and computational biology (Box 3), education and training (Box 4), and genomics and society (Box 5).

Understanding the biology of genomes

Substantial progress in understanding the structure of genomes has revealed much about the complexity of genome biology. Continued acquisition of basic knowledge about genome structure and function will be needed to illuminate further those complexities (Fig. 2). The contribution of genomics will include more comprehensive sets (catalogues) of data and new research tools, which will enhance the capabilities of all researchers to reveal fundamental principles of biology

Comprehensive catalogues of genomic data

Comprehensive genomic catalogues have been uniquely valuable and widdy used. There is a compelling need to improve existing catalogues and to generate new ones, such as complete collections of genetic variation, functional genomic elements, RNAs, proteins, and other biological molecules, for both human and model organisms.

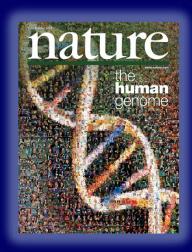
Genomic studies of the genes and pathways associated with disease related traits require comprehensive catalogues of genetic variation, which provide both genetic markers for association studies and variants for identifying candidate genes. Developing a detailed catalogue of variation in the human genome has been an international effort that began with The SNP Consortium20 and the International HapMap Project2 (http://hapmap. nchi.nlm.nih.gov), and is ongoing with the 1000 Genomes Project (http://www.1000genomes.org).

Over the past decade, these catalogues have been critical in the discovery of the specific genes for roughly 3,000 Mendelian (monogenic) diseases

¹National Human Genome Research Institute, National institutes of Health, 31 Center Dr., Bethesda, Maryland 20892-2152, USA.
*Usts of participants and their affiliations appear at the end of the paper.

February 2011 NHGRI Published New Vision for Genomics

The Path to Genomic Medicine

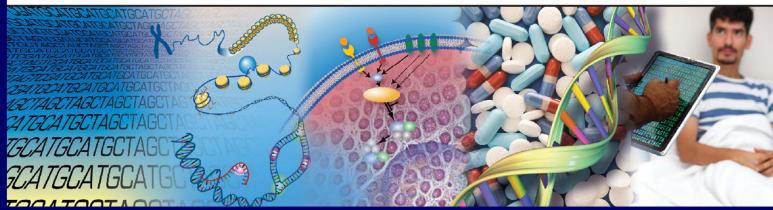


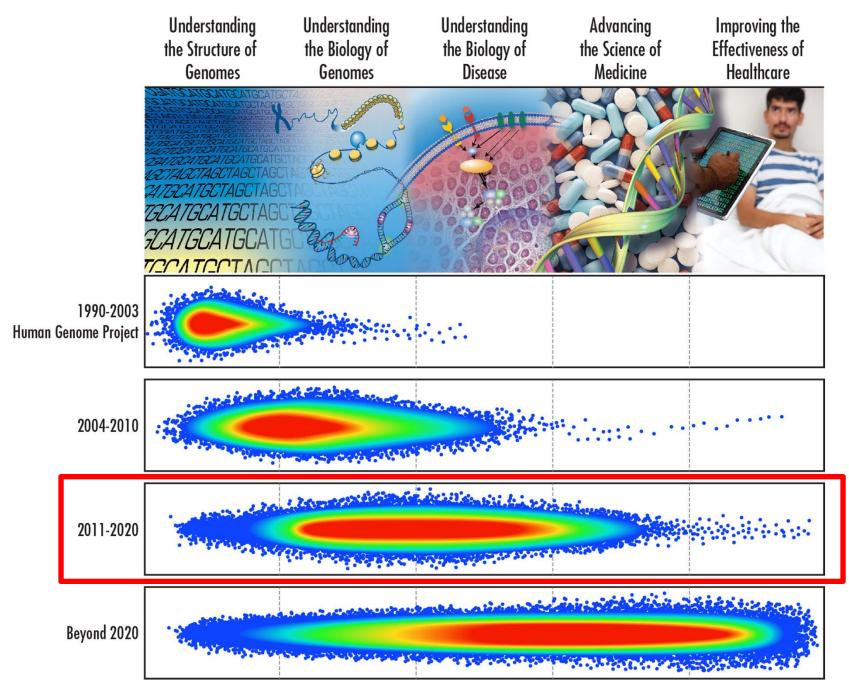


Human Genome Project

Realization of Genomic Medicine

Understanding the Structure of Genomes Understanding the Biology of Genomes Understanding the Biology of Disease Advancing the Science of Medicine Improving the Effectiveness of Healthcare





Green et al. (2011)

NHGRI Extramural Research Program: Circa 'Then'



Human Genome Project

NHGRI Extramural Research Program: Circa 'Now'

\$1000 Genome

Technology Development Program

TCGA

The Cancer Genome Atlas

KOMP

Knockout Mouse Project

PAGE

Population Architecture using Genomics and Epidemiology

1000 Genomes

ENCODE

Encyclopedia of DNA Elements Project

modENCODE

Model Organism ENCODE

Clinical Sequencing

Mendelian Disorders
Sequencing

Large-Scale Sequencing Program

eMERGE

Electronic Medical Records and Genomics

PhenX

Consensus Measures for Phenotypes and eXposures

GENEVA

Gene Environment Association Studies

CEGS

Centers of Excellence in Genomic Science

FI SI

Ethical Legal Social Implications Program

GARNET

Genomics and Randomized
Trials Network

CEER Program

Centers for Excellence in ELSI Research

KOMP2

KOMP Phenotyping

HMP

Human Microbiome **Project**

GTEx

Genotype-Tissue Expression

H3Africa

Human Heredity and Health in Africa

Protein Capture Reagents

LINCS Library of Integrated Net

Library of Integrated Networkbased Cellular Signatures

MLP

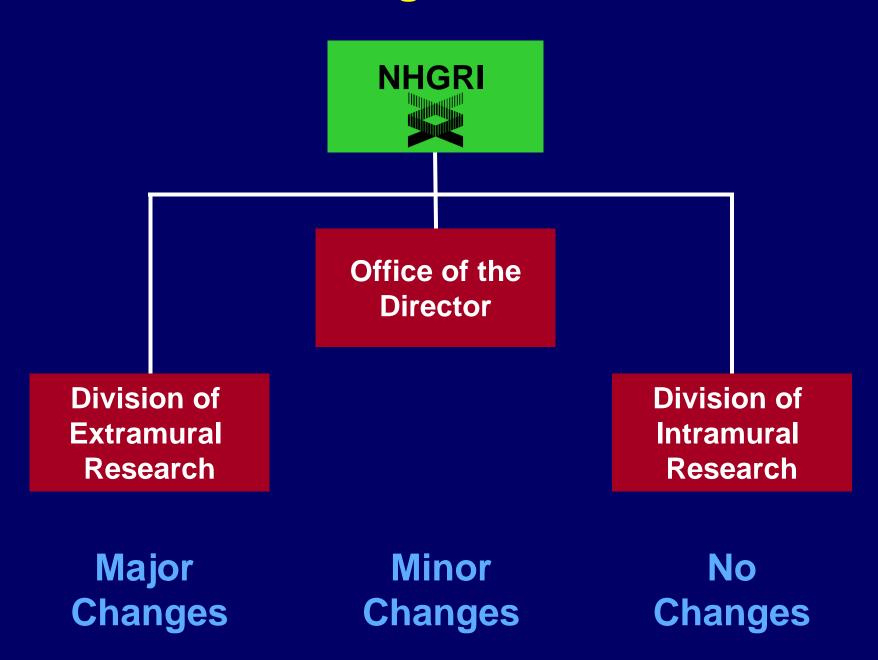
Molecular Libraries
Program

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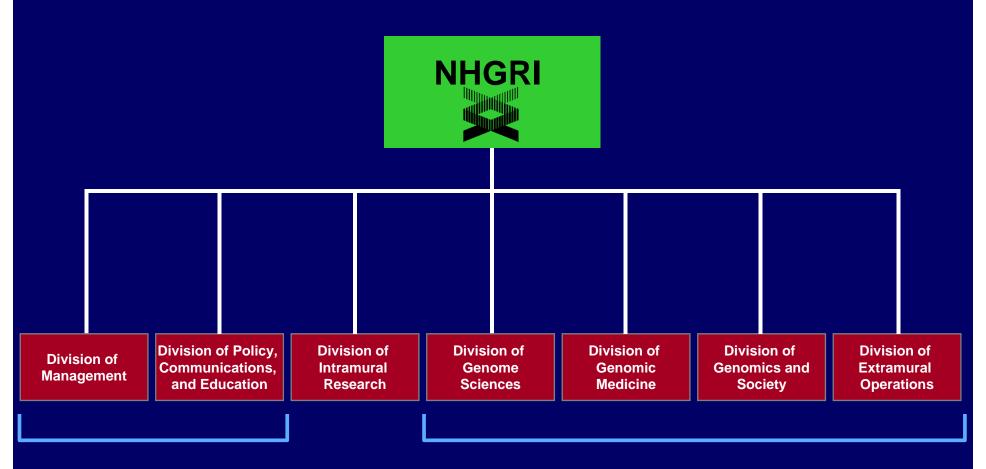




Current NHGRI Organizational Structure

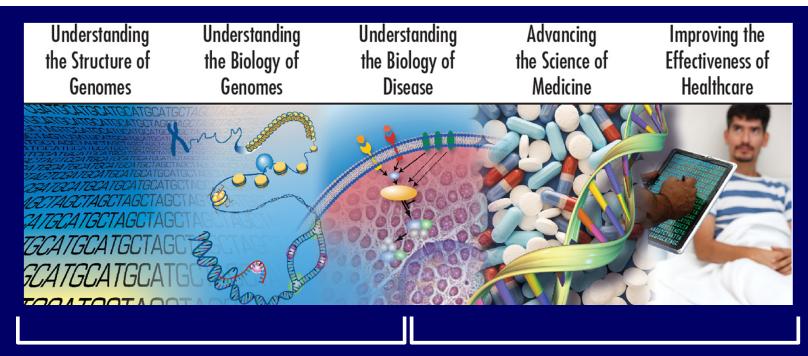


Proposed NHGRI Organizational Structure



From the Office of the Director

Extramural Research Program



Division of Genome Sciences

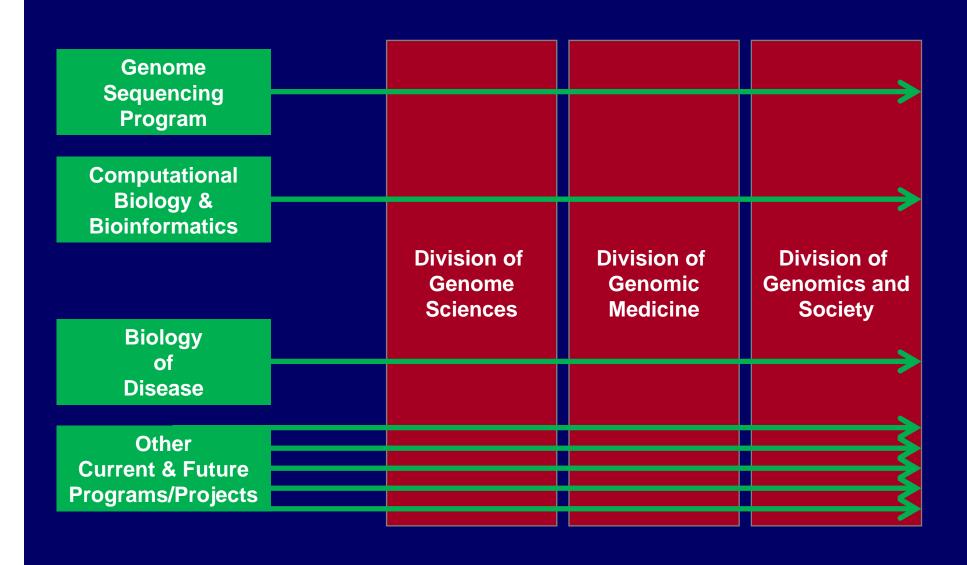
Division of Genomic Medicine

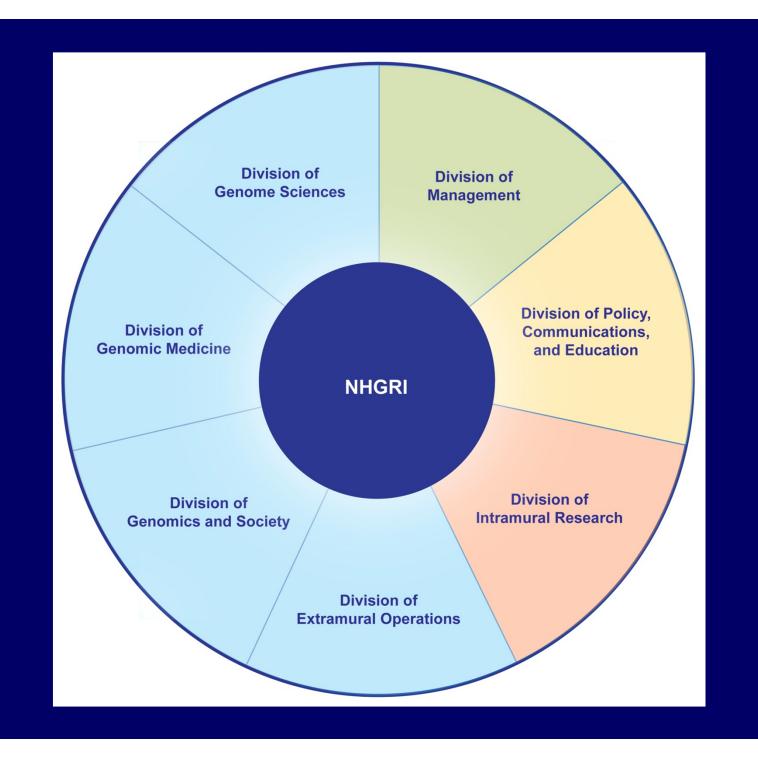


Genomics & Society

Division of Genomics and Society

Program/Project Oversight and Execution





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NIH Reform Act of 2006

H. R. 6164

One Hundred Minth Congress of the United States of America

AT THE SECOND SESSION

Begun and held at the City of Washington on Tuesday, the third day of January, two thousand and six

An Act

To amend title IV of the Public Health Service Act to revise and extend the authorities of the National Institutes of Health, and for other purposes.

Be it enacted by the Senate and House of Representatives of the United States of America in Congress assembled,

SECTION 1. SHORT TITLE.

This Act may be cited as the "National Institutes of Health Reform Act of 2006".

TITLE I-NIH REFORM

TITLE I-NIH REFORM

Reform Act of 2006".

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the United States of America in Congress assembly Representatives of



Federal Register Posting



Federal Register/Vol. 77, No. 10/Tuesday, January 17, 2012/Notices

Time: 2 p.m. to 3 p.m. Agenda: To review and evaluate grant

Place: National Institutes of Health, 6701 Rockledge Drive, Bethesda, MD 20092, (Telephone Conference Call).

Contact Person: Lynn E Luethke, Ph.D., Scientific Review Officer, Center for Scientific Review, National Institutes of Health, 6701 Rockledge Drive, Room 5166, MSC 7844, Bethesda, MD 20892, (301) 806– 3223. Jueth kellSer, nil, nov.

Name of Committee: Center for Scientific Review Special Emphasis Panel, Collaborative Applications in Adult Psychopathology and Disorders of Aging, The Wales of Aging,

Time: 4 p.m. to 5 p.m. Agendo: To review and evaluate grant

Place: Doubletree Guest Suites Santa Monica, 1707 Fourth Street, Santa Monica, CA 90401.

Contact Person: Mark Lindner, Ph.D., Scientific Review Officer, Center for Scientific Review, National Institutes of Health, 6701 Rockledge Drive, Room 3182, MSC 7770, Bethasda, MD 20892, (201) 435– 0913, mark lin desr@cc.nih,gov.

Name of Committee: Center for Scientific Review Special Emphasia Panel, Hypertension and Microcirculation A. Date: February 7, 2012.

Time: 1 p.m. to 1:45 p.m. Agenda: To review and evaluate grant

applications.
Place: National Institutes of Health, 6701
Rockledge Drive, Bethesda, MD 20092,
(Telephone Conference Call).

Contact Person: Larry Pinkus, Ph.D., Scientific Roview Officer, Center for Scientific Roview, National Institutes of Health, 6701 Rockledge Drive, Room 4132, MSC 7802, Bethneda, MD 20092, (301) 435– 1214, piskus/Norr aith.sov.

(Catalogue of Federal Demestic Assistance Program Nos. 23.306, Comparative Medicines 23.333, Clinical Rossarch, 23.306, 23.333, 23.337, 23.393–23.396, 23.837–23.844, 23.846–23.878, 23.892, 23.893, National lastitutes of Health, HHS)

Dated: January 9, 2012.

Jennifer S. Spaeth,

Director, Office of Federal Advisory
Committee Policy.

[FR Doc. 2012–732 Piled 1–13–12; 8:45 am] BELING CODE 4146-01-P

DEPARTMENT OF HEALTH AND

National Institutes of Health

National Human Genome Research Institute; Notice of Meeting

Public Health Service Act (42 U.S.C. 281(d)(4)), notice is hereby given that the National Human Genome Research Institute (NHCRI) will host a series of meetings to enable public discussion of the Institute's proposal to roorganize its

internal structure. The proposed reorganization reflects the expanding scope of NHGRI's research portfolio in response to the priorities detailed in the institute's new strategic plan for genomics research, titled "Charting a Course for Genomic Medicine from Base Pairs to Bedside." (Green, E.D., Guyer, M.S. Nature, (470) 204–213.2011.)

The first public meeting will be a webinar and teleconference on January 18, 2012. The second public meeting will be on February 13, 2012, during the open seeston of the 64th meeting of the National Advisory Council for Human Cenome Research. Background materials on the proposed reorganization and logistical information regarding the two public meetings are available at genome.gov/reorg. Additional information and updated details on these public meetings will be added to this Web site as the dates approach.

Organizing Institute: National Human Genome Research Institute. Dates and Times: January 18, 2012, at 2:30

February 13, 2012, at 1 p.m. EST.

Agenda: The agenda for each meeting will include presentations and discussion about the proposed internal reorganization. Members of the public will have the opportunity to ask questions and provide comments on NHGB's proposal. Draft agendas, background materials, and instructions for joining the meetings will be made available at genome, gov/morg, ledividuals wishing to submit written questions or comments should send them via

email to NHGRIcomusentoSmih.gov. Constact Pursos: Laura Lyman Rodriguez, Ph.D., Office of Policy, Communications, and Education, National Human Genome Rossarch Institute, National Institutes of Health, 31 Center Drive, Room 4B09, Bethesda, MD 20622–2153, (201) 594–7185, NHGRIcom mentr@n ih.gov.

Dated: January 10, 2012. Eric D. Green, Director, NHGH. JPR Doc. 2012–729 Piled 1–13–12; 8:45 am] BLIMG COOK 446-01-P

DEPARTMENT OF HEALTH AND

National Institutes of Health

National Institute on Alcohol Abuse and Alcoholism; Notice of Closed Meeting

Pursuant to section 10(d) of the Federal Advisory Committee Act, as amended (5 U.S.C. App.), notice is hereby given of the following meeting

The meeting will be closed to the public in accordance with the provisions set forth in sections 552b(c)(4) and 552b(c)(6), Title 5 U.S.C., as amended. The grant applications and the and the discussions could disclose confidential trade socrets or commercial property such as patentable material, and personal information concerning individuals associated with the grant applications, the disclosure of which would constitute a clearly unwarranted invasion of personal privacy.

Name of Committee: National Institute on Alcohol Abuse and Alcoholism; Special Emphasis Panel, Review of the Collaborative Initiative on Fetal Alcohol Spectrum Disorders.

Date: March 28–29, 2012.
Time: 4 p.m. to 5 p.m.
Agenda: To review and evaluate
cooperative agreement applications.
Place: Legocy Hotel and Moeting Genter,
1778 Rockville Pike, Rockville, MD 20352.
Contact Prace: Beats Banss, Ph.D.,
Scientific Review Officer, National Institute
on Alcohol Abuse and Alcoholism, National
lastitutes Of Health, 5625 Fishers Lane, Rm.
2081, Rockville, MD 20852, (201) 443–6800.

biomas@mail.nih.gov.
(Catalogue of Fedoral Domestic Assistance Program Nos. 93.271, Alcohol Research
Garser Development Awards for Scientists
and Clinicians, 93.272, Alcohol National
Research Service Awards for Research
Training 93.273, Alcohol Research
Training 93.273, Alcohol Research
Training 93.273, Alcohol Research
Gassarch Geriter Grants,
93.991, Alcohol Research Center Grants,
93.791, ARA Related Biomedical Research
and Research Support Awards, National
Institutes of Health, HES

Dated: January 9, 2012. Jennifer S. Spaeth, Director, Officer of Federal Advisory Consentator Policy.

[PR Doc. 2012–733 Piled 1–13–12; 8:45 am]

DEPARTMENT OF HOMELAND SECURITY

Federal Emergency Management

[Docket ID: FEMA-2011-0042; OMB No. 1660-0083]

Agency Information Collection Activities: Proposed Collection; Comment Request, Application for Community Disaster Loan (CDL)

AGENCY: Federal Emergency Management Agency, DHS. ACTION: Notice.

SUMMARY: The Federal Emergency Management Agency, as part of its continuing effort to reduce paperwork and respondent burden, invites the general public and other Federal agencies to take this opportunity to comment on a proposed extension,

Steps to Reorganization

Public meetings:

Webinar (January 18)
NACHGR Meeting (February 13)

- Submission of reorganization package
- If approved, pursue next steps of appointing Division Directors and implementing new organizational structure

Anticipated Benefits of Reorganizing

- Organizational structure will more effectively align with the Institute's research portfolio (i.e., 'structure will match function')
- New divisions and anticipated substructures will improve succession planning of senior leadership
- New structure commensurate with Director's vision for organizational management

Additional information:

genome.gov/reorg

To provide feedback:

NHGRIcomments@nih.gov

