DIRECTOR'S REPORT

National Advisory Council for Human Genome Research

February 2015

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





Director's Report-Related Documents: February 2015



No.	Relevant Documents
1	New Video Spotlights NHGRI Programs
2	New NHGRI Executive Officer
3	New Branch Chiefs, NHGRI Division of Policy, Communications, and Education New Chief, Genomic Healthcare Branch New Chief, Education and Community Involvement Branch
4	Changing Role for Vence Bonham
5	Upcoming NHGRI Recruitment: Division of Genomic Medicine

genome.gov/DirectorsReport

Open Session Presentations

Update on the Genomic Medicine Working Group

Teri Manolio

The Alzheimer's Disease Sequencing Project

Eric Boerwinkle

Concept Clearance:

Centers of Excellence in ELSI Research

Joy Boyer

Open Session Presentations

Concept Clearances:

- Genome Sequencing Program Analysis
 Satellites
- Genome Sequencing Program Coordinating Center
- High-Quality Human and Primate Genomes as Foundational Resources
- Comparative and Evolutionary Genomics

Adam Felsenfeld

Open Session Presentations

 Biennial Report on the Inclusion of Women and Minorities in NHGRI-Supported Research

Rongling Li
Jacqueline Odgis

 Review of the Statement of Understanding between NACHGR and NHGRI

Rudy Pozzatti

Director's Report Outline

- I. General NHGRI Updates
- II. General NIH Updates
- **III. General Genomics Updates**
- IV. NHGRI Extramural Research Program
- V. NIH Common Fund/Trans-NIH
- VI. NHGRI Division of Policy,

 Communications, and Education
- VII. NHGRI Intramural Research Program

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New Video Spotlights NHGRI Programs





New NHGRI Executive Officer



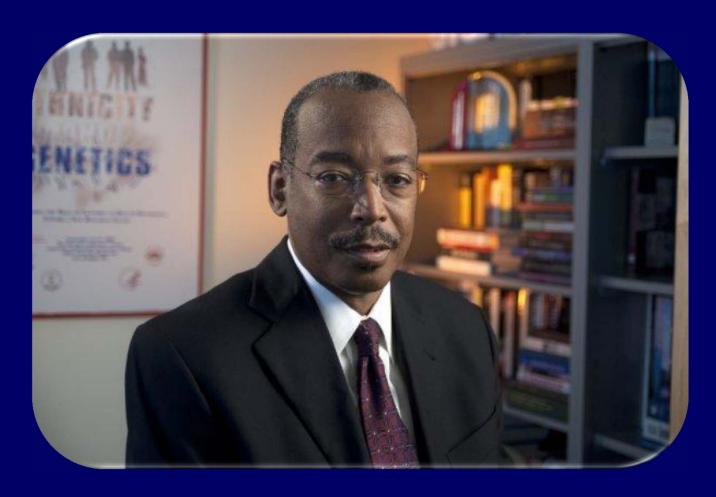
Ellen Rolfes, M.A.

New Chief, Genomic Healthcare Branch



Bob Wildin, M.D.

Changing Role for Vence Bonham



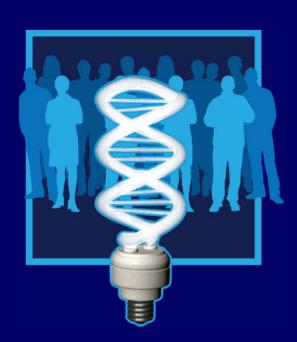
Vence Bonham, Jr., J.D.

New Chief, Education and Community Involvement Branch



Carla Easter, Ph.D.

Upcoming NHGRI Recruitment



Chief, Communications and Public Liaison Branch

Division of Policy, Communications, and Education

Contact: Dr. Laura Lyman Rodriguez laura.rodriguez@nih.gov or 301-594-7185

Upcoming NHGRI Recruitment



Medical Officer

Division of Genomic Medicine

Vacancy opening soon: Feb. 16 – Feb. 25, 2015

Contact: Dr. Teri Manolio manolio@mail.nih.gov or 301-402-2915

NHGRI Implementation of NIH Genomic Data Sharing Policy



- January 25 is effective date for policy
- Consistent implementation across NHGRI portfolio
- Coordinating with NIH-wide implementation

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President Obama Visits NIH







Secretary Burwell Visits NIH



President Obama: A Long-Standing Interest in Genomics



109TH CONGRESS 2D SESSION S. 3822

To improve access to and appropriate utilization of valid, reliable and accurate molecular genetic tests by all populations thus helping to secure the promise of personalized medicine for all Americans.

IN THE SENATE OF THE UNITED STATES

AUGUST 3, 2006

Mr. Obama introduced the following bill; which was read twice and referred to the Committee on Finance

A BILL

- To improve access to and appropriate utilization of valid, reliable and accurate molecular genetic tests by all populations thus helping to secure the promise of personalized medicine for all Americans.
- 1 Be it enacted by the Senate and House of Representa-
- 2 tives of the United States of America in Congress assembled,
- 3 SECTION 1. SHORT TITLE.
- 4 This Act may be cited as the "Genomics and Person-
- 5 alized Medicine Act of 2006".

Senator Obama, 2006

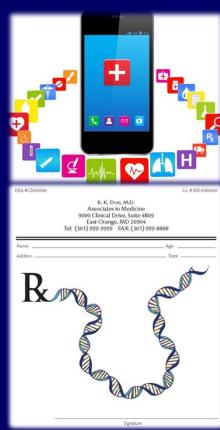


A broader context for 'individualizing' medical care to advance human health

Precision Medicine

 Today: most medical care based on expected response of the <u>average</u> patient

 Tomorrow: medical care based on individual in genomic, environmental, and lifestyle differences that enable more precise ways to prevent and treat disease



How do we get from today to tomorrow?

President Obama's State of the Union Address: January 20, 2015







"And that's why the budget I send this Congress on Monday will include a new Precision Medicine Initiative that brings America closer to curing diseases like cancer and diabetes, and gives all of us access, potentially, to the personalized information that we need to keep ourselves and our families healthier."

> President Barack Obama January 30, 2015





The NEW ENGLAND JOURNAL of MEDICINE

January 30, 2015

Perspective

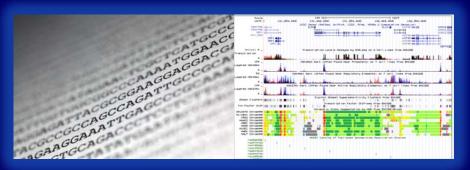
A New Initiative on Precision Medicine

Francis S. Collins, M.D., Ph.D., and Harold Varmus, M.D.

66 Tonight, I'm launching a new Precision Medicine Initiative to bring us closer to curing diseases like cancer and diabetes — and to give all of us access to the personalized information we need to keep ourselves and our families healthier."

- President Barack Obama, State of the Union Address, January 20, 2015

The proposed initiative has two main components: a nearterm focus on cancers and a longer-term aim to generate knowledge applicable to the whole range of health and disease. Both components are now within our reach because of advances in basic research, including molecular biology, genomics, and bioinformatics. Furthermore, the initiative



Genomics

EHRs



Technologies

Data Science



Patient Partnerships

Precision Medicine Initiative: The Vision

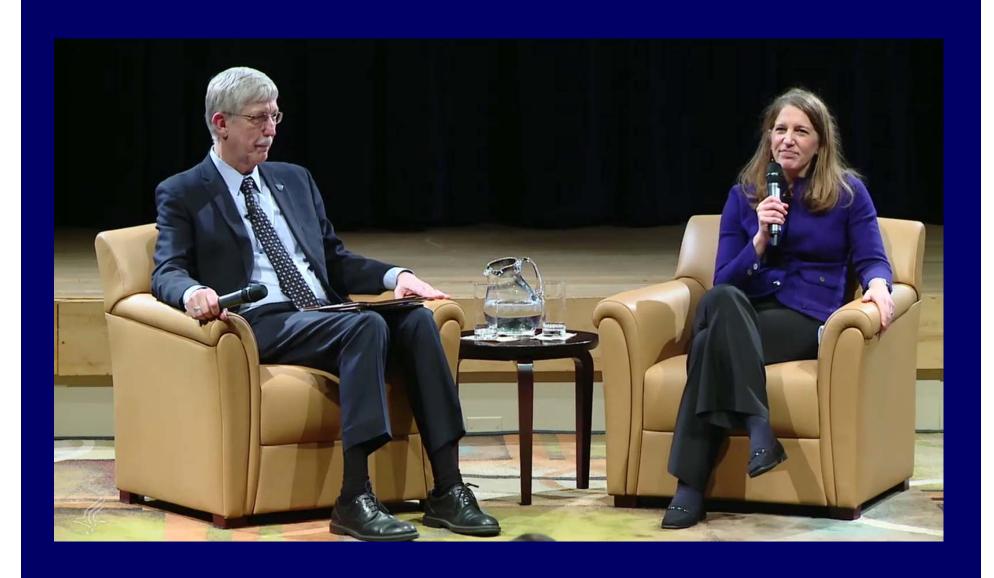
- NEAR TERM: Cancer as a Model of Precision Medicine
 Leading edge of precision medicine, yet more to learn
 Ramp up current efforts to include more cancer types
- LONGER TERM: Expanding the Model to Other Diseases
 Create national research cohort of >1 million volunteers
 Generate knowledge base for precision medicine
- POLICY CHANGES: Remove Barriers to Clinical Implementation
 Update federal rules protecting research participants
 Advance FDA oversight of precision medicine products



Precision Medicine Initiative: Proposed Fiscal Year 2016 Funding

Agency	\$ Million
National Institutes of Health	\$200
Food and Drug Administration	\$10
Office of the National Coordinator for Health Information Technology	\$5
TOTAL	\$215

Secretary Burwell Speaks Candidly at NIH



Obama seeks \$215 million for personalized medicine effort

Washington Post

Obama to Unveil Research Initiative to Develop
Tailored Medical Treatments

New York Times

Obama Announces \$215 Million Precision-Medicine Genetic Plan

Wall Street
Journal

Obama's \$215 Million DNA Sequencing Project Is A Great Idea

Forbes

A Path for Precision Medicine

New York Times

Obama Seeks Millions for 'Precision Medicine'

NBC News

White House fleshes out Obama's \$215 million plan for precision medicine

Science

Obama Enumerates Precision Medicine Initiative

The President requests \$215 million to launch his push for personalized clinical care.

The Scientist

Obama to seek \$215 million for precision-medicine plan

Details emerge as White House prepares to release budget request to Congress.

Nature

U.S. to Develop DNA Study of One Million People

An Obama initiative seeks to channel a torrent of gene information into treatments for cancer, other diseases.

MIT Tech Review



Senator Bill Cassidy (R-LA)

"This is an incredible area of promise," said Senator Bill Cassidy, Republican of Louisiana and a gastroenterologist. "There will be bipartisan support."

Obama to Request Research Funding for Treatments Tailored to Patients' DNA

By ROBERT PEAR JAN. 24, 2015

Precision Medicine Initiative: The Vision

- NEAR TERM: Cancer as a Model of Precision Medicine
 Leading edge of precision medicine, yet more to learn
 Ramp up current efforts to include more cancer types
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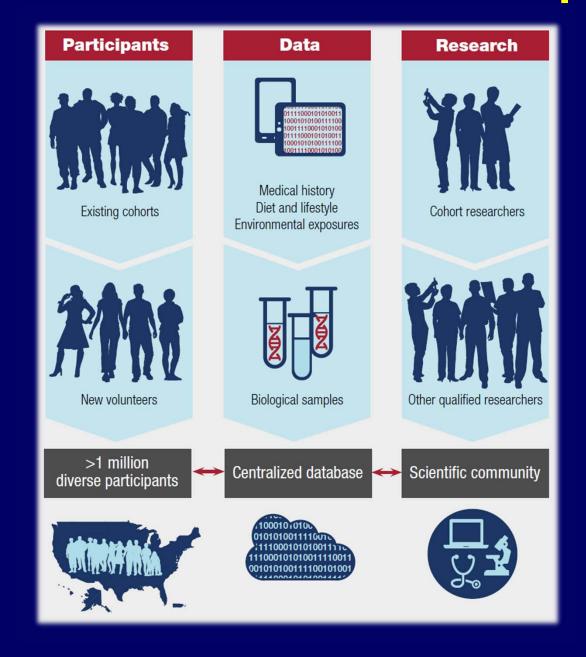


National Research Cohort



- >1 million U.S. volunteers
 - Numerous existing cohorts (many funded by NIH)
 New volunteers
- Participants to share genomic data, lifestyle information, biological samples – all linked to their EHRs
- Provide scientists with a ready platform for myriad new studies to propel understanding of health and disease
- Forge new model for 'doing science' that emphasizes engaged participants and open, responsible data sharing with strong privacy protections

National Research Cohort: Components



Building a Large U.S. Cohort for Precision Medicine Research





















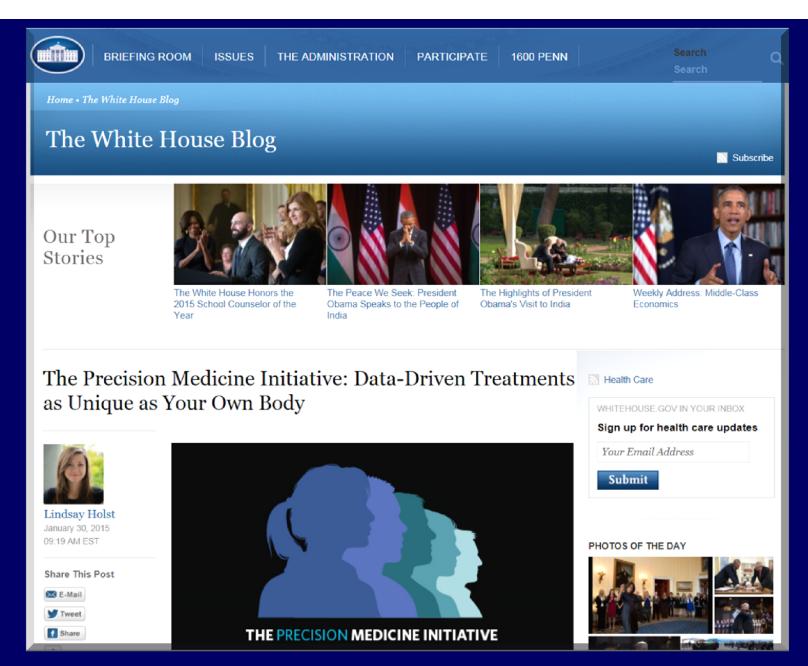
- NIH workshop to be held February 11-12, 2015
- Representatives from a wide variety of fields
- Major areas of discussion:

Cohort identification and participant recruitment

Participant engagement, data privacy, and novel ways of returning information to participants

Data collection, including mobile technologies

Informatics and electronic health records



www.whitehouse.gov/precisionmedicine



www.nih.gov/precisionmedicine

New U.S. Surgeon General





Vivek Murthy, M.D.

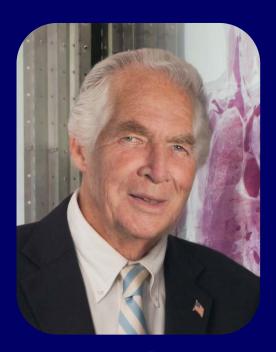
U.S. FDA Commissioner to Step Down



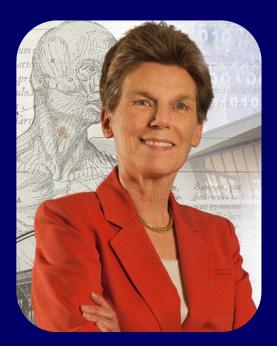


Margaret Hamburg, M.D.

Retirement of Donald Lindberg



Donald Lindberg, M.D.



Betsy Humphreys, M.L.S.



National Library of Medicine Working Group



Co-Chairs: Eric Green &

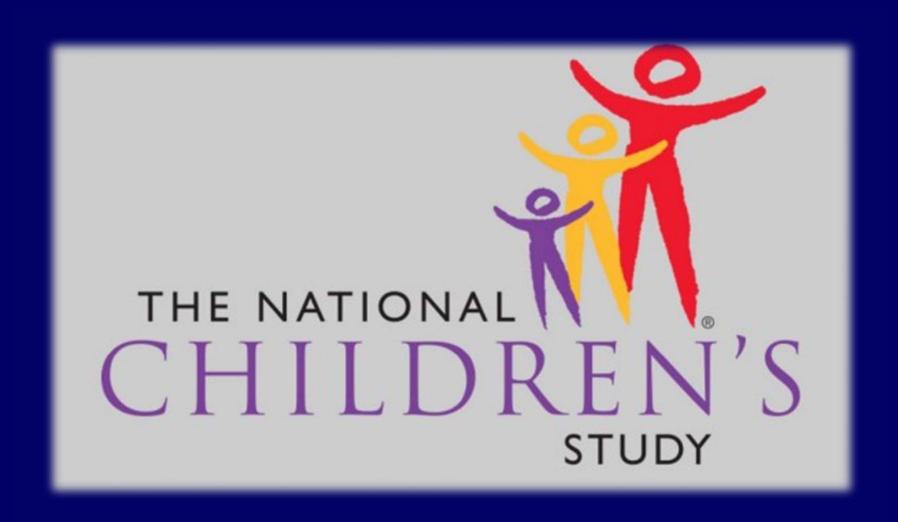
Harlan Krumholz

Charge:

- Review mission, organization, and programmatic priorities
- Articulate a strategic vision for NLM

Report Due Date:
June 2015 ACD Meeting

Cessation of the National Children's Study



Renaming of an NIH Center: NCCIH



New Associate Director for Science Policy, NIH



Carrie Wolinetz, Ph.D.



New Deputy Director, National Institute of General Medical Sciences





Judith Greenberg, Ph.D.

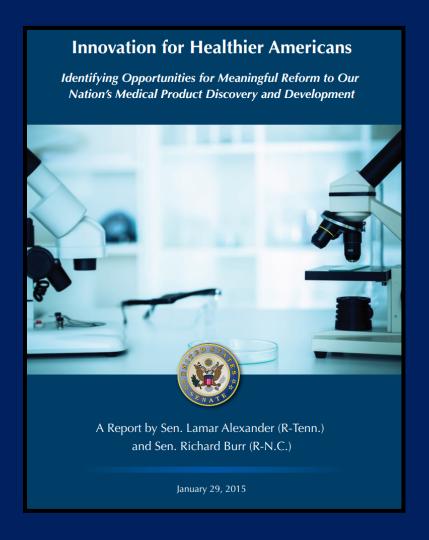
21st Century Cures Act







Health, Education, Labor, and Pensions (HELP) Committee Report



Annual Appropriations

	Fiscal Year 2014	Fiscal Year 2015	Fiscal Year 2016 President's Budget
NIH	\$30.2 B	\$30.3 B	\$31.1 B
NHGRI	\$498 M	\$499 M (+0.3%)	\$515 M (+3.2%)

- 'CRomnibus' passed, establishing NHGRI Fiscal Year (FY) 2015 funding
- On February 2, President's Fiscal Year 2016 budget sent to Congress

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Mourning the Loss of Mary Lyon





National Medal of Science and National Medal of Technology and Innovation



Bruce Alberts, Ph.D.



Douglas Lowy, M.D. and John Schiller, Ph.D.





Lasker~Koshland Special Achievement Award in Medical Science





Mary-Claire King, Ph.D.

Awards at 2014 ASHG Annual Meeting



David Valle, M.D.



Gonçalo Abecasis, D.Phil.



Mark Daly, Ph.D.



2015 Breakthrough Prize in Life Sciences





Elected to the Institute of Medicine

Goncalo Abecasis
Todd Golub
Julie Johnson
Harry Orr
Joe Takahashi



Elected to AAAS

Nancy Allbritton Russ Altman **Steven Benner** Lon Cardon A. Chakravarti Ronald DePinho **Geoffrey Duyk** Irene Eckstrand Sean Eddy

Jeffrey Friedman Dan Graur Chuan He Trey Ideker **Christine Keating Bruce Korf David Landsman David Ledbetter Brendan Lee**

Joseph Loscalzo
Karen Mohlke
Jeffrey Murray
David Nelson
Michael Snyder
William Talbot



New Director, Woods Hole Marine Biological Laboratory





Huntington Willard, Ph.D.

New Executive VP of Global Research and Chief Scientific Officer, Vertex





David Altshuler, M.D., Ph.D.

Opening of the Jackson Laboratory of Genomic Medicine





The Scientist's Top Ten Innovations 2014





DRAGEN Bio-IT Processor • Edico Genome



MiSeqDx • Illumina



HiSeq X Ten • Illumina

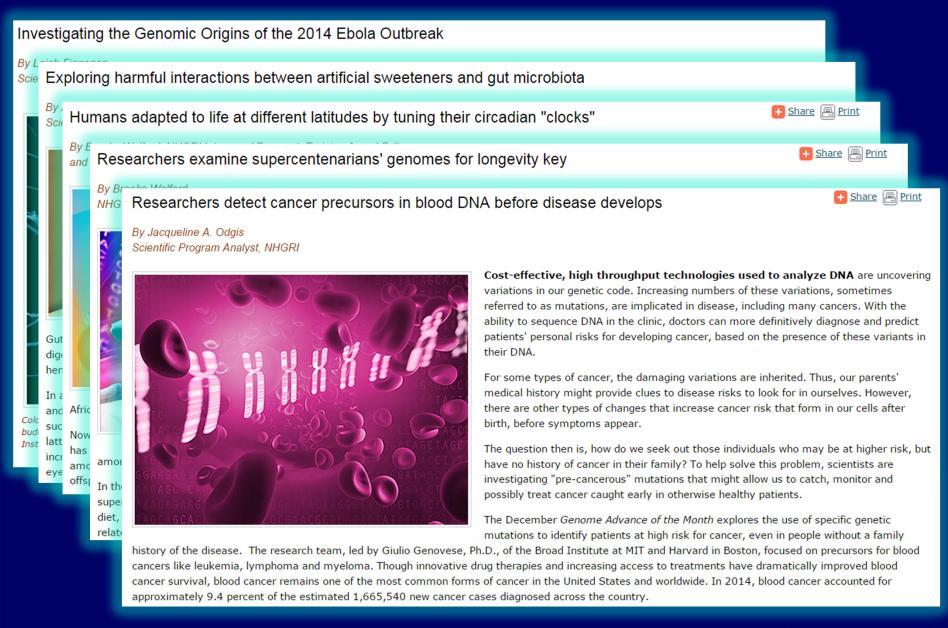


IrysChip V2 • BioNano Genomics



RainDrop Digital PCR System
• RainDance Technologies

NHGRI Genome Advance of the Month





Genomics In The News...



Forbes

The (Unmet) Potential Value of Cancer Genome Testing

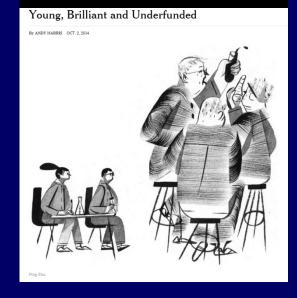
THE WALL STREET JOURNAL.

Genome Sequencing in Babies to Begin as Part of Study



Sequencing in newborns can prevent disease and lead to scientific breakthroughs later in life. WSJ's Amy Dockser Marcus discusses with Tanya Rivero. Photo: Children's Mercy Kansas City

The New York Times



genomeweb **Foundation Medicine Reimbursement Progress** Marked by Google

TheScient EXPLORING LIFE, INSPIRING INNOVATION

Why, Oh Y?
A toothpick and a bit of chance shaped David Page's career, which he has dedicated to understanding the mammalian Y chromosome and fetal germ cell development.



absolutely at random from what was then the first library of the human genome, the Maniatis lambda phage library." Page says, "I was literally pickingwith a toothpick—lambda phage plaques that contained 15-kilobase segments of the human genome. And it turns out that one of my first toothpickings was of a lambda phage clone that contained a segment of DNA that derived from the human X and Y chromosomes." Page has now spent more than three decades researching the chromosome, defending it against hypotheses that it was slowly disappearing, and demonstrating its role both within and now outside the reproductive tract. "[For] every experiment that we've done since, I can trace an unbroken line back to that

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Genome Sequencing Program

Two RFAs released in mid-December 2014:

Centers for Common Disease Genomics (UM1): RFA-HG-15-001

Centers for Mendelian Genomics (UM1): RFA- HG-15-002

Applicant information webinar on February 18

Submission deadline for both RFAs is April 7

Large-Scale Genome Sequencing and Analysis Centers

54 new papers in most recent quarter

Microbiome

Comparative Genomics

Inherited Disease

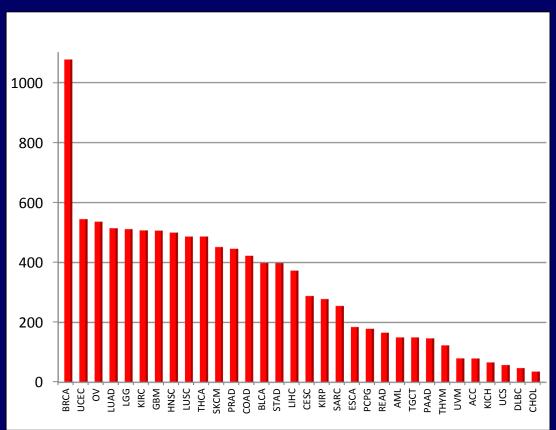
Cancer

Technology Development



TCGA Exome Sequencing Progress

TCGA Tumor Publication Status



Cancer Type	Status	Year
Glioblastoma Multiforme 1	Nature	2008
Ovarian Carcinoma	Nature	2010
Breast Cancer	Nature	2012
Colorectal Adenocarcinoma	Nature	2012
Lung Squamous Cell Ca	Nature	2012
Acute Myeloid Leukemia	NEJM	2013
Uterine Corpus Endometrial Ca	Nature	2013
Kidney Renal Clear Cell Ca	Nature	2013
Glioblastoma Multiforme 2	Cell	2013
Bladder Urothelial Carcinoma	Nature	2014
Lung Adenocarcinoma	Nature	2014
Stomach Adenocarcinoma	Nature	2014
Chromophobe Renal Cell Ca	Cancer Cell	2014
Papillary Thyroid Carcinoma	Cell	2014
Head and Neck Squamous Ca	Nature	2015
Brain Lower Grade Glioma	Submitted	
Skin Cutaneous Melanoma	Submitted	

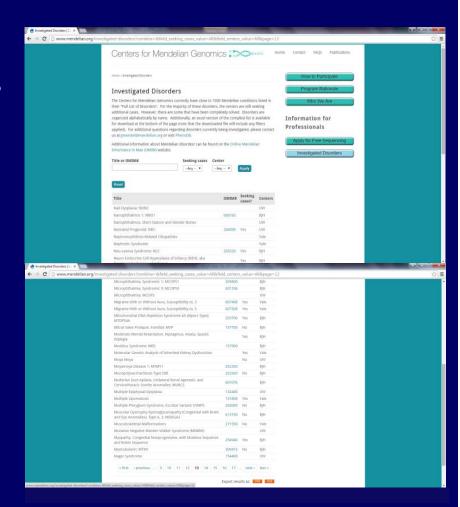
Finding the genes underlying human Mendelian conditions

Discovery

- Discovery of over 600 causal genes for Mendelian conditions
- Over 125 publications

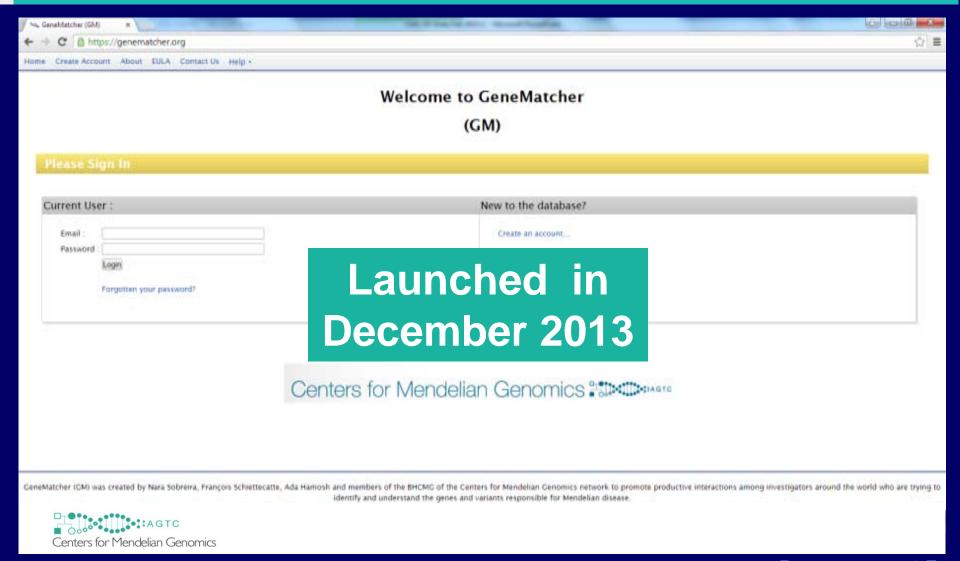
Public Project List

- Downloadable project list
- Annotation OMIM #, project site, interested in additional cases or not
- About 1000 Mendelian disorders currently listed



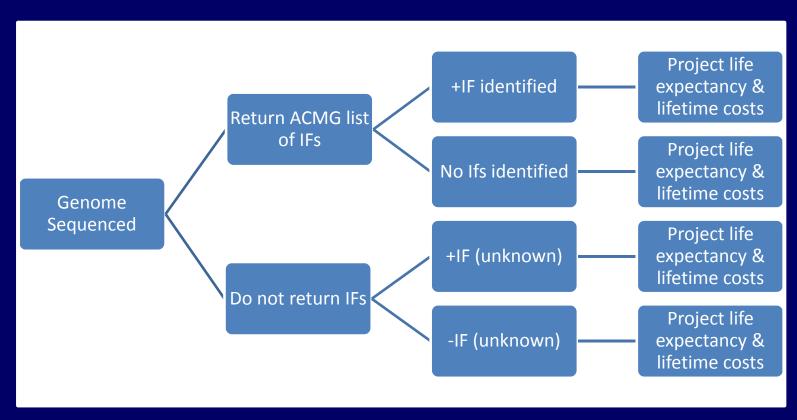
Centers for Mendelian Genomics :: Illi III III III

Finding the genes underlying human Mendelian conditions



Clinical Sequencing Exploratory Research (CSER) Program

Enrolled 3,058 adults, 707 children;
 2,661 germline, 510 tumor sequences



Bennette et al., Genetics in Medicine, 2014

Clinical Sequencing Exploratory Research (CSER) Program

Examples of recent impact:

- >130 publications, 9 working group publications
- >200 presentations/posters



ASBH 16th Annual Meeting

October 16-19, 2014

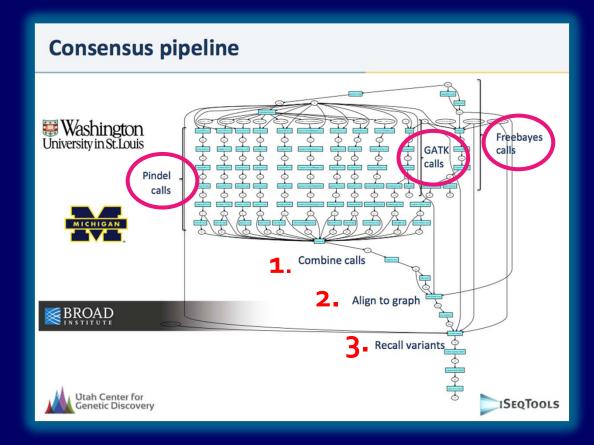
Hilton San Diego Bayfront Hotel • San Diego, CA



Genome Sequencing Informatics Tools



 iSeqTools provides GKNO pipelines that integrate results from multiple variant or mutation callers



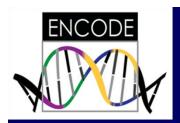
- Uses graph realignment to make best call
- IOBIO 'apps' for rapid, easy, visual analysis of data

DNA Sequencing Technology Development

Grantee meeting and public meeting in May 2015

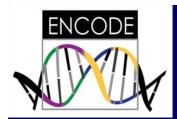


Sequencing Technology Grantees



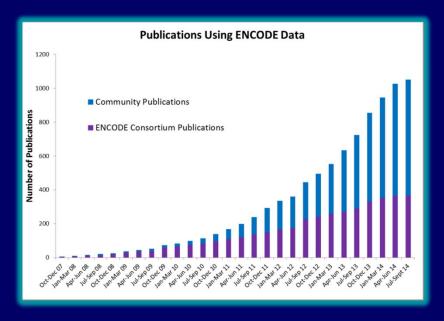
ENCyclopedia Of DNA Elements (ENCODE) Project

- Planning workshop: "From Genome Function to Biomedical Insight: ENCODE and Beyond"
 March 10-11, 2015; NIH Campus
- Annual ENCODE Consortium Meeting
 March 14-17, 2015; Cold Spring Harbor Laboratory
- ENCODE User's Meeting with hands-on workshops
 June 29-July 1, 2015; Bolger Center, Potomac, MD



ENCyclopedia Of DNA Elements (ENCODE) Project

■ ENCODE Publications →

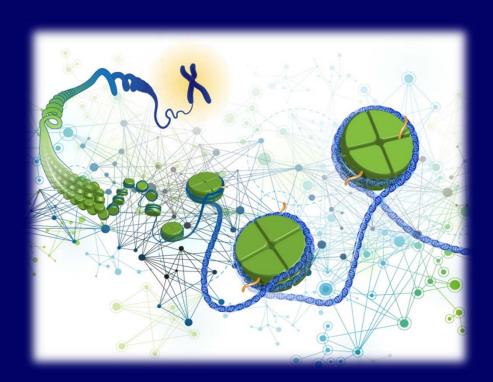


 Mouse ENCODE integrative and companion papers published in late 2014



Genomics of Gene Regulation Project

- Goal: To learn how to derive predictive gene regulatory networks starting from genomic data
- Five awards issued in January 2015
- Biological systems include immune system, skin, and nuclear hormone receptor response



Centers of Excellence in Genomic Science (CEGS) Program

Two new CEGS awards:

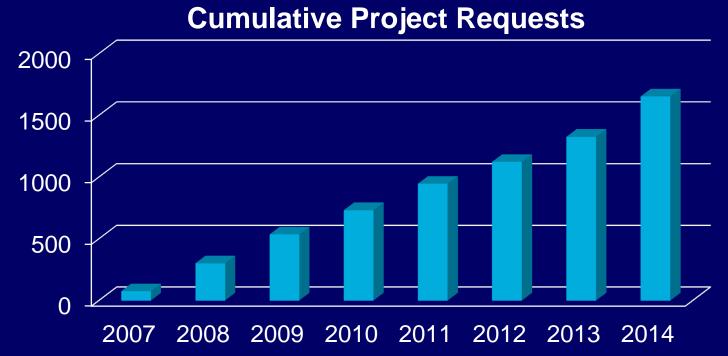
- Stanford University, Center for Personal Dynamic Regulomes
- Harvard Medical School, Neuropsychiatric Genome-Scale and RDoC Individualized Domains (N-GRID)



GAIN Data Access Committee



- GAIN DAC retired in late 2014 and all GAIN datasets transferred to relevant NIH DACs
- Nearly 1,700 project requests from 878 investigators over the course of 8 years

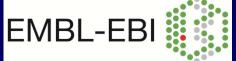


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GWAS Catalog

■ 15,000th variant





■ 2,100th paper





Beginning early March 2015, the GWAS Catalog infrastructure will be migrating to the European Bioinformatics Institute (EMBL-EBI). Content prior to this time will continue to be available on this NHGRI web page, but additional content updates will be limited. We will provide updates here as they are available; interested users can sign up for email updates at gwas-announce@ebi.ac.uk.

Current uses of and future directions for the Genome-Wide Association Studies Catalog

On Thursday, July 18th, 2013, the Division of Genomic Medicine held a webinar to highlight current uses and explore priorities and future directions for the GWAS catalog. See archived video and presentations.

The NHGRI GWAS Catalog, a curated resource of SNP-trait associations

Click here to read our recent article from the Nucleic Acids Research Database Issue.





AMIA 2014 Distinguished Paper & Homer R. Warner Awards

Development and validation of an electronic phenotyping algorithm for chronic kidney disease

Girish N. Nadkarni, Omri Gottesman, James G. Linneman, Herbert Chase, Richard L. Berg, Samira Farouk, Rajiv Nadukuru, Vaneet Lotay, Steve Ellis, George Hripcsak, Peggy Peissig, Chunhua Weng, Erwin P. Bottinger

SOEMPI: A Secure Open Enterprise Master Patient Index Software Toolkit for Private Record Linkage

Csaba Toth, Elizabeth A. Durham, Murat Kantarcioglu, Yuan Xue, Bradley Malin



- NIMH suicide and PTSD measures released in December 2014
- New NHLBI funding for sickle cell disease
- PhenX protocols released in REDCap

REDCap PhenX Demo	
Actions: Download PDF of instrument(s) Ethnicity	Assign record to a Data Access Group? selec
Record ID	100001
Do you consider yourself Hispanic/Latino? [Where did your ancestors come from?]	☐ Yes☐ No☐ Refused☐ Don't Know
Please give me the number of the group that represents your Hispanic origin or ancestry. Please select 1 or more of these categories.	 □ Puerto Rican □ Dominican Mexican/Mexicano □ Mexican American □ Chicano □ Cuban □ Cuban American □



PAGE Multi-Ethnic Genotyping (MEGA) Array

PAGE-related Traits

Multiethnic Exomic Variants

Functional Variants

GWAS Scaffold

African Diaspora Power Chip

Human Core Array

Human Exome Array

Custom Content



■ 36,000 Multiethnic Exomes

ClinVar, OMIM

1000 Genomes Project

Existing

ClinGen: Sharing Data. Building Knowledge. Improving Care.

- Standardizing clinical assessment of genomic variants and their deposition into ClinVar
- GenomeConnect: ClinGen's Patient Portal

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ClinGen-DECIPHER Public
 Meeting in May 2015



Newborn Sequencing In Genomic Medicine and Public HealTh (NSIGHT)



Genomics and Society Working Group

- Next in-person meeting in April 2015
- Outgoing members: Tim Caulfield, Jeff Long, Andrea
 Patenaude, and David Williams
- New members: Chanita Hughes-Halbert, Barbara Bernhardt, and David Veenstra
- Change in leadership:



Pamela Sankar, Ph.D. University of Pennsylvania



Lisa Parker, Ph.D.
University of Pittsburgh
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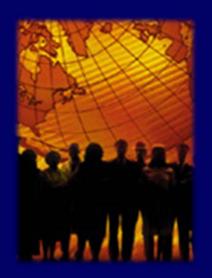
ELSI Research Program

- New ELSI NRSA Institutional Training Grant (T32) mechanism
- Centers of Excellence in ELSI Research (CEER):

First Regional CEER Networking Meeting February 23-24, 2015

Annual CEER Investigator Meeting March 16-18, 2015

New CEER RFA Concept Clearance



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Human Microbiome Project (HMP)

- HMP Phase 2: 'integrative HMP' (iHMP)
- iHMP marker paper (open access)
- iHMP Data Coordination Center
- 2nd iHMP Consortium Meeting

June 2015

Bethesda

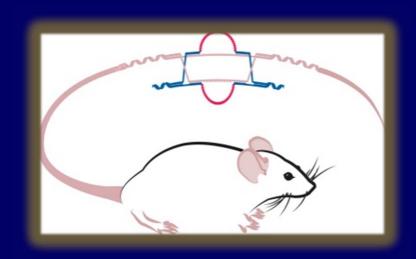


 5th International Human Microbiome Consortium Congress

March 2015

Luxembourg

Knockout Mouse Phenotyping Project (KOMP2)





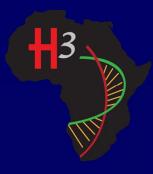
- International Mouse Phenotyping Consortium annual meeting in November
- Planning for renewal of KOMP2 program
- Sexual dimorphism evident in mutant phenotypes in the KOMP2 database



- Six investigator-initiated LINCS Data and Signature Generation Centers
- One NIH-initiated BD2K- LINCS Data Coordination and Integration Center
- First trans-LINCS meeting in October
- Data/tools release milestones for all centers developed
- Collaborative data science research opportunity announcement published

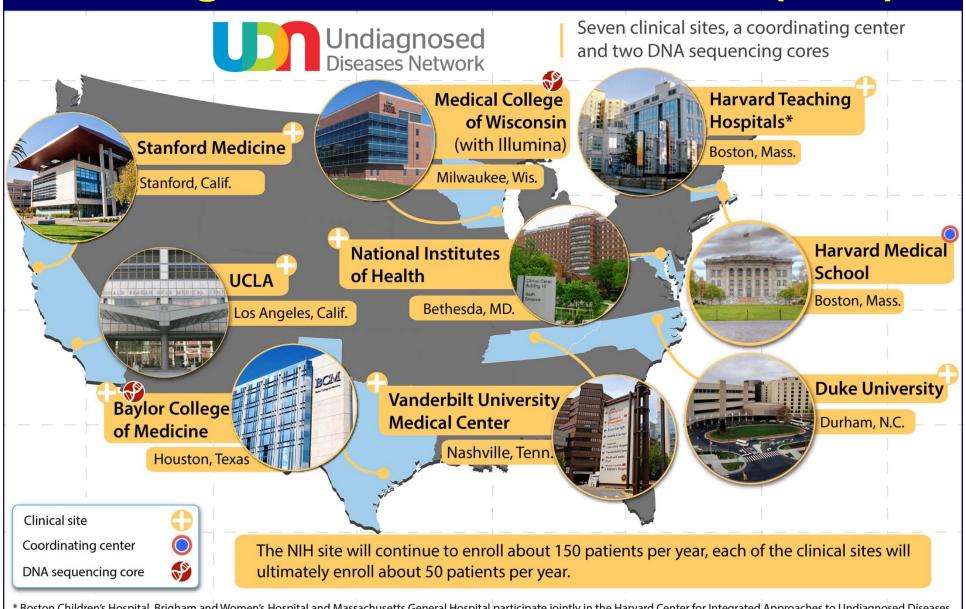
H3Africa





- 5th Consortium Meeting in November (Tanzania)
 Grant Writing Workshop
 Study Coordinators Session
 Sickle Cell Disease Workshop
- Supplement for whole-genome sequencing
- Progress on developing custom genotyping chip
- Supplement for research ethics training
- 6th Consortium Meeting in May (Zambia)

Undiagnosed Diseases Network (UDN)



* Boston Children's Hospital, Brigham and Women's Hospital and Massachusetts General Hospital participate jointly in the Harvard Center for Integrated Approaches to Undiagnosed Diseases

Undiagnosed Diseases Network (UDN)

PI Name	Gene	Model	Patient Phenotype
Worley	FRMPD4	mouse, human male cohort	developmental delay and regression, seizures
Lin	HK1, SUSD4, CCDC89, and BAI2	zebrafish	neurodegeneration and spastic tetraplegia like syndrome
Antonellis	GARS, AARS, and DARS	yeast, cell culture, zebrafish	recessive disease phenotypes
Graham	ATP1A3, SYNE1, SNAP29, ARHGAP22, KIF4B and XRN1	patient primary fibroblasts, drosophila	neurological and/or metabolic phenotypes
Chen	50 genes	zebrafish	many
Slusarski	15 genes	zebrafish	epilepsy







Big Data to Knowledge (BD2K) Initiative



Funded BD2K Elements

- Centers of Excellence, Data Discovery Index Coordination Consortium, Short Courses, Open Educational Resources, Mentored Career Development
- Joint kickoff meeting in November

Recently Reviewed Elements

- Targeted Software Development
- Institutional Pre-Doctoral Training Programs

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Future BD2K Efforts



- The Commons
- NIH Standards Information Resource
- Database Sustainability
- Training Coordination Center
- Diversity

Genome Privacy Challenge



- iDASH center and collaborators develop solutions to problems in data sharing and privacy protection
- 2nd iDASH privacy challenge in March 2015

Challenge 1: Homomorphic encryption

Challenge 2: Secure multiparty computing for secure genomic data analysis across institutions

Director's Report Outline

- I. General NHGRI Updates
- II. General NIH Updates
- III. General Genomics Updates
- IV. NHGRI Extramural Research Program
- V. NIH Common Fund/Trans-NIH
- VI. NHGRI Division of Policy,

 Communications, and Education
- VII. NHGRI Intramural Research Program

Inter-Society Coordinating Committee for Practitioner Education in Genomics

American College of Medical Genetics and Genomics

COMMENTARY

Genetics inMedicine

The growing role of professional societies in educating clinicians in genomics

Teri A. Manolio, MD, PhD¹ and Michael F. Murray, MD²; for the Inter-Society Coordinating Committee for Practitioner Education in Genomics

- Physicians and Dentists
- Pharmacists and Nurses
- Genetic Counselors





- In-person meeting in November 2014
- New working groups:

 Physician-patient Communications
 Innovative Approaches to Education

Physician Resources Now on G2C2

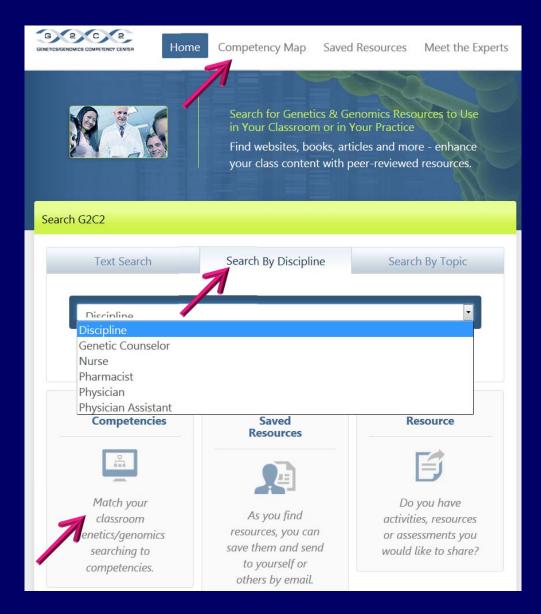












Newborn Screening Saves Lives Reauthorization Act of 2014



Clinical Trials Policy Update



ClinicalTrials.gov

A service of the U.S. National Institutes of Health

ClinicalTrials.gov is a registry and results database of publicly and privately supported clinical studies of human participants conducted around the world. Learn more <u>about</u> clinical studies and about this site, including relevant history, policies, and laws.

Now Available for Public Comment: Notice of Proposed Rulemaking (NPRM) for FDAAA 801 and NIH Draft Reporting Policy for NIH-Funded Trials

- New definition of "clinical trial"
- Reporting requirements
- Centralized Institutional Review Boards (IRBs)

FDA Lab-Developed Test (LDT) Guidance



Should the FDA regulate laboratory-developed diagnostic tests? -No.

In November 2014, the US Food and Drug Administration (FDA) revealed its intent to regulate thousands of medical diagnostic tests being performed in as many as 11 000 clinical laboratories throughout the United States, focusing especially on genomic medicine.1 Although the FDA is well intentioned, the current plan for regulation is unnecessary and, if carried out, could result in the closure of many laboratories, undermine innovation, and potentially limit patient choice. Moreover, the proposed regulation, if unchanged, is likely to lead to thousands of laboratory submissions to the FDA, for which its own staffing capacity is tenuous at best. If implemented, the requirements may have the unintended effect of derailing the long-awaited emergence of genomic medicine.

The last several years have seen substantial expansion in genetic testing, resulting from advances in technology that allow rapid and accurate sequencing of large fractions of an individual's DNA. Such analyses have begun to inform patient care in spheres ranging from carrier screening and the diagnosis of birth defects to individualized diagnosis and treatment of cancer.^{2,3} These developments have occurred in the span of just a few years, in large part because of the nimbleness of relatively small clinical and academic laboratories that can quickly respond to new medical findings and patient needs by rapidly and safely developing and improving laboratory-developed tests. The resulting landscape is one of vibrant competition in which laboratories that offer genetic testing now compete on the basis of quality, service, innovation, and

The FDA now proposes to regulate laboratory-developed tests as "medical devices," mandating that laboratories be treated as manufacturers that must meet formal FDA manufacturer requirements for each test developed-a costly and time-consuming process. However, this approach has little valid or even apparent justification. The FDA has failed to cite more than a few anecdotal examples of patient harm to justify its proposed actions. During a congressional hearing on the draft regulation,4 the director of the



Should the FDA regulate laboratory-developed diagnostic tests? -Yes.

In April 2014, the Centers for Disease Control and Prevention and the US Food and Drug Administration (FDA) published a warning in Morbidity and Mortality Weekly Report about a commercially available test for Lyme infection. The test returned the result "culture positive," when in fact the procedure was far more complex than a routine culture. There also were "serious concerns about false-positive results caused by laboratory contamination," leading to "the potential for misdiagnosis,"1

The questionable assay was a laboratory-developed test, meaning an "in vitro diagnostic test that is designed, manufactured, and used within a single laboratory."2 Laboratory-developed tests exist in a regulatory crevice. Because of its broad statutory authority over products "intended for use in the diagnosis of disease or other conditions, "3 the FDA considers laboratory-developed tests under its jurisdiction. Yet for many years, the FDA has taken the position that there were too few of these tests, and that they were of sufficiently low risk, to merit oversight. As a result, tests "designed, manufactured, and used within a single laboratory" are not subject to the standards for quality and validity applicable to other diagnostic tests, such as those made by medical device manufacturers.

Recently, however, the FDA has expressed concern with the proliferation of laboratory-developed tests, their marketing, and their potential to mislead physicians and patients and undermine clinical care. On July 31, 2014, the agency notified Congress that the agency would shortly release a draft guidance document containing a framework for the application of agency standards for quality, safety, and validity to laboratory-developed tests. On September 30, the agency posted the draft guidance document to its website and opened a comment period lasting until February 2, 2015.4 The agency will hold a public meeting on the topic on January 8 and 9, 2015, at the National Institutes of Health in Bethesda, Maryland.

FDA Workshop on Oversight of 'Next-Generation' Genome Sequencing





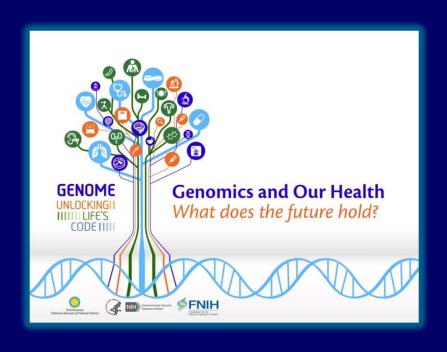
Public Workshop: Optimizing FDA's Regulatory Oversight of Next-Generation Sequencing Diagnostic Tests
February 20, 2015

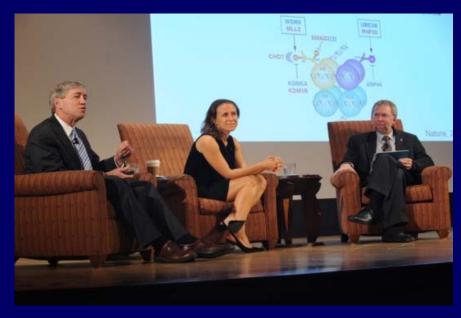
Informed Consent Resource for Genomics



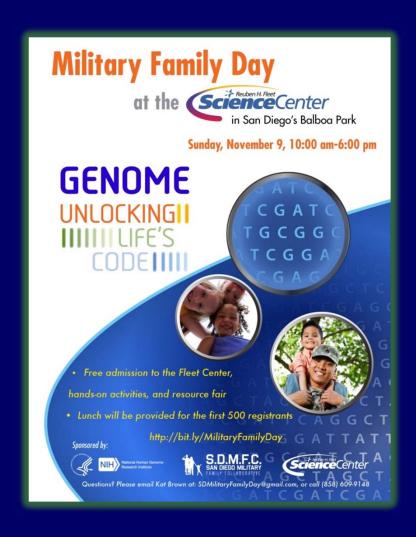
- Discussion of topics essential to genomics research
- Sample language and consent forms
- Relevant regulations and policies
- Email: informedconsent@mail.nih.gov

Genome: Unlocking Life's Code Exhibition Closing Symposium





Genome: Unlocking Life's Code Exhibition Military Family Day









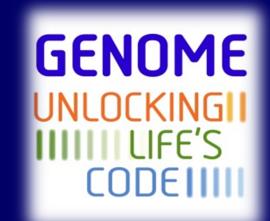


Genome: Unlocking Life's Code Exhibition Travel Schedule

2015

January 22-April 27:
The Tech Museum of Innovation
San Jose, CA

May 15-September 10
The Saint Louis Science Center
St. Louis, MO



October 2-January 3
Oregon Museum of Science and Industry
Portland, OR

2016

January 18-April 25
Discovery World Milwaukee
Milwaukee, WI

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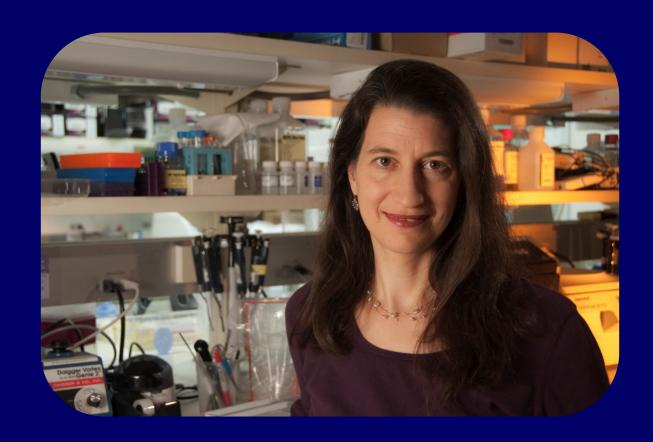
 Communications, and Education
- VII. NHGRI Intramural Research Program

Report of the NIH Intramural Research Program Working Group

National Institutes of Health **Advisory Committee to the Director** Long-Term Intramural Research Program (LT-IRP) **Planning Working Group Report** December 12, 2014 National Institutes of Health



2014 CHANEL-CERIES Research Award



Julie Segre, Ph.D.

2014 Rare Voice Award



William A. Gahl, M.D., Ph.D.

2014 NSGC Leadership Award

Natalie Weissberger Paul National Achievement Award



Barbara B. Biesecker, M.S., Ph.D., C.G.C.

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2014 South African Medical Research Council Scientific Merit Award



Charles Rotimi, Ph.D.

2014 AJHG C.W. Cotterman Award





Shurjo Sen, Ph.D.

NHGRI Intramural Research Highlights



nature International weekly journal of science

The African Genome Variation Project shapes medical genetics in Africa

nature International weekly journal of science

Biogeography and individuality shape function in the human skin metagenome



The Journal of Clinical Investigation

Vector design influences hepatic genotoxicity after adeno-associated virus gene therapy



The Genomics Landscape

A monthly update from the NHGRI Director



To receive *The Genomics Landscape*, go to list.nih.gov

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Past issues can be accessed at: genome.gov/27527308



Thanks!



Special Thanks!