

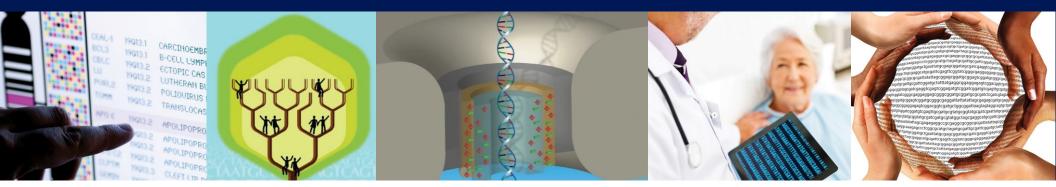
NIH National Human Genome Research Institute



# **DIRECTOR'S REPORT**

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

## September 2019





# genome.gov/DirectorsReport



## **Open Session Presentations**

 Presentation: Inter-Society Coordinating Committee for Practitioner Education in Genomics (ISCC-PEG) – Promoting Collaboration in Genomics Education
 Richard Haspel

Concept Clearances:

Genomic Community Resources Chris Wellington

ELSI Program Announcements for Unsolicited Applications Joy Boyer

# **Open Session Presentations**

Concept Clearances:

Genomic Technology Development Michael Smith

Technology Development Michael Smith

Mendelian Genomics: Solving the Unsolved Lisa Chadwick

Polygenic Risk Scores from Populations of Diverse Ancestry Lucia Hindorff

# **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 ASHG-NHGRI Fellows**



Samata Katta, Ph.D. Genetics and Public Policy Fellow



Sarah Robbins, Ph.D. Genetics Education and Engagement Fellow



# NIH-ACMG Fellowship in Genomic Medicine Program Management

- Increase pool of physicians trained in managing research and implementation programs in genomic medicine
- Up to two qualified physicians selected annually to acquire credentials and experience at NIH and other organizations
- Applications for two-year fellowship due annually December 1





Marie-Luise Brennan, M.D., Ph.D.

Hwaida Hannoush, M.D.

# Genomics and Health Disparities Lecture Series



## Minoli Perera, Pharm. D., Ph.D. Associate Professor. Department of Pharmaco

Associate Professor, Department of Pharmacology Northwestern University March 26, 2019



## Katrina Armstrong, M.D.

Chair, Department of Medicine Physician-in-Chief, Massachusetts General Hospital July 16, 2019



# Ali Gharavi, Ph.D.

Chief, Division of Nephrology Institute of Genomics Medicine, Columbia University December 5, 2019

# Genomic Medicine Working Group Lancet Series

# THE LANCET

#### Series from the Lancet journals

View all Series

### **Genomic medicine**

Published: August 5, 2019

#### **Executive Summary**

The acceleration of documenting genomic variation in disease states and health outcomes has led to an upswell of clinical insights and avenues of exploration. The exciting and maturing area of genomic medicine, encompassing the use of individuals' genetic information to guide clinical treatment, is featured in this Series. In these five, timely reviews, authors from the National Human Genome Research Institute at the National Institutes of Health (MD, USA) present an overview of efforts to implement genomic medicine, including contemporary tools and resources; the role of pharmacogenomics in drug safety and efficacy; the application of genomics to identifying rare and undiagnosed diseases; enhancing risk assessment through family health history records; and improving the collection of outcomes and evidence to better incorporate genomic techniques in clinical care.



Log in

#### **Related Content**

Mapping a route to Indigenous engagement in cancer genomic research Kimiora L Henare, Kate E Parker, Helen

# 'Genomics2020' Strategic Planning Process

- Community Town Hall at NIH
- Two ELSI Strategic Planning Webinars:

Appropriate Use & Implementation of Genomics in Diverse Communities Genomic Decision-Making: Power and Identity

- Town Hall at Massachusetts General Hospital
- Workshop: Perspectives in Comparative Genomics and Evolution





# 'Genomics2020' Strategic Planning Process

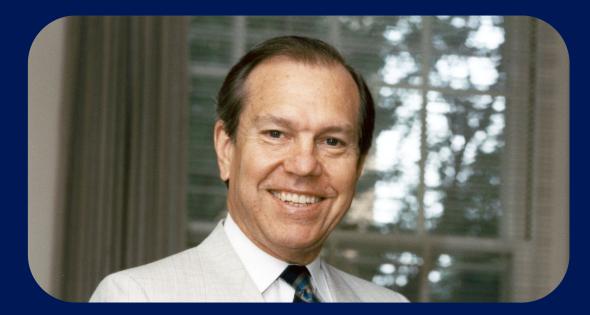
- Workshop: Genomics in Medicine & Health
- ASHG Feature: Strategic Planning Update with Eric Green, NHGRI



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# Mourning the Loss of James Wyngaarden





# Mourning the Loss of Donald Lindberg





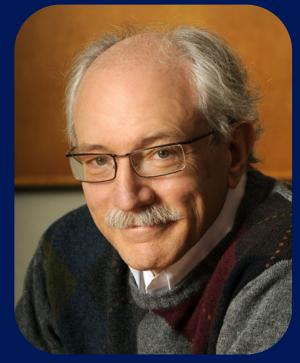
# Paul Sieving Departs as Director, National Eye Institute



# Linda Birnbaum Departs as Director, National Institute of Environmental Health Sciences



# Linda Birnbaum, Ph.D., D.A.B.T., A.T.S.



Richard Woychik, Ph.D.

# **New NIH Associate Director for Data Science**





## Susan Gregurick, Ph.D.

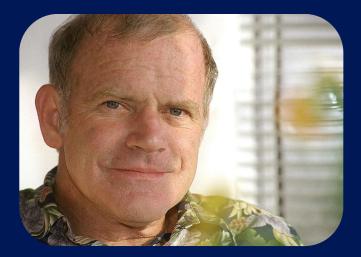
# **Fiscal Year 2020 Appropriations**

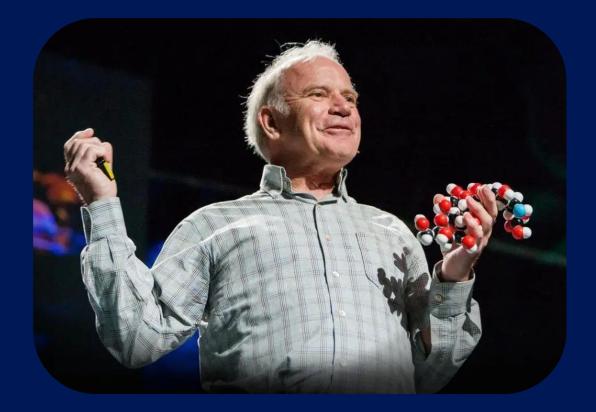
	Actual FY 2019 Labor-HHS Appropriations	Proposed FY2020 Labor-HHS Appropriations					
		House	%Δ	Senate	%Δ		
NIH	\$39.1B	\$41.1B	+5%	\$42.1	+7.7%		
NHGRI	\$575M	\$604M	+5%				

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# Mourning the Loss of Kary Mullis





# Presidential Early Career Awards for Scientists and Engineers



Neville Sanjana, Ph.D. Adam Phillippy, Ph.D. David Zhang, Ph.D.

# **Genomicist Receives British Knighthood**



Peter Donnelly, FRS FmedSci.



# **Director's Report Outline**

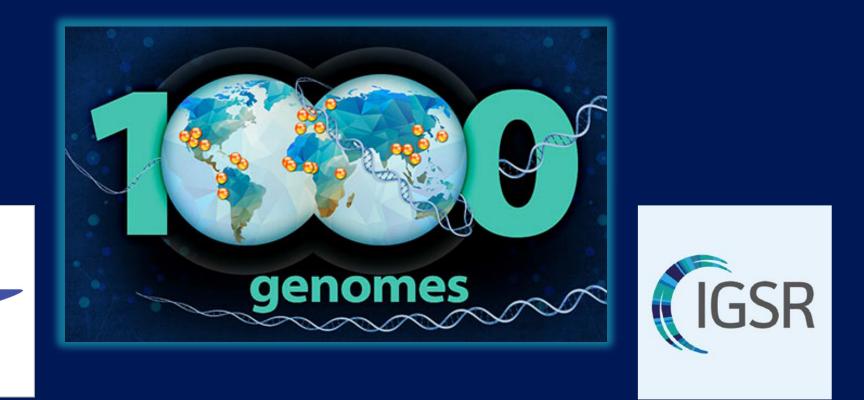
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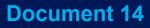
# **Genome Sequencing Program** Centers for Common Disease Genomics

	Disease Category	Sample Type	Samples Sequenced	
	Cardiovascular	Genomes	58,718	Samples
	cardiovascular	Exomes	61,799	Projected by End
	mmune-Mediated	Genomes	10,271	of Y5
		Exomes	15,135	
	Neuropsychiatric	Genomes	21,728	
	Neuropsychiatric	Exomes	28.911	
	TOTAL GENOMES		~91,000	~141,000
	TOTAL EXOMES		~105,000	~212,000
				Document

14

# **Genome Sequencing Program** 1000 Genomes High-Quality Sequences Available





# **Genome Sequencing Program** Centers for Mendelian Genomics

Tie	Tier 1		Tier 2				
Total		3,048	Tota	al		1,	595
Novel	1,213	(40%)	Νον	el	1,55	50 (9	7%)
Not Novel	1,829	(60%)	Not	Novel		45 (	3%)
Dat	a Sharing				<i>-</i>		
Matchmaker Exchange		125,0			NOX .	ClinVar	
VariantMatcher		5,7		Matchm Exchar		Clinically relevant variation	
ClinVar		3,9	986		IGE		
			V	ariantN	1atch	er	Docume

# Genome Sequencing Program Genome Sequencing Program Analysis Centers

## nature neuroscience

A Bayesian framework that integrates multi-omics data and gene networks predicts risk genes from schizophrenia GWAS data

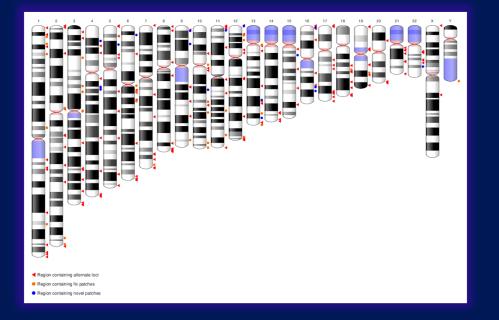
Quan Wang, Rui Chen, Feixiong Cheng, Qiang Wei, Ying Ji, Hai Yang, Xue Zhong, Ran Tao, Zhexing Wen, James S. Sutcliffe, Chunyu Liu, Edwin H. Cook, Nancy J. Cox & Bingshan Li 🏁

### nature genetics

Genome sequencing analysis identifies Epstein–Barr virus subtypes associated with high risk of nasopharyngeal carcinoma

Miao Xu, Youyuan Yao, Hui Chen, Shanshan Zhang, Su-Mei Cao, Zhe Zhang, Bing Luo, Zhiwei Liu, Zilin Li, Tong Xiang, Guiping He, Qi-Sheng Feng, Li-Zhen Chen, Xiang Guo, Wei-Hua Jia, Ming-Yuan Chen, Xiao Zhang, Shang-Hang Xie, Roujun Peng, Ellen T. Chang, Vincent Pedergnana, Lin Feng, Jin-Xin Bei, Rui-Hua Xu, Mu-Sheng Zeng, Weimin Ye, Hans-Olov Adami, Xihong Lin, Weiwei Zhai X, Yi-Xin Zeng & Jianjun Liu - Show fewer authors

# Human Genome Reference Program



# **Two centers funded:**

Human Genome Reference Center (HGRC)



# High-Quality Human Reference Genomes Center (HGRQ)



# **Technology Development Program** Advanced Genomic Technology Development Meeting



 Novel Genomic Technology Development Program Announcement PAR-HG-18-777 (R01, also linked R21 and R43/44) Next application due date: October 2, 2019

# Technology Development Program Recent Papers

# DNA Microscopy: Optics-free Spatio-genetic Imaging by a Stand-Alone Chemical Reaction

Joshua A. Weinstein,<sup>1,8,\*</sup> Aviv Regev,<sup>1,2,3,4,\*</sup> and Feng Zhang<sup>1,3,5,6,7,\*</sup>

# Increasing the accuracy of nanopore DNA sequencing using a time-varying cross membrane voltage

Matthew T. Noakes<sup>1,2</sup>, Henry Brinkerhoff<sup>1,2</sup>, Andrew H. Laszlo<sup>1</sup>, Ian M. Derrington<sup>1</sup>, Kyle W. Langford<sup>1</sup>, Jonathan W. Mount<sup>1</sup>, Jasmine L. Bowman<sup>1</sup>, Katherine S. Baker<sup>1</sup>, Kenji M. Doering<sup>1</sup>, Benjamin I. Tickman<sup>1</sup> and Jens H. Gundlach<sup>1</sup>



# **ENCyclopedia of DNA Elements (ENCODE)**





 Research Applications & Users Meeting in Seattle, WA

Accessing ENCODE's 15,000+ Datasets

Using the GENCODE reference annotation of human and mouse genes and transcripts

Neural networks and deep learning: applications to genomics and gene regulation

 Next Users Meeting planned for Summer 2020 in Barcelona, Spain

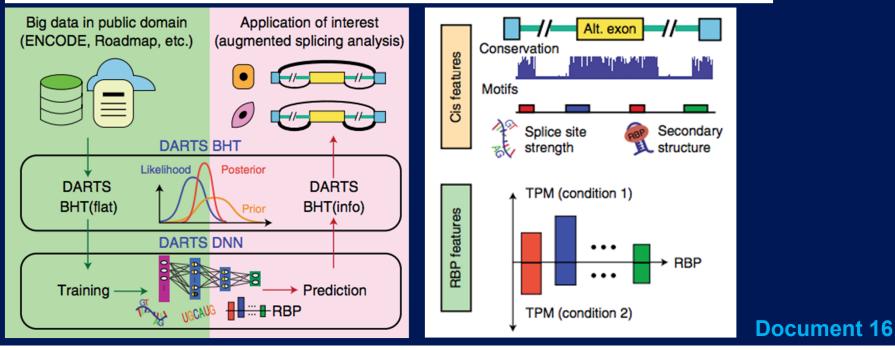




# **ENCyclopedia of DNA Elements (ENCODE)**

# Deep-learning augmented RNA-seq analysis of transcript splicing

Zijun Zhang<sup>1,8</sup>, Zhicheng Pan<sup>1,8</sup>, Yi Ying<sup>2</sup>, Zhijie Xie<sup>2</sup>, Samir Adhikari<sup>2,3</sup>, John Phillips<sup>4</sup>, Russ P. Carstens<sup>5</sup>, Douglas L. Black<sup>2</sup>, Yingnian Wu<sup>6</sup> and Yi Xing<sup>1,2,3,7\*</sup>



# **Centers of Excellence in Genomic Science (CEGS)**



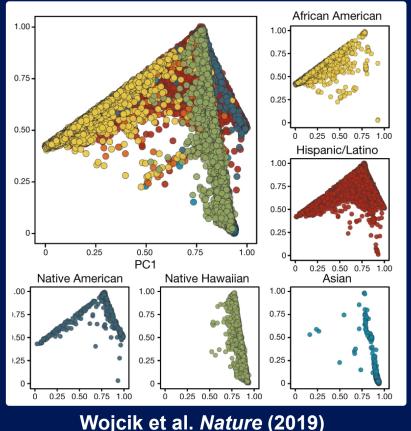


Center for Personal Dynamic Regulomes Howard Chang Stanford University Center for the Multiplexed Assessment of Phenotype Stanley Fields & Douglas Fowler University of Washington



# Population Architecture using Genomics and Epidemiology (PAGE)

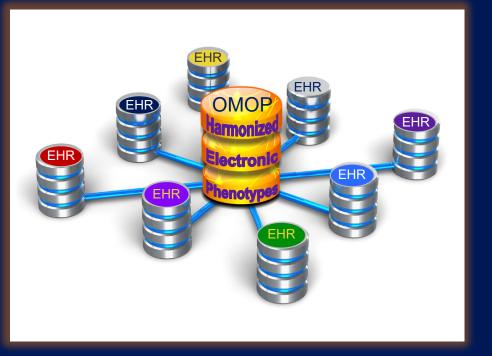
## **Principal Component Analysis**



## Added value of diversity in GWAS:

- Analysis of European ancestry (EA) height GWAS: more variation in height explained in EA vs. non-EA populations
- Adding 50K EA participants: widens gap
- Adding 50K non-EA participants: narrows gap
- Risk of exacerbating disparities in genetic knowledge

# emerge network



Contents lists available at ScienceDirect

Journal of Biomedical Informatics

journal homepage: www.elsevier.com/locate/yjbin

Facilitating phenotype transfer using a common data model

George Hripcsak<sup>a,b,\*</sup>, Ning Shang<sup>a</sup>, Peggy L. Peissig<sup>c</sup>, Luke V. Rasmussen<sup>d</sup>, Cong Liu<sup>a</sup>, Barbara Benoit<sup>e</sup>, Robert J. Carroll<sup>f</sup>, David S. Carrell<sup>g</sup>, Joshua C. Denny<sup>f,h</sup>, Ozan Dikilitas<sup>i</sup>, Vivian S. Gainer<sup>e</sup>, Kayla Marie Howell<sup>j</sup>, Jeffrey G. Klann<sup>e</sup>, Iftikhar J. Kullo<sup>i</sup>, Todd Lingren<sup>k</sup>, Frank D. Mentch<sup>l</sup>, Shawn N. Murphy<sup>e</sup>, Karthik Natarajan<sup>a,b</sup>, Jennifer A. Pacheco<sup>d</sup>, Wei-Qi Wei<sup>f</sup>, Ken Wiley<sup>m</sup>, Chunhua Weng<sup>a</sup>

- Clinical sites converted local data model to the Observational Medical Outcomes Partnership (OMOP) Common Data Model (CDM)
- Conversion helped identify errors
- Value of using the OMOP data model when generating e-phenotypes



## **Clinical Genome (ClinGen) Resource**



#### Variant Curation Panels

- Very Long-Chain Acyl-CoA Dehydrogenase Deficiency
- Hereditary Hemorrhagic Telangiectasia
- FBN1 for Marfan & Hereditary Thoracic Aortic Aneurysm and Dissection
- Coagulation Factor Deficiency
- Cerebral Creatine Deficiency Syndromes

#### **Volunteer to Curate**

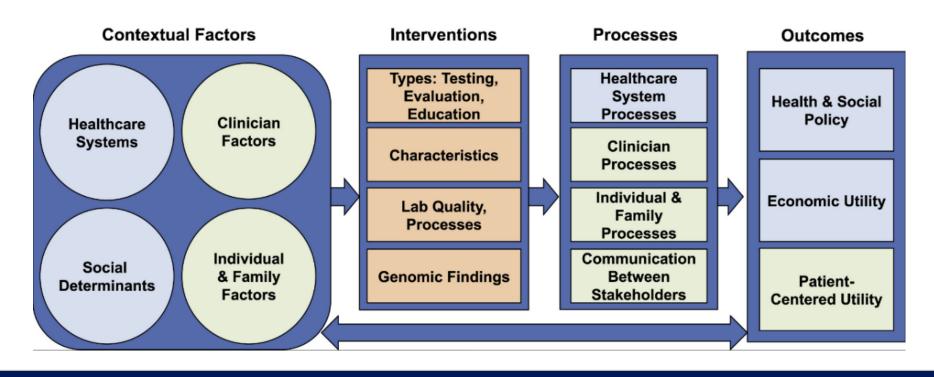


- 248 people completed the survey
- 116 volunteers completed training



#### **Clinical Sequencing Evidence-Generating Research Program**

The Genomic Medicine Integrative Research Framework: A Conceptual Framework for Conducting Genomic Medicine Research



Horowitz et al. Am J Hum Genet (2019)

#### Newborn Sequencing In Genomic medicine and public HealTh (NSIGHT)

Jennifer Puck, M.D. University of California, San Francisco

NIH

Cynthia Powell, M.D., M.S. University of North Carolina at Chapel Hill

NIH

# **NSIGHT Final Public** Session

#### June 24, 2019

YOUR HEALTH

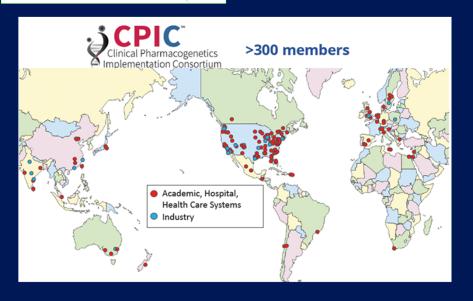
The Promises And Pitfalls Of Gene Sequencing For Newborns

July 8, 2019 · 5:00 AM ET Heard on Morning Edition

RICHARD HARRIS

Robert Green, M.D., M.P.H. Brigham and Women's Hospital Stephen Kingsmore, M.D., D.Sc. Rady Children's Institute for Genomic Medicine

#### Clinical Pharmacogenetics Implementation Consortium (CPIC®)



CPIC

#### June 2019 Meeting:

- Identify practical considerations for choosing a genetic testing platform and clinical laboratory
- Summarize the risks and benefits of returning pharmacogenetic test results
- Describe barriers to implementation of pharmacogenetics
- Define successful implementation of pharmacogenetics and associated improvements

Menu 💻

Clinical PGx Consortium Taking Aim at Non-Actionable Genes, Countering 'Inappropriate' Marketing

Jun 17, 2019 | Turna Ray

enomeweb

#### **ELSI Research Program** Center for ELSI Resource and Analysis (CERA)

Co-Pls:

Mildred Cho (Stanford University)

Sandra Lee (Columbia University)

Goals:

Resource Sharing and Coordination Data Curation and Synthesis Increase availability and visibility of ELSI resources



## ELSI Research Program 5<sup>th</sup> ELSI Congress



COLUMBIA DIVISION OF ETHICS DEPARTMENT OF MEDICAL HUMANITIES AND ETHICS June 15-17, 2020

- Hosted by Columbia Division of Ethics (New York City)
- Abstracts due in late fall

#### **Research Training and Career Development**

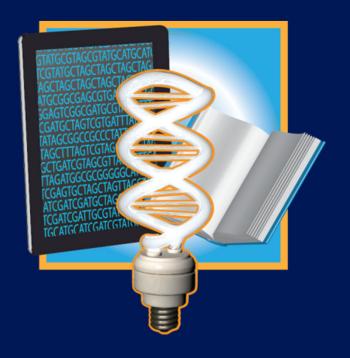
New DAP Awards:

Stanford University PI: Michael Snyder

Baylor College of Medicine Pls: Richard Gibbs & Debra Murray

New T32 Training Award:

Massachusetts General Hospital Pls: Jordan Smoller & Heidi Rehm



#### Genomic Innovator Awards 2019 Awardees

PI	Institution
Eric Gamazon	Vanderbilt University Medical Center
Stacy Gray	Beckman Research Institute/City Of Hope
Channabasavaiah Gurumurthy	University Of Nebraska Medical Center
Timothy O'Connor	University of Maryland Baltimore
Luca Pinello	Massachusetts General Hospital
Jason Vassy	Harvard Medical School

 Supports early-career researchers to do highly innovative work on important problems in genomics

• Annual receipt date October 30

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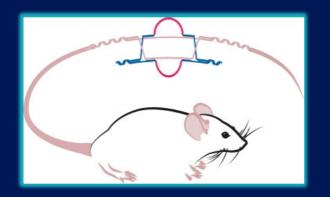


#### Human Microbiome Project (HMP)

- HMP Phase 2 Collection: Nature May 29, 2019
- Four flagship papers, 20+ companion papers, & perspective
- All HMP data, tools, & protocols at HMP Data Coordination Center



#### **Knockout Mouse Phenotyping Project (KOMP2)**





#### KOMP2 Community Workshop – June 2019

Prioritize completing mouse protein-coding genome Strong support for generating conditional alleles Valuable to the rare disease community Increase outreach and interaction with research community

#### KOMP2 Annual Meeting – November 2019

Focus on collaborations with human disease gene discovery programs

#### **Genotype-Tissue Expression Project (GTEx)**



**Version 8 Data Release:** 

**SNP/CNV** genotypes: 900 subjects

**RNA-Seq expression: 17,000 samples** 

Allele-specific expression: 16,000 samples





Analysis, Visualization and Informatics Labspace (AnVIL)



## Human Heredity and Health in Africa (H3Africa)

#### 48 projects, 34 countries



#### 14<sup>th</sup> Consortium Meeting in Accra, Ghana



**25 Trainee Travel Fellowships** 





#### Human Heredity and Health in Africa (H3Africa)

#### H3Africa Biorepositories Renewed for Phase II



#### Kampala, Uganda



Johannesburg, South Africa

Abuja, Nigeria

#### H3Africa Biorepository Website Launching Soon

Our Services H3Africa Biorepository Services

#### Useful Links

- The Human Heredity and Health in Africa
- Integrated Biorepository of H3Africa Uganda
- > Clinical Laboratory Services, South Africa
- Institute of Human Virology Nigeria
- > Pan African Bioinformatics Network for H3Africa
- International Society for Biological and Environmental Repositories (ISBER)
- The Alliance for Accelerating Excellence in Science in Africa (AESA)
- National Institutes of Health (NIH)

#### **Recent Posts**

Our Mission is to develop, maintain and sustain state-of-the-art world class Biorepositories for the responsible storage, maintenance, and custodianship of well-annotated high-quality Biospecimens and to make these specimens available to researchers for genomic discovery and other biomedical research. The H3africa Biorepositories offer biological specimens collection, storage, processing, and shipment as well as Biorepository quality control/assurance related services, Training in Biorepository Science and Management, Sanger and Next Generation sequencing services.

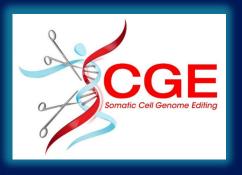
#### **Biospecimen Collection/Reception**

☑ MAY 02, 2019 SIOREPOSITORY SERVICES

**F** 



#### **Somatic Cell Genome Editing**



#### Six components:

New Delivery SystemsIn Vivo TrackingAnimal ReportersNovel Genome EditorsTissue PlatformsData Coordination CenterSecond round of awards now in process

• Annual Program Meeting in December 2019

# RESEARCH PROGRAM

#### SPECIAL REPORT



# The "All of Us" Research Program

The All of Us Research Program Investigators



# RESEARCH PROGRAM



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### **Congressional Outreach Activities**





#### New Resource Celebrating Henrietta Lacks and the Impact of HeLa Cells on Biomedical Research



#### HeLa Cells: A Lasting Contribution to Biomedical Research

In 1951, Henrietta Lacks, a 31-year-old African-American woman, went to Baltimore's Johns Hopkins Hospital to be treated for cervical cancer. Some of her cancer cells began being used in research due to their unique ability to continuously grow and divide in the laboratory. These socalled "immortal" cells were later named "HeLa" after the first two letters of **He**nrietta **La**cks first and last name.

Since Ms. Lacks' untimely death in 1952, HeLa cells have been a vital tool in biomedical research, leading to an increased understanding of the fundamentals of human health and disease. Some of the research involving HeLa cells also served as the underpinning of several Nobel Prize winning discoveries.

While Henrietta Lacks' story has been known in the research community for some time, it raised further awareness after the publication of the best-selling book *The Immortal Life of Henrietta Lacks* (Crown, 2010).

To honor Ms. Lacks' and her family's continued support of biomedical research, NIH analyzed and evaluated the scientific literature involving HeLa cells and found over 10,000 publications that cited the use of HeLa cells between 1953 to 2018. This analysis further highlights the persistent impact of HeLa cells in science and medicine, proving that they have been a consistent, essential tool that has allowed researchers to expand the knowledge base in fields such as cancer biology, infectious disease, and many others.



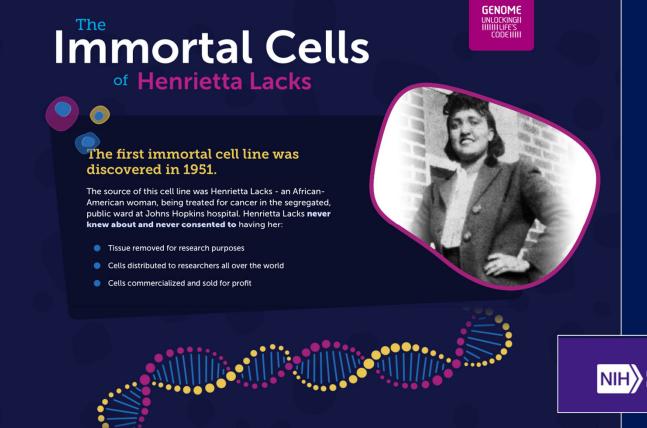
Policy Areas Outreach

Library

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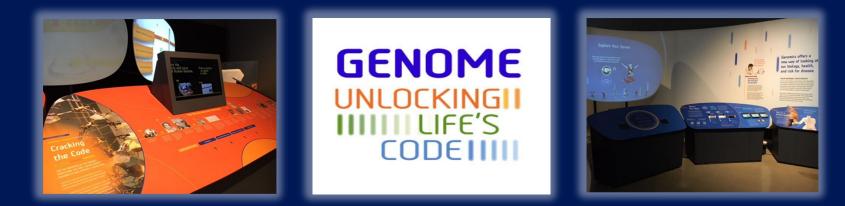
#### **New Lesson Plans Featuring Henrietta Lacks**





NATIONAL MUSEUM of AFRICAN AMERICAN HISTORY & CULTURE

#### Genome: Unlocking Life's Code Exhibition Travel Schedule



September 12, 2019 – January 2, 2020
 Turtle Bay Exploration Park, Redding, CA

 January 17 – April 12, 2020
 Museum of Science & History, Jacksonville, FL

#### **NHGRI Short Course in Genomics**



National Human Genome Research Institute Short Course in Genomics

# **NHGRI Summer Outreach Program**







#### NHGRI's South by Southwest Conference Proposals



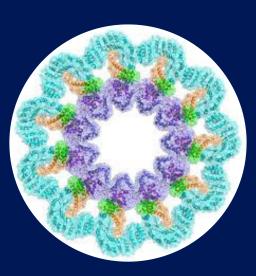
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#### **Ross Prize in Molecular Medicine**







# **Departure of NHGRI Intramural Investigator**



Max Muenke, M.D.

#### International Summit in Human Genetics and Genomics



#### **NHGRI Intramural Research Highlights**





ZRANB3 is an African-Specific Type 2 Diabetes Locus Associated with Beta-Cell Mass and Insulin Response



# **Science** Advances

*De novo* assembly of the goldfish genome and evolution of genes after whole-genome duplication



Genetics inMedicine

Constructing identities: the implications of DTC ancestry testing for tribal communities

# The Genomics Landscape

#### **Email Updates**

Sign up to receive National Human Genome Research Institute (NHGRI) updates and stay informed about our latest science, research, news, upcoming events and website content.

Email Address



# **Thanks!**



# **Special Thanks!**

