



From the Human Genome Project to Genomic Medicine

A Journey to Advance Human Health

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



The Origin of “Genomics”: 1987

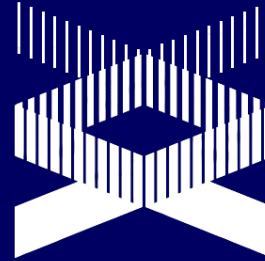
EDITORIAL

A New Discipline, A New Name, A New Journal

Genomics (1987)

“For the newly developing discipline of [genome] mapping/sequencing (including the analysis of the information), we have adopted the term GENOMICS...

‘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

NHGRI: *Circa 1990-2003*

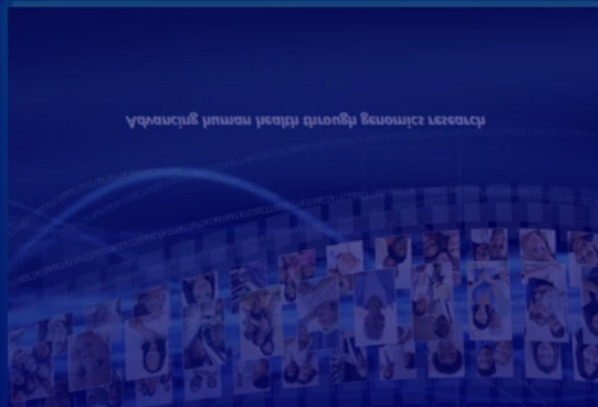


Human Genome Project

NHGRI Today: Characteristic Features



- Relatively young (~28 years)
- Relatively small (~1.7% of NIH)
- Unusual historical origins (think 'Human Genome Project')
- Emphasis on 'Team Science' (think managed 'consortia')
- Rapidly disseminating footprint (think 'genomics')
- Novel societal/bioethics research component (think 'ELSI')
- Over-achievers for trans-NIH initiatives (think 'Common Fund')
- Vibrant (and large) Intramural Research Program



A Quarter Century of Genomics

**Human Genome Sequenced for First Time
by the Human Genome Project**



Genomic Medicine

An emerging medical discipline that involves using genomic information about an individual as part of their clinical care (e.g., for diagnostic or therapeutic decision-making) and the other implications of that clinical use



The Path to Genomic Medicine



**Human
Genome
Project**



**Realization of
Genomic
Medicine**

A blueprint for the genomic era.

The completion of a high-quality, comprehensive sequence of the human genome, in this fiftieth anniversary year of the discovery of the double-helical structure of DNA, is a landmark event. The genomic era is now a reality.

the creativity and
of talented sci-
this project the
the initial obje-
been achieved
expectation, a
research has be-

NATURE/VOL. 422

Our ability to explore genome function is increasing in specificity as each subsequent genome is sequenced. Microarray technologies have catapulted many laboratories from studying the expression of one or two genes in a month to studying the expression of tens of thousands of genes in a single afternoon¹⁰. Clinical opportunities for gene-based pre-symptomatic prediction of illness and adverse drug response are emerging at a rapid pace, and the therapeutic targeting of molecular biology has replaced

836

1000

© 2000 Blackwell Science Ltd

836

Base Pairs to Bedside

Heli2011 Health

doi:10.1038/nature09764

Eric D. Green¹, Mark S. Guyer³ & National Human Genome Research Institute⁴

Since the first of the Human Genome Project (HGP) in 2003 and the publication of a reference human genome sequence^{1,2}, genomics has become a mainstay of biomedical research. The scientific community's foresight in launching this ambitious project is evident in the broad range of scientific advances that the HGP has enabled, as shown in Figure 1. The HGP has not only provided a framework for understanding the molecular basis of inherited diseases (<http://www.ncbi.nlm.nih.gov/omim>) and <http://www.genome.gov/GWASStudies>) and the discovery of structural variation in disease³, some of which have already led to new therapies^{4,5}. Other advances are already changing medical practice (for example, microarrays are now used in the diagnosis of genomic imbalances^{6,7}), and the use of genotyping is routinely performed by health administrators to certify certain medications^{8,9}. Together, these achievements (see accompanying paper)¹⁰ document that genomics is contributing to a better understanding of human biology and to improving human health.

This 2011 vision for genomics, from basic research to clinical applications, is that, over time, the modern Genome Project will help us understand normal biology and disease, and how to understand disease better to improve human health. At the same time, the Genomics offers opportunities for understanding of disease and health based on genomic profiles. Genomic profiles can lead back to the next level of understanding.

quidly. Although genomics has already begun to improve diagnostics and treatments in a few circumstances, profound improvements in the effectiveness of healthcare cannot realistically be expected for many years (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).

...uniquely valuable and
...rove existing catalogues
...tions of genetic variation,
...at, and other biological

associated with disease of genetic variation, which has and variants for identification of variation in the that began with The SNP Project* (<http://hapmap.org>) and the 1000 Genomes Project*.

Nature

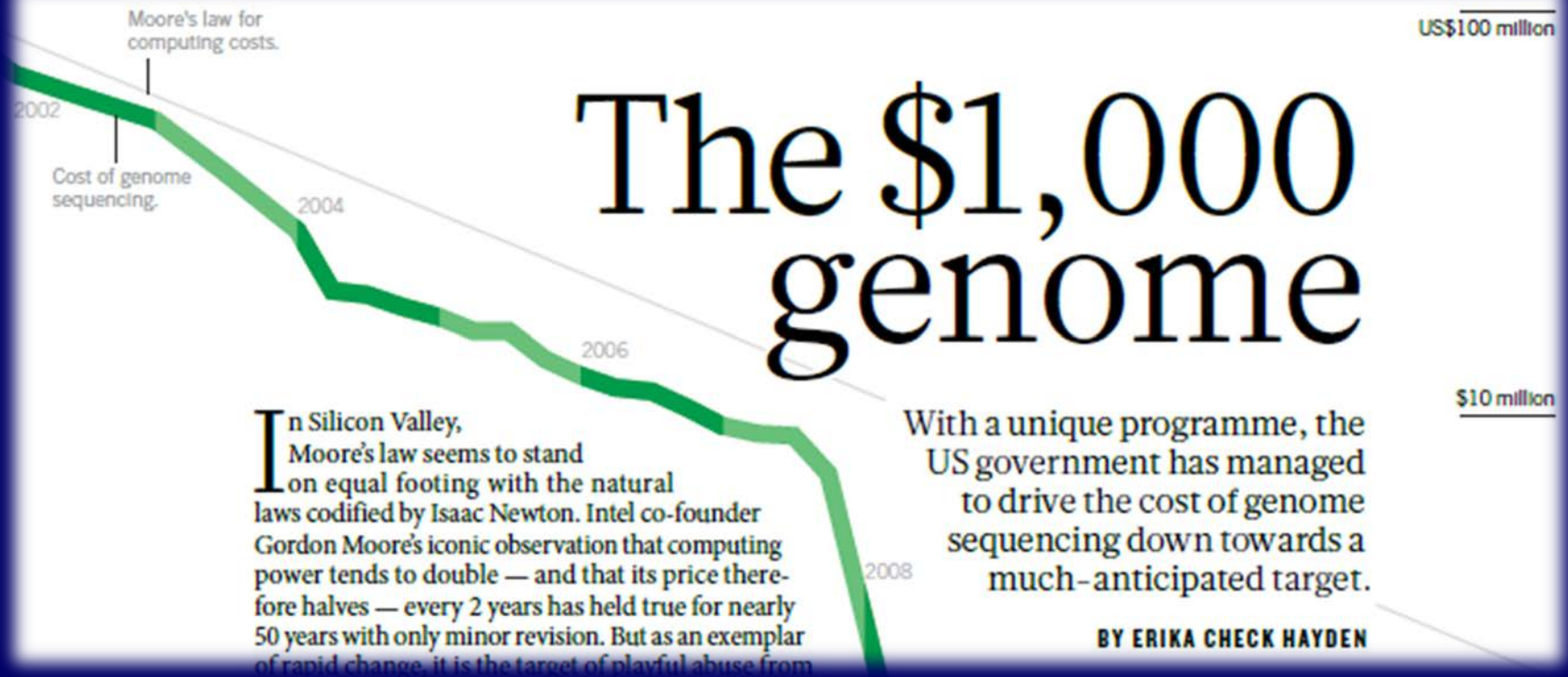


A Quarter Century of Genomics

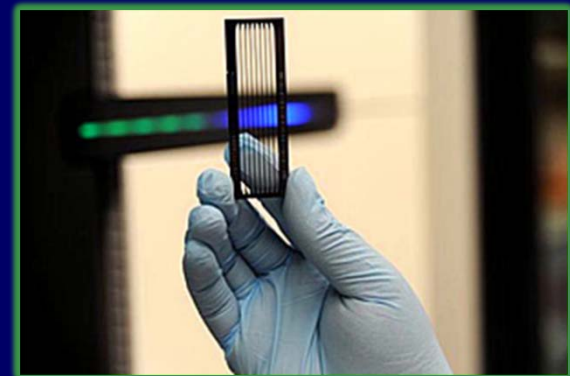
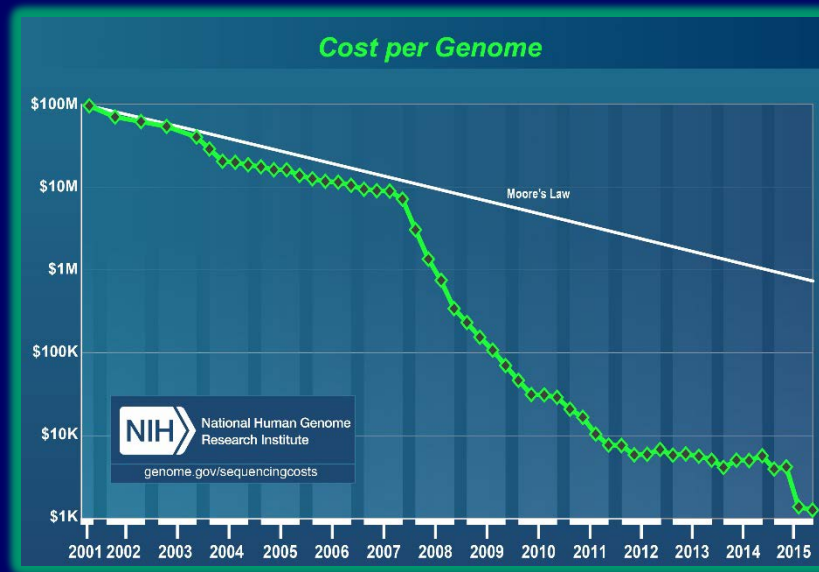


**Human Genome Sequenced for First Time
by the Human Genome Project**

**Cost of Sequencing a Human Genome
Reduced Nearly ~1 Million-Fold**



Nature (2014)



A Quarter Century of Genomics



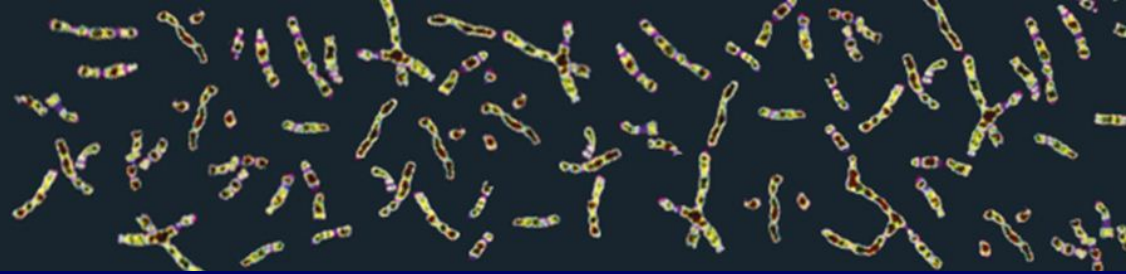
**Human Genome Sequenced for First Time
by the Human Genome Project**

**Cost of Sequencing a Human Genome
Reduced Nearly ~1 Million-Fold**

**Tens of Thousands of Human
Genomes Sequenced**

1000 Genomes

A Deep Catalog of Human Genetic Variation



ARTICLE

A global reference for human genetic variation

The 1000 Genomes Project Consortium*

Nature (2015)

A Quarter Century of Genomics



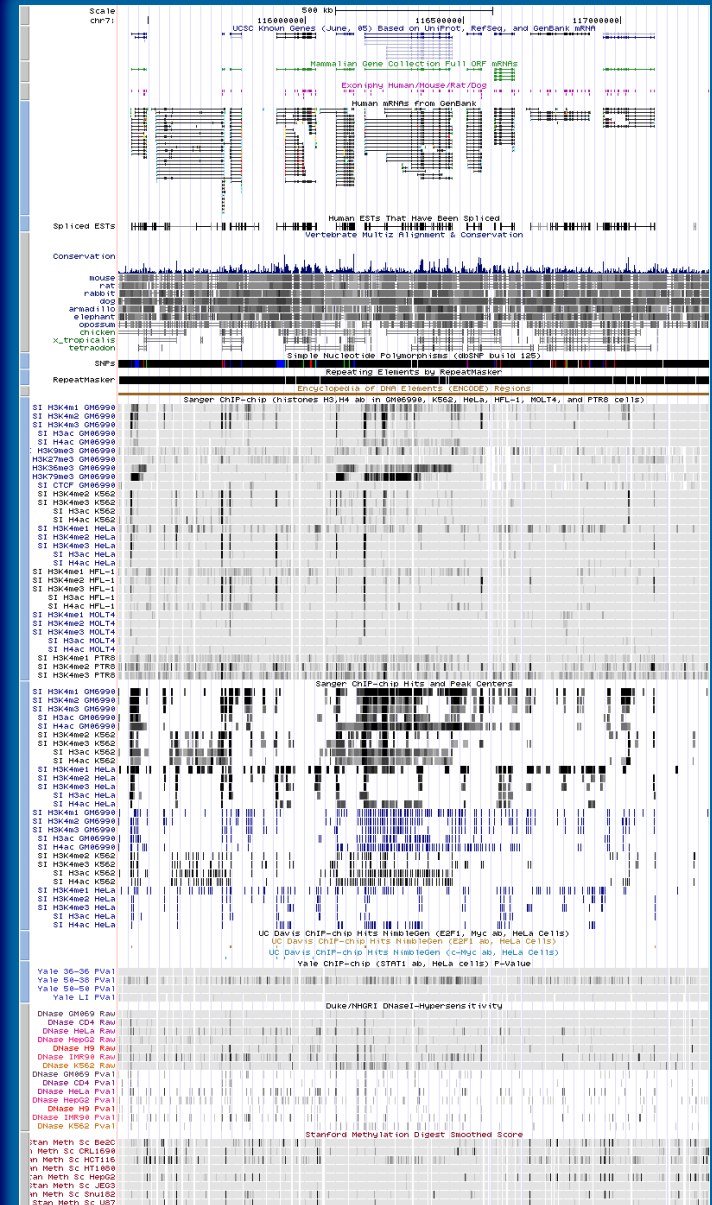
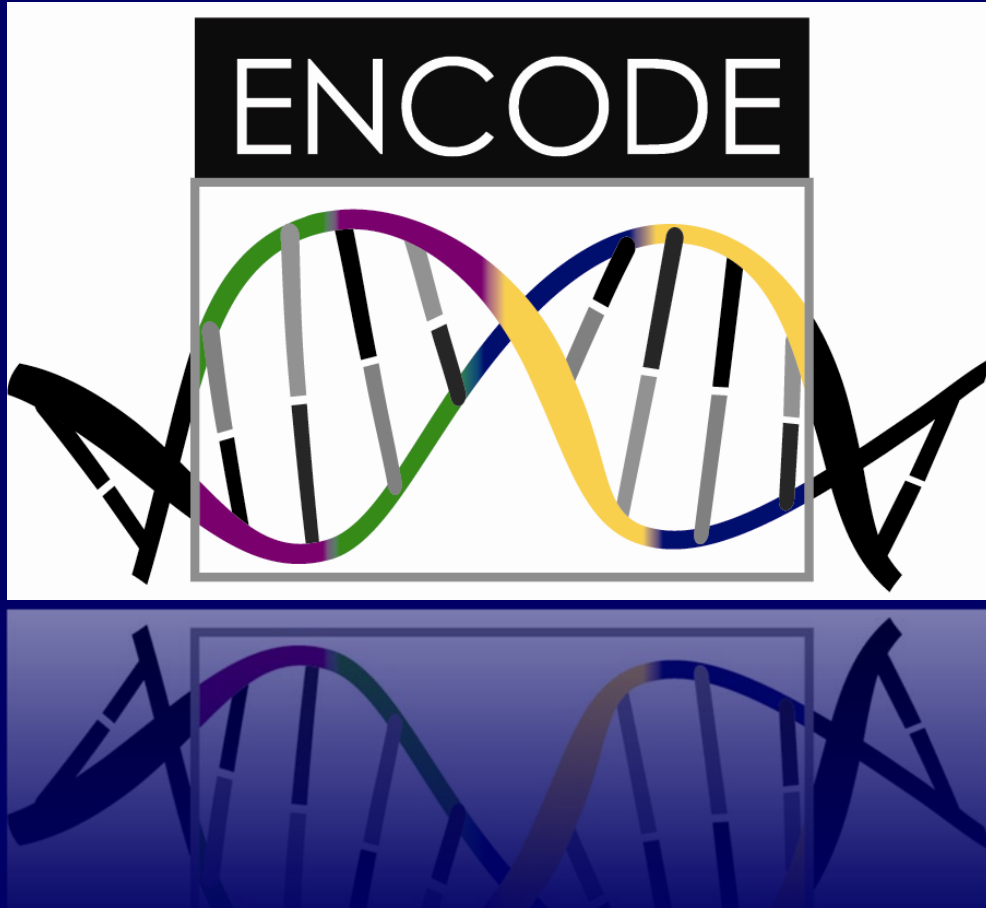
**Human Genome Sequenced for First Time
by the Human Genome Project**

**Cost of Sequencing a Human Genome
Reduced Nearly ~1 Million-Fold**

**Tens of Thousands of Human
Genomes Sequenced**

**Profound Advances in Understanding
How the Human Genome Functions**

ENCODE: Giving 'GPS' Views of Genomes



A Quarter Century of Genomics



**Human Genome Sequenced for First Time
by the Human Genome Project**

**Cost of Sequencing a Human Genome
Reduced Nearly ~1 Million-Fold**

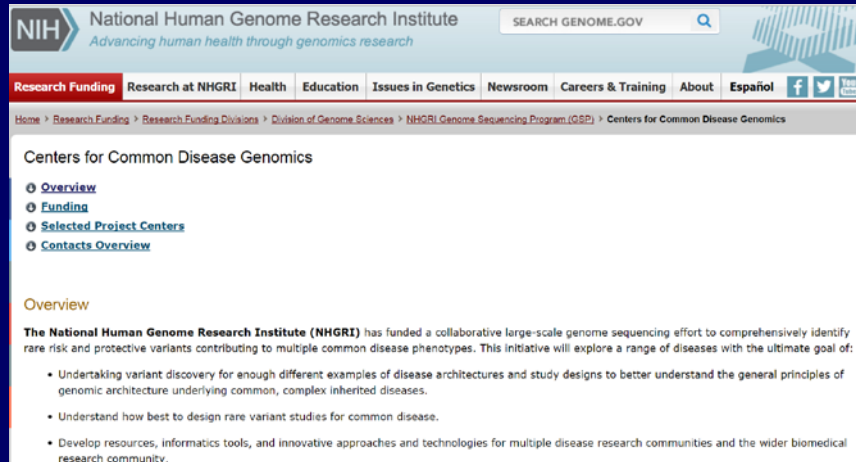
**Tens of Thousands of Human
Genomes Sequenced**

**Profound Advances in Understanding
How the Human Genome Functions**

**Significant Advances in Unraveling the
Genomic Bases of Human Disease**

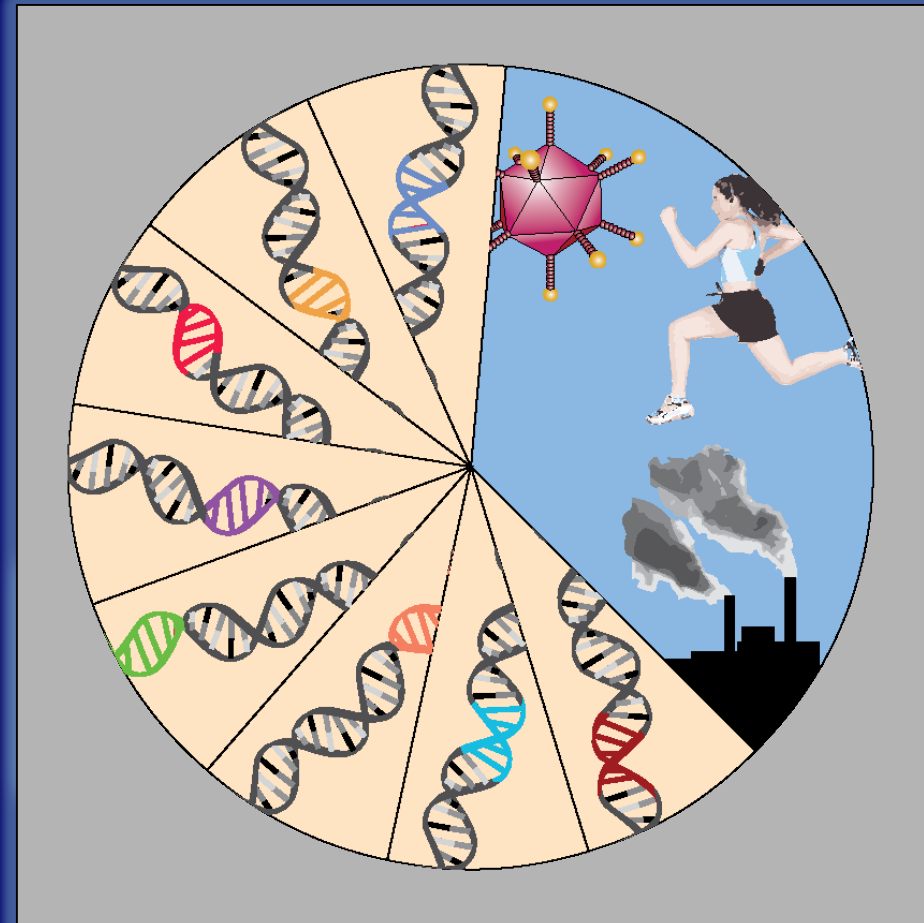
Genomic Architecture of Genetic Diseases

NHGRI Centers for Common Disease Genomics



genome.gov/27563570

Cardiovascular Disease
Hemorrhagic Stroke
Autism
Epilepsy
Asthma



**Common, Complex, Multigenic,
Non-Mendelian...**

Manolio et al., J Clin Invest (2008)

A Quarter Century of Genomics



**Human Genome Sequenced for First Time
by the Human Genome Project**

**Cost of Sequencing a Human Genome
Reduced Nearly ~1 Million-Fold**

**Tens of Thousands of Human
Genomes Sequenced**

**Profound Advances in Understanding
How the Human Genome Functions**

**Significant Advances in Unraveling the
Genomic Bases of Human Disease**

**Vivid Examples of Genomic Medicine
in Action Now Emerging**

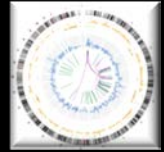
Bringing Genomic Medicine Into Focus



'Hot Areas' in Genomic Medicine



Cancer Genomics



Pharmacogenomics



Electronic Medical Records and Genomics (eMERGE) Network

LOGIN TO EMERGE

emerge network
ELECTRONIC MEDICAL RECORDS AND GENOMICS



451

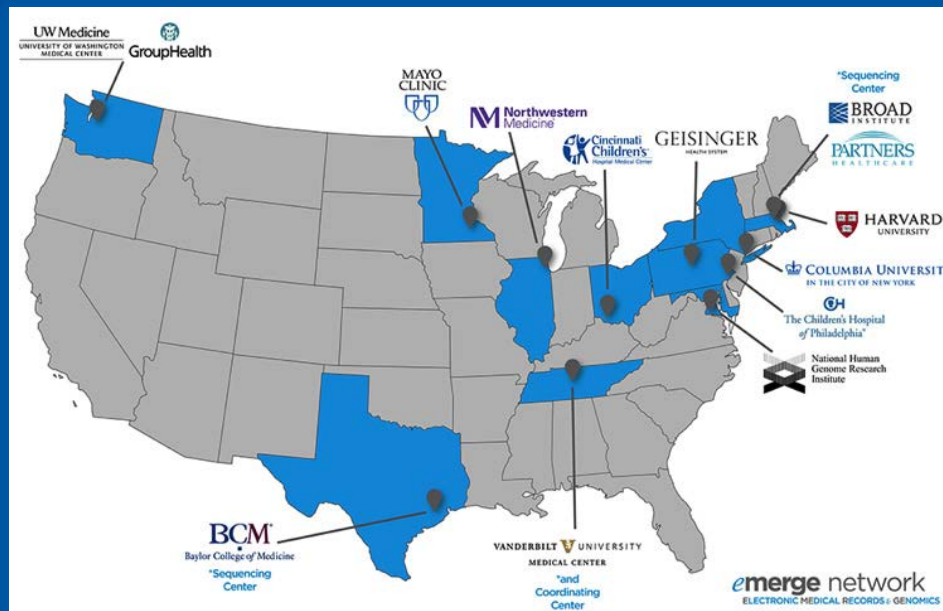
Number of
network
publications

47

Number of
phenotypes
developed

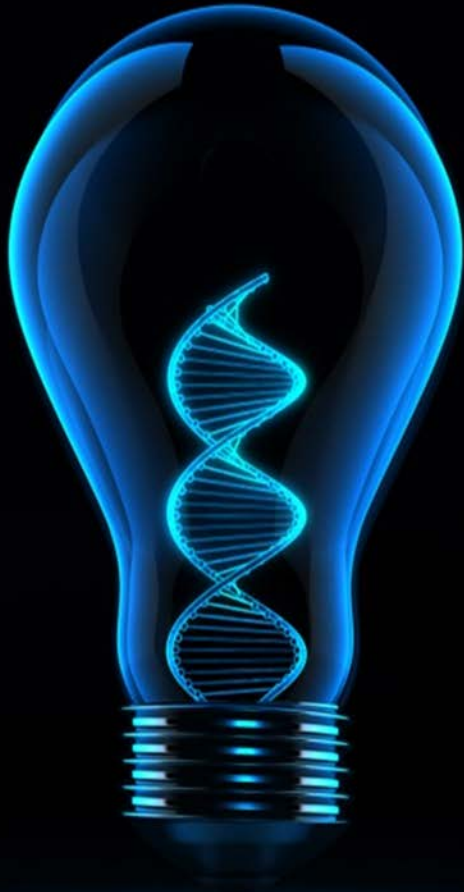
55,028

Number of participants in the Network Cohort

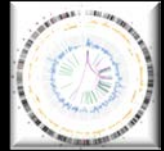


emerge.mc.vanderbilt.edu

'Hot Areas' in Genomic Medicine



Cancer Genomics



Pharmacogenomics



**Rare Genetic Disease
Diagnostics**



TECHNOLOGY FEATURE

WHEN DISEASE STRIKES FROM NOWHERE

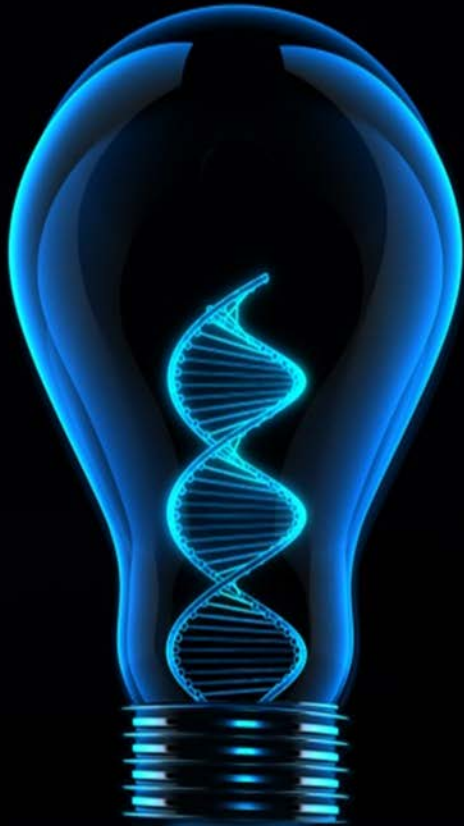
When healthy parents have a child with a genetic disorder, the cause is sometimes a new mutation. Tools are emerging to meet the challenge of finding such changes.



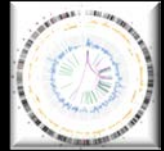
“ ...disorders not readily explained by standard tests can sometimes be diagnosed through genome sequencing and analysis.”

Nature (2014)

'Hot Areas' in Genomic Medicine



Cancer Genomics



Pharmacogenomics



**Rare Genetic Disease
Diagnostics**



**Newborn Genome
Sequencing**



Newborn Genome Sequencing

HEALTH RESEARCH

In 2025, Everyone Will Get DNA Mapped At Birth

Alice Park @aliceparkny | June 30, 2014



Scientists have scoured trends in research grants, patents and more to come up with these 10 innovations that will be reality in 10 years (or so they think)

Everybody likes to blue-sky it when it comes to technology. Driverless cars! Fat-burning pills! Telepathic butlers! But the folks at Thomson Reuters Intellectual Property & Science do it for a living—and they do it with data.



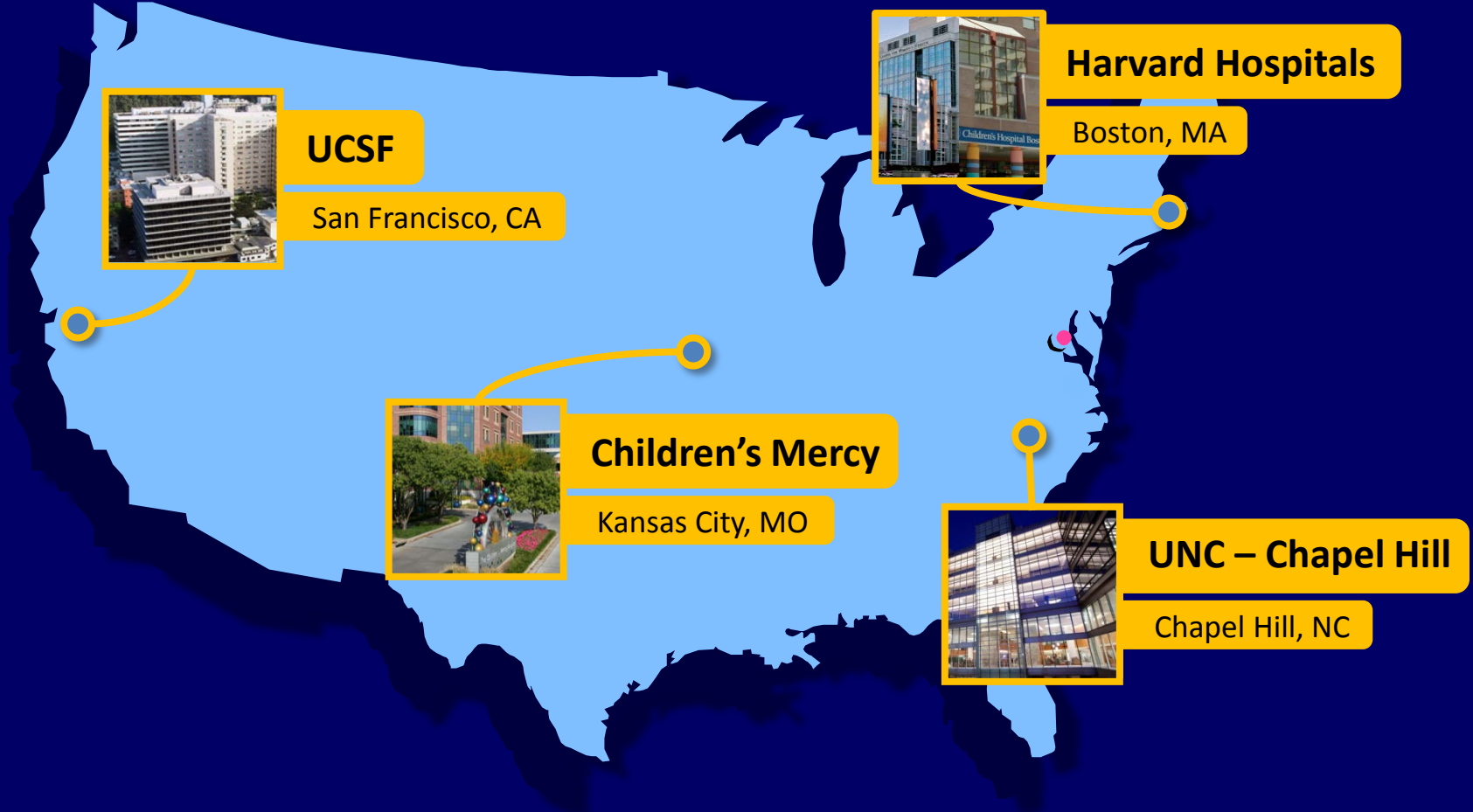
What will the future hold?

REB Images—Getty Images/Blend Images

Time (2014)



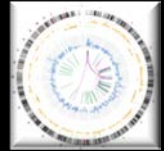
Newborn Sequencing In Genomic medicine and public Health (NSIGHT)



'Hot Areas' in Genomic Medicine



Cancer Genomics



Pharmacogenomics



**Rare Genetic Disease
Diagnostics**



**Newborn Genome
Sequencing**



**Clinical Genomics
Information Systems**

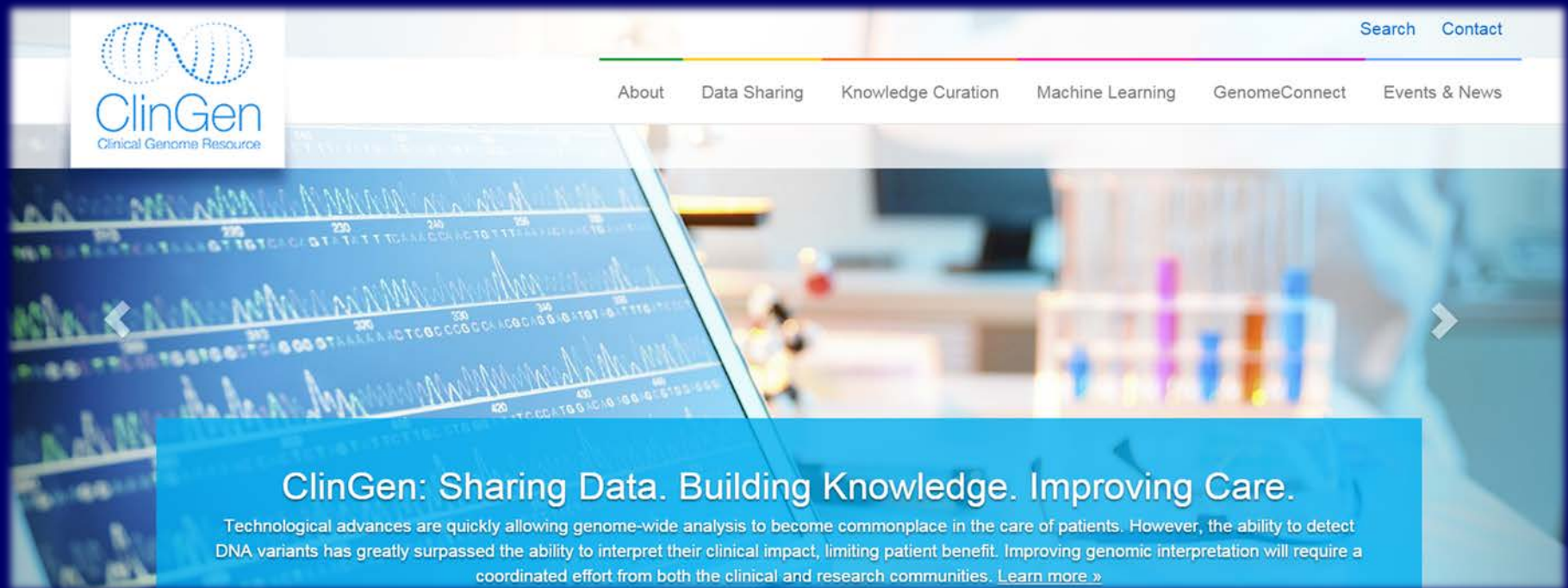




Clinical Genomics Information Systems



Clinical Genome Resource (ClinGen)



clinicalgenome.org

ClinGen — The Clinical Genome Resource

Heidi L. Rehm, Ph.D., Jonathan S. Berg, M.D., Ph.D., Lisa D. Brooks, Ph.D.,
Carlos D. Bustamante, Ph.D., James P. Evans, M.D., Ph.D., Melissa J. Landrum, Ph.D.,
David H. Ledbetter, Ph.D., Donna R. Maglott, Ph.D., Christa Lese Martin, Ph.D.,
Robert L. Nussbaum, M.D., Sharon E. Plon, M.D., Ph.D., Erin M. Ramos, Ph.D.,
Stephen T. Sherry, Ph.D., and Michael S. Watson, Ph.D., for ClinGen

NEJM (2015)

Genomics and Society



NHGRI Extramural Research Program: Circa 2016

\$1000 Genome

Technology Development
Program

TCGA

The Cancer Genome
Atlas

PAGE

Population Architecture using
Genomics and Epidemiology

ENCODE

Encyclopedia of DNA
Elements Project

GGR

Genomics of Gene
Regulation

eMERGE

Electronic Medical
Records and Genomics

PhenX

Consensus Measures for
Phenotypes and eXposures

NoVa

Non-Coding Variants
Program

CEGS

Centers of Excellence in
Genomic Science

CSER

Clinical Sequencing
Exploratory Research

IGNITE

Implementing Genomics in
Practice Network

CCDG

Centers for Common
Disease Genomics

CMG

Centers for Mendelian
Genomics

NSIGHT

Newborn Sequencing In Genomic
medicine and public Health

ClinGen

Clinical Genome
Resource

ELSI

Ethical, Legal, and Social
Implications Program

CEERs

Centers for Excellence
in ELSI Research

BD2K

Big Data to Knowledge
Initiative

PMI

Precision Medicine
Initiative

KOMP2

KOMP
Phenotyping

HMP

Human Microbiome
Project

GTEx

Genotype-Tissue
Expression

UDN

Undiagnosed Diseases
Network

**Protein Capture
Reagents**

LINCS

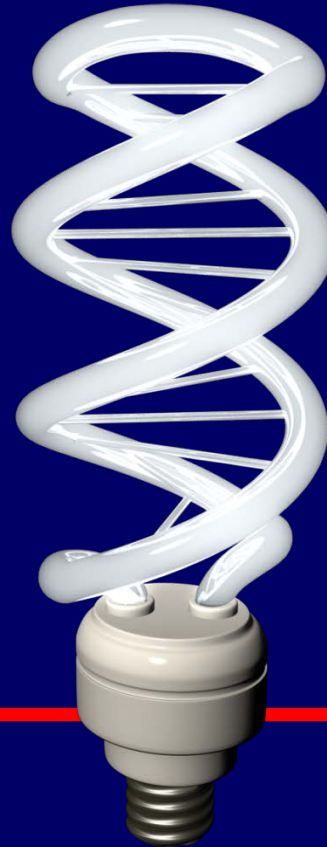
Library of Integrated Network-
based Cellular Signatures

H3Africa

Human Heredity and
Health in Africa

Kids First

Gabriella Miller Pediatric
Research Program





October 6, 2015

This month brought a historic 'odometer moment' for the field of genomics – October 1, 2015, marked the 25th anniversary of the launch of the Human Genome Project. I, for one, cannot believe a quarter-century has now passed since many of us started working on the Project. At the same time, it is truly incredible to think about how far genomics has progressed since that time. I thought the significance of this anniversary warranted making this topic the lead story in this month's *The Genomics Landscape*; in addition, I reflect on this important anniversary in a recent video interview now available on the NHGRI web site.

To subscribe, follow link from:
genome.gov/Director



NATIONAL HUMAN GENOME RESEARCH INSTITUTE



***Advancing human health
through genomics research***