



The Human Genome: Unlocking Life's Code

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



The Relevance of Genomics



Biomedical Researchers

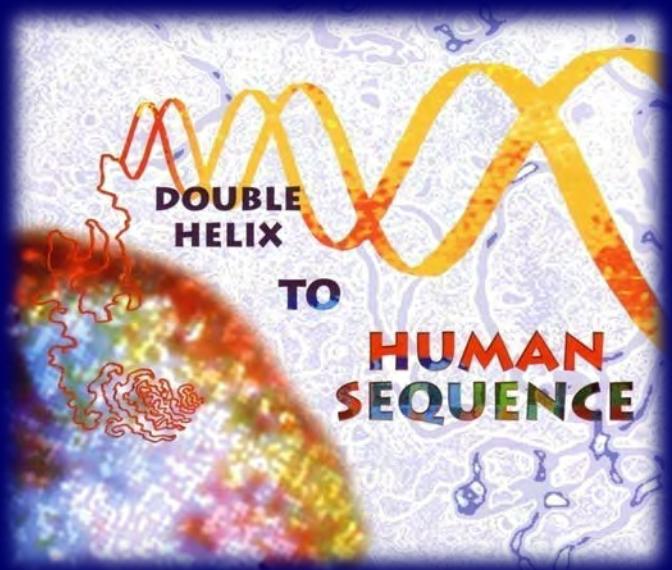


Healthcare Professionals



Patients (and Friends & Relatives of Patients)

Human Genome Project 1990-2003



HUMAN GENOME
GCCAAAGTATACT
TTTCAGCCAACAT
ATCTCCACTCTCT
AACGAGGGAAAT
ATCTGTATGTATC
AGGGAAAAAA





NIH

*Turning discovery into
health*

*Advancing human health
through genomics research*



Genomic Medicine

An emerging medical discipline that involves using an individual's genomic information as part of their clinical care



February, 2011

nature

genome.gov/sp2011

THE FUTURE IS BRIGHT

Reflections on the first ten years of the human genomics age

GENOMICS
THE END OF THE BEGINNING
Eric Lander on the impact of the human genome sequence
PAGE 187

METHODS
MORE BASES PER DOLLAR
Elaine Mardis on the march of sequencing technology
PAGE 198

HEALTH
FROM LAB TO CLINIC
A road map to genomic medicine
PAGE 204

NATURE ASIA.COM
10 February 2011
Vol. 470, No. 7333

PERSPECTIVE

doi:10.1038/nature09764

Charting a course for genomic medicine from base pairs to bedside

omim and <http://www.genome.gov/GWAStudies>) and the role of structural variation in disease⁸, some of which have already led to new therapies^{9–13}. Other advances have already changed medical practice (for example, micro-arrays are now used for clinical detection of genomic imbalances¹⁴ and pharmacogenomic testing is routinely performed before administration of certain medications¹⁵). 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 updated vision 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 genetics 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

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 widely 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 Project²¹ (<http://hapmap.ncbi.nlm.nih.gov>), and is ongoing with the 1000 Genomes Project²² (<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

Figure 1 | Genomic achievements since the Human Genome Project (see accompanying roll-off). ►

¹National Human Genome Research Institute, National Institutes of Health, 33 Center Dr., Bethesda, Maryland 20892-2152, USA.

²Lists of participants and their affiliations appear at the end of the paper.

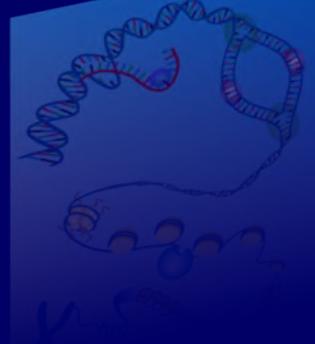
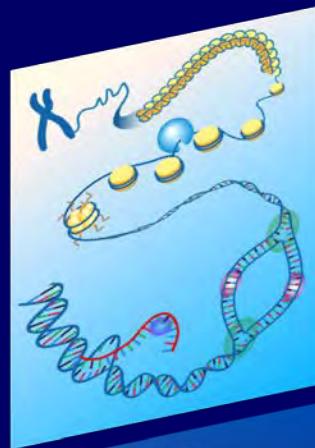
New NHGRI Vision for Genomics Published

Five Domains of Genomics Research

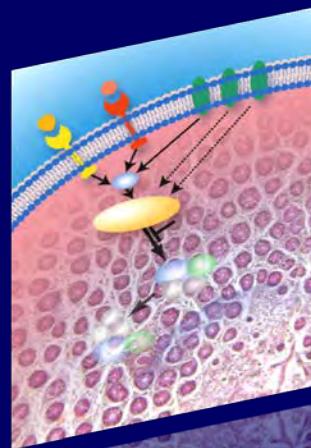
Understanding
the Structure of
Genomes



Understanding
the Biology of
Genomes



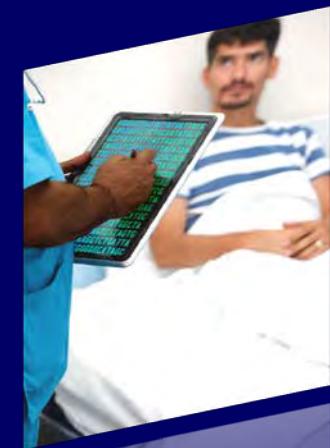
Understanding
the Biology of
Disease



Advancing
the Science of
Medicine

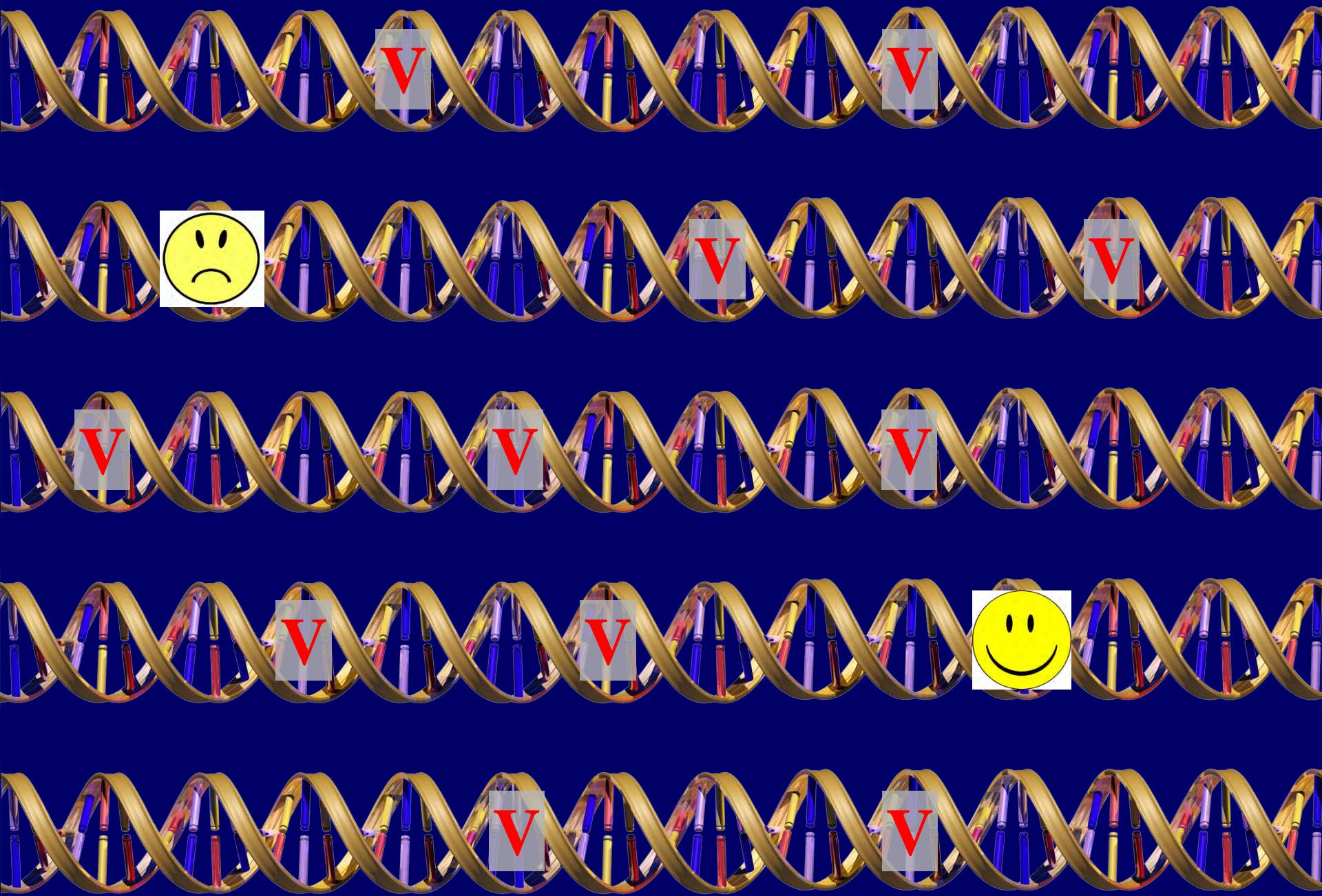


Improving the
Effectiveness
of Healthcare



~3,000 bp (0.0001%) of Human Genome Sequence

TGCGCGGAACCTTCGGCTCTCTAAGGCTGTATTTGATATACGAAAGGCACATTTCCCTTCAAAATGCACCTTGCAAACGTAACAG
GAACCCGACTAGGATCATCGGGAAAAGGAGGAGGAGGAAGGCAGGCTCCGGGAAGCTGGTGGCAGCGGGCTGGTCTGGCGGACCCCTGA
CGCGAAGGAGGGTCTAGGAAGCTCTCCGGGAGCCGGTTCTCCGCCGGTGGCTCTGTCCAGCGTTGCCAACTGGACCTAAAGAGAGG
CCCGACTGTGCCAACCTCGGGATGGGCCTGGTCTGGCGGTAAAGGACACGGACCTGGAAGGAGCGCGCGAGGGAGGGAGGCTGGGAGTC
AGAATCGGGAAAGGAGGTGGGGCGCGAGGGAGCGAAGGAGGAGGAGGAAGGAGCGGGAGGGTGTGGCGGGGTGCGTAGTGGGTGGA
GAAAGCCCTAGAGCAAATTGGGGCCGGACCAGGCAGCAGTCTGGCTTTAACCTGGCAGTGAAGGCGGGGAAAGAGCAAAGGAAGGGTGG
TGTGCGGAGTAGGGTGGGTGGGGGAAATTGGAAGCAAATGACATCACAGCAGGTAGAGAAAAAGGGTGTAGCGGCAGGCACCCAGAGTAGTAG
GTCTTGGCATTAGGAGCTTGAGCCCAGACGCCCTAGCAGGACCCCAGCGCCCGAGAGACCATGCAGAGGTGCGCTCTGGAAAAGGCCAGCGT
TGTCTCCAAACTTTTCAGGTGAGAAGGTGCCAACCGAGCTCGGAAAGACACGTGCCACGAAAGAGGAGGGCGTGTGTATGGGTTGGGTT
TGGGGTAAAGGAATAAGCAGTTTTAAAAGATGCGCTATCATTCAATTGTTGAAAGAAAATGTGGGTATTGAGAATAAAACAGAAAGCATT
AGAAGAGATGGAAGAATGAACGTGAGCTGATTGAATAGAGAGCCACATCTACTTGCAACTGAAAAGTTAGAATCTCAAGACTCAAGTACGCTACT
ATGCACTTGTTTATTCATTCTAAGAAACTAAAATACTTGTAAATAAGTACCTAAGTATGGTTATTGGTTTCCCCCTCATGCCTTGG
ACACTTGATTGCTTCTGGCACATACAGGTGCCATGCCTGCATATAGTAAGTGCTCAGAAAACATTCTTGACTGAATTCAAGCCAACAAAATT
TTGGGGTAGGTAGAAAATATGCTTAAAGTATTGTTATGAGACTGGATATCTAGTATTGTCACAGGTAAATGATTCTTCAAAAATTG
AAAGCAAATTGTTGAAATATTGAAAAAGTTACTTCACAAGCTATAAATTAAAAGCCATAGGAATAGATAACCGAAGTTATCCAA
CTGACATTAAATAAAATTGATTCATAGCCTAATGTGATGAGCCACAGAAGCTGCAAACCTTAATGAGATTTTAAAATAGCATCTAAGTTCGG
AATCTTAGGCAAAGTGTGTTAGATGTGACTTCATATTGAAGTGTCTTGGATATTGCATCTACTTGTTCTGTTATTATACTGGTGTGA
ATGAATGAATAGGTACTGCTCTCTGGGACATTACTTGACACATAATTACCAATGAATAAGCATACTGAGGTATCAAAAAGTCAAATATGT
TATAAAATAGCTCATATATGTTAGGGGGAGGAATTAGCTTCACTCTCTTATGTTAGTTCTCTGCATGTCAGTTAACCTGGAAAC
TCCGGTCTAAGGAGAGACTGTTGCCCTTGAAGGAGAGCTCCCTGTGGATGAGAGAGAAGGACTTTACTCTTGGAAATTATCTTTGTGT
TGATGTTATCCACCTTGTACTCCACCTATAAAATCGGCTTATCTATTGATCTGTTCTAGTCCTTATAAAAGTCAAATGTTAATTGGCAT
AAATTATAGACTTTTTAGCAGAGAACCTTGAGGAACCTAAATGCCAACAGTCTAAAATGCAGTTCAAGAAGAATGAATAATTGATGGATA
GTTCTAAATACTAATGAACCTTAAAGCTTACTATTGATCTGCTAAAGTGGTTTATATAATTCTTTACAAATCACCTGACACATT
AATATAGGTAAAAATGCTATCAGGCTGGTTGCAAAGAAAATGTATTACAAAGGCTGCTAAGTGTGTTAAGAGCATACTCATTTCTGTTCTCC
AAAATTTCTACAGGTGCTTAAGAATAGGTATGTTTAAAAGTTAACCTACTATTATAGGAACCTGACAATCACCTAAAATACCAATGA
TTACAAACTCCTCTGGCCTCTGGACTGCAATTCTAAAAGTGTAAAAAACATATTCTGCATTAAGTTAGGCAGTATTGCTTAGTTCTGAA
GTGGTAGGCTTGGAGTCAGATTATTTGATTGATCAGATCCTACATCTACTGTTAGTAGCTCTGTTGCCTGAGGCAGGTCCCTAACATCTGTG
TGTGACTTGACCTTAAAATTGGAGACTGTCATAGGGTTAACCTTGAGAAAATGAATGTGAAAAGTTAGCCTAATGTTAAGTCTATT
ATGGATTACCATATTTCACATTGATCACAGTACATGCACCTGTTAATATAAGATGCTCAATTGATCTTGTGACTCTCAAT
CTGGATATGCAATGAGTGGCCTGTATGAGAATTAAATTGTTACATGGCCTTACAGATACAGGAAACACGTACATG
TTCTATTGATGTTAAATGCCTAGAATTACTTCTGAATAGGATCCCTCAGTTGAGAGTCATAAAAGAGTAAAATTATTGTT

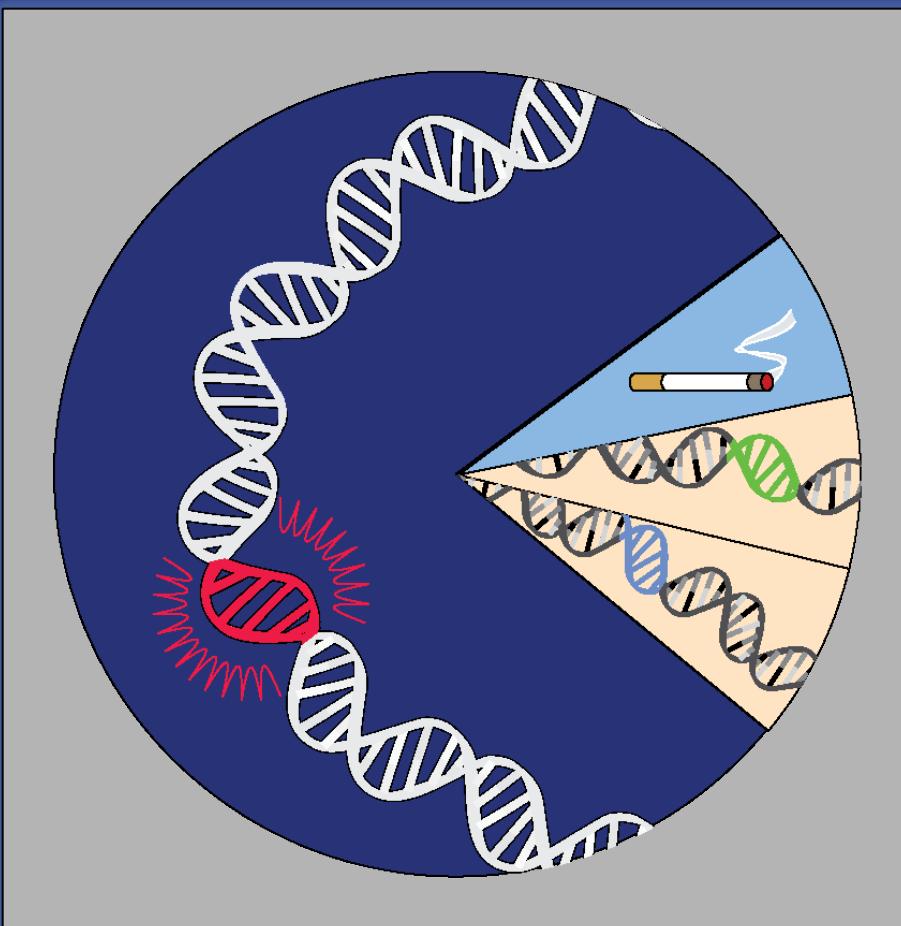


Your Genome: By the Numbers



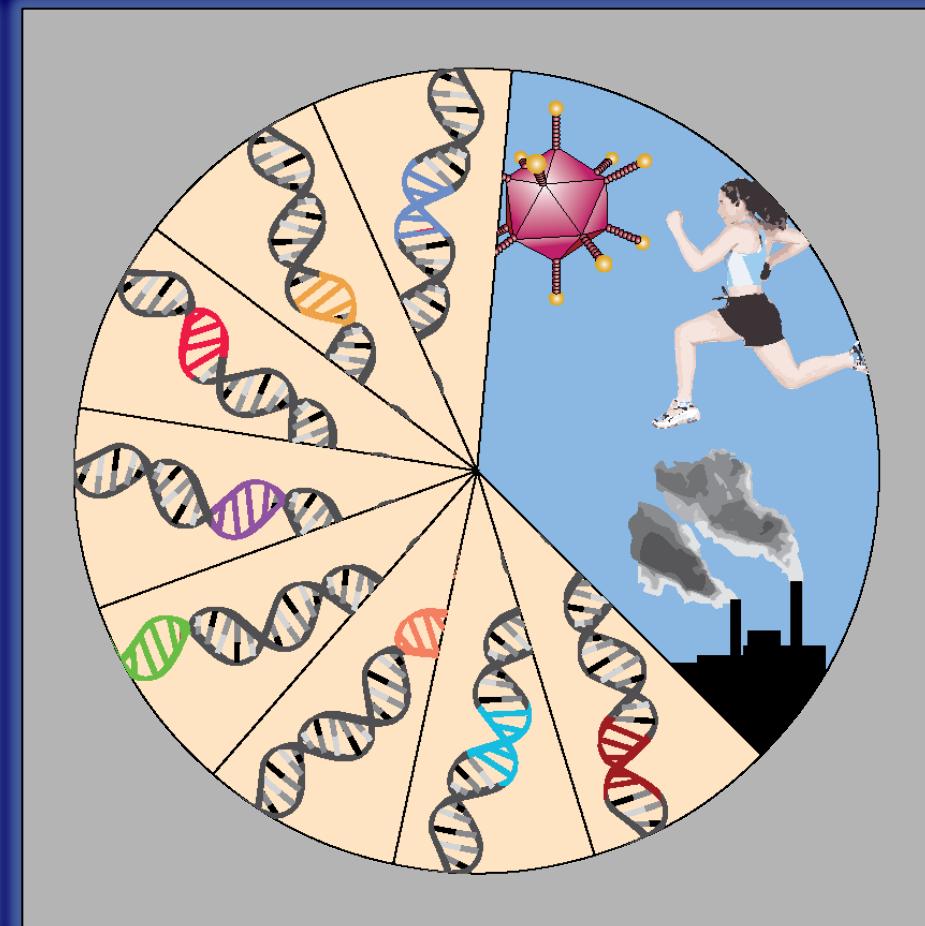
- ~6B nucleotides
- ~3-5M single-nucleotide variants
 - ~150K not in databases
 - ~60 not in either parent
- ~10-20K structural variants
- ~100 'disruptive' variants in genes
- ~20 completely inactivated genes (both copies)

Genomic Architecture of Genetic Diseases



Rare, Simple, Monogenic,
Mendelian...

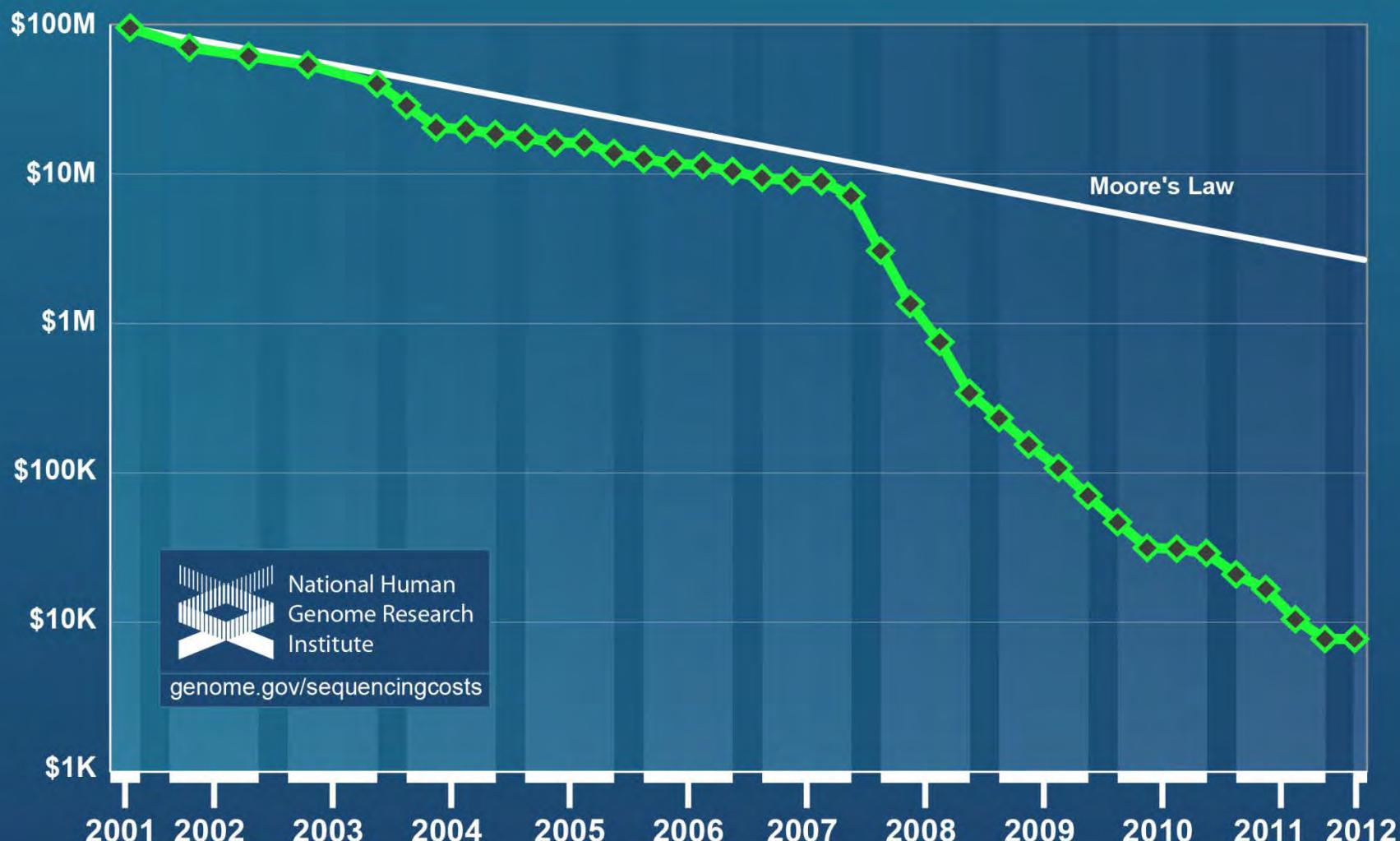
Mostly *Coding* Mutations



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

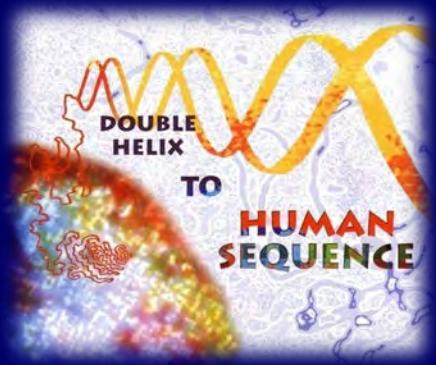
Mostly *Non-Coding* Mutations

Cost per Sequenced Human Genome

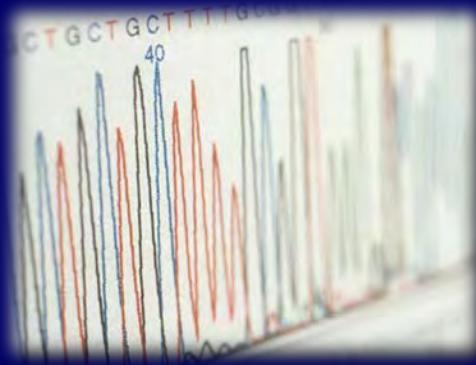


Sequencing a Human Genome

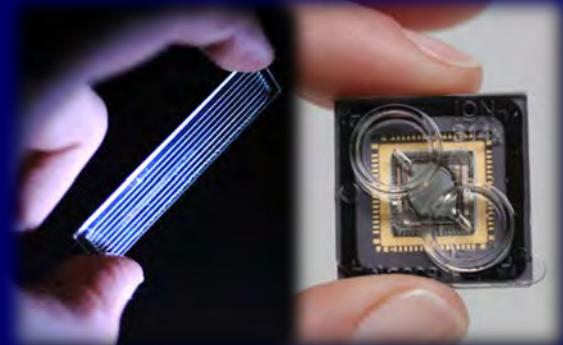
**HGP
(1st Sequence)**



Immediate Post-HGP



Today



~6-8 years

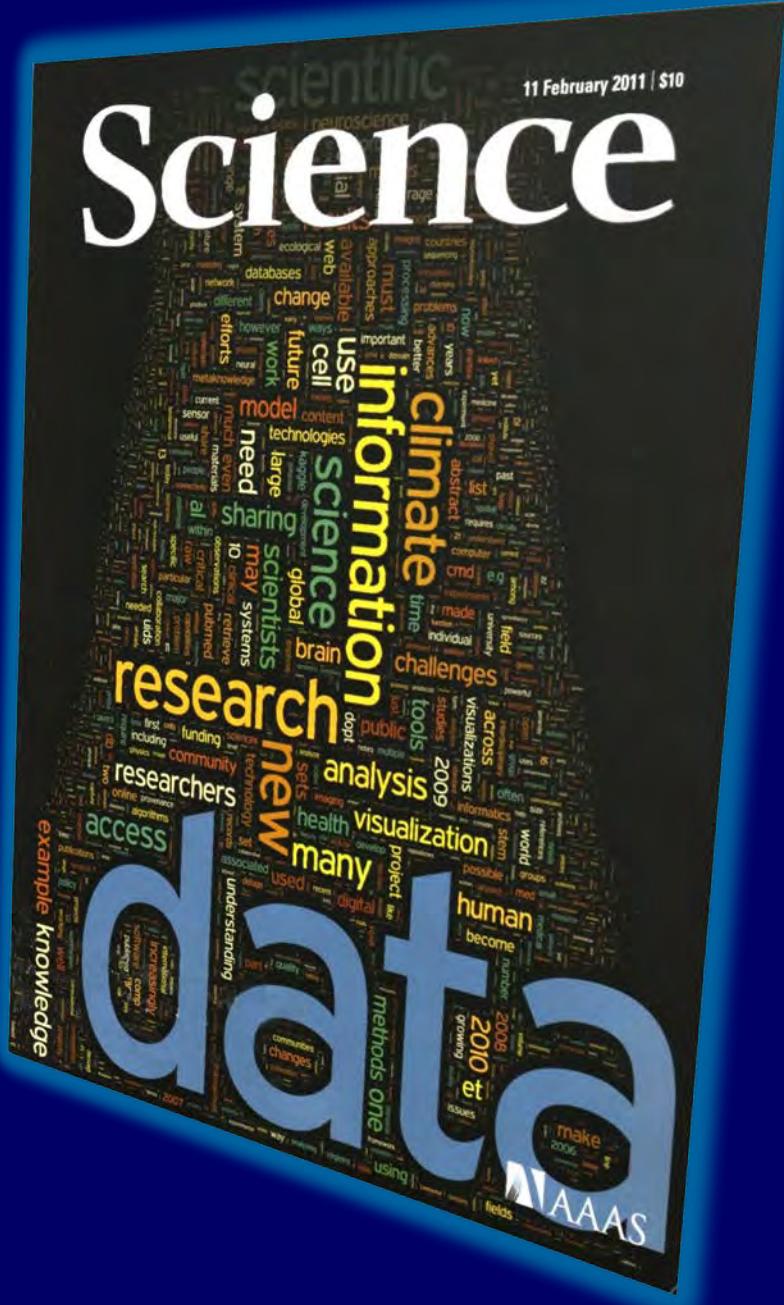
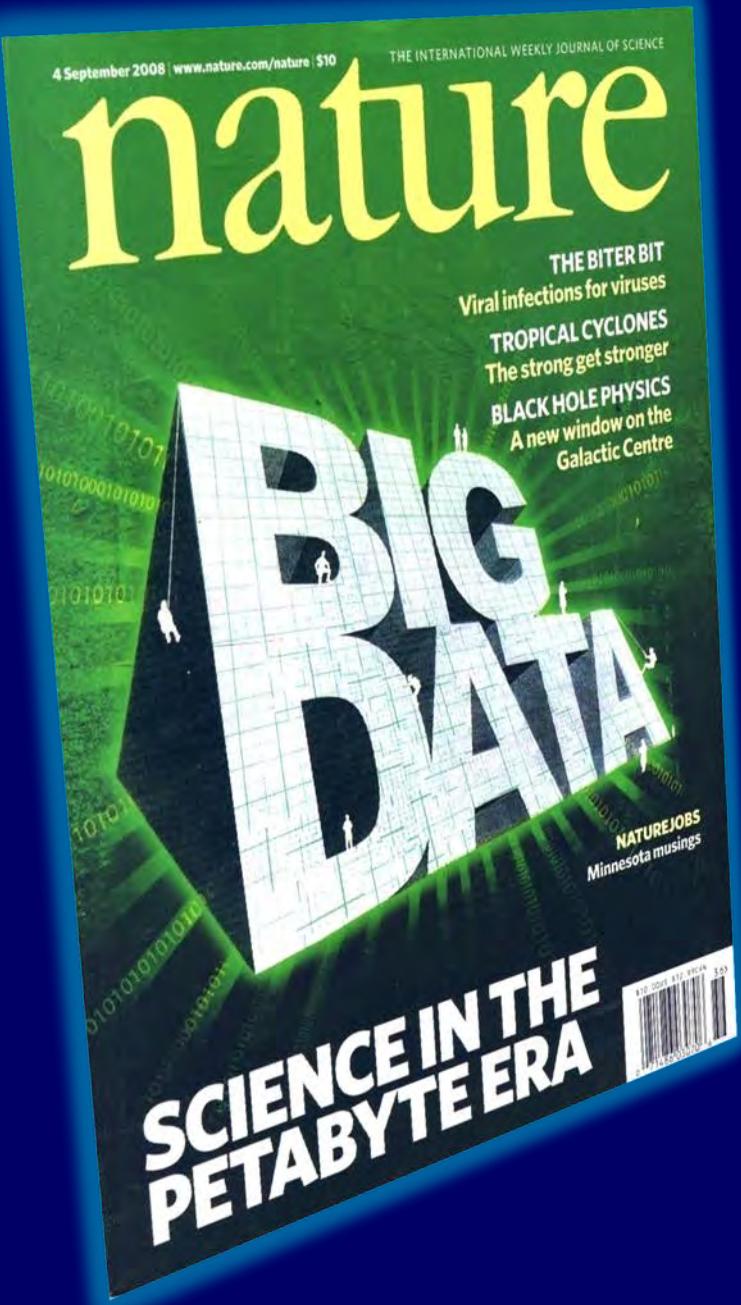
~3-4 months

~2-3 days

~\$1B

~\$10-50M

~\$4-6K



Genomics and the Economy



The Impact of Genomics on the U.S. Economy

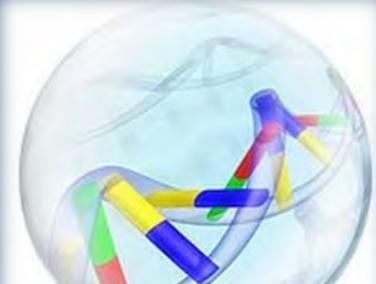
*How a federal research investment has
changed medical science, improved public health,
boosted the U.S. economy and created American jobs*

Prepared by Battelle Technology Partnership Practice
for United for Medical Research (UMR)

- From 1988-2012, genomic activities generated \$965 billion in economic output
- In 2012, genomic activities generated \$65 billion in economic output
- Genomics has a remarkable demonstrated return-on-investment

The Future

NEXT EXIT 

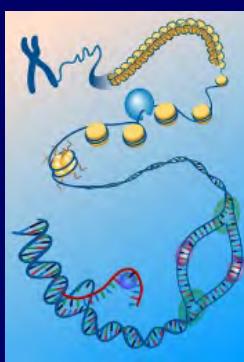


Genomic Accomplishments Across Domains

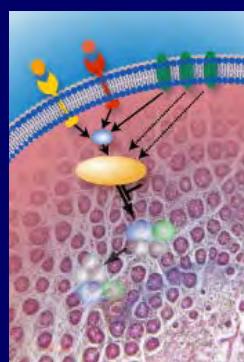
Understanding the Structure of Genomes



Understanding the Biology of Genomes



Understanding the Biology of Disease



Advancing the Science of Medicine



Improving the Effectiveness of Healthcare

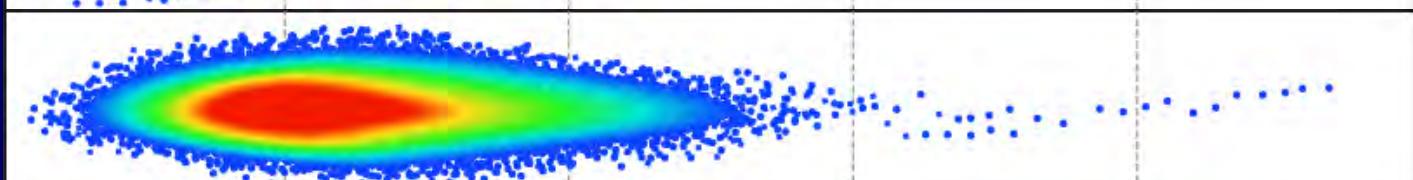


1990-2003

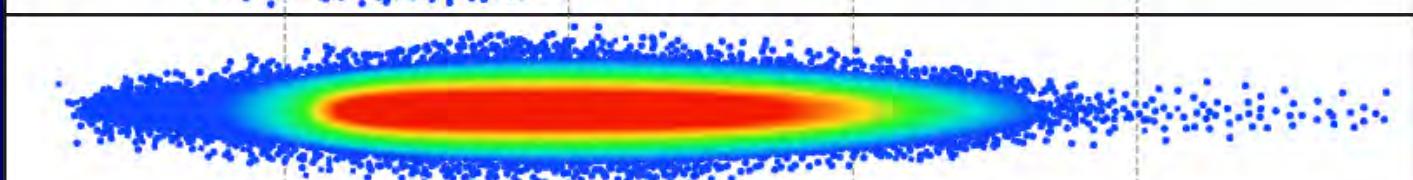
Human Genome Project



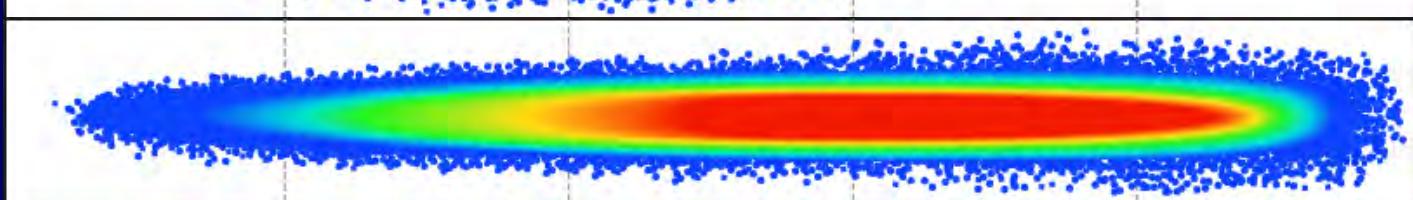
2004-2010



2011-2020

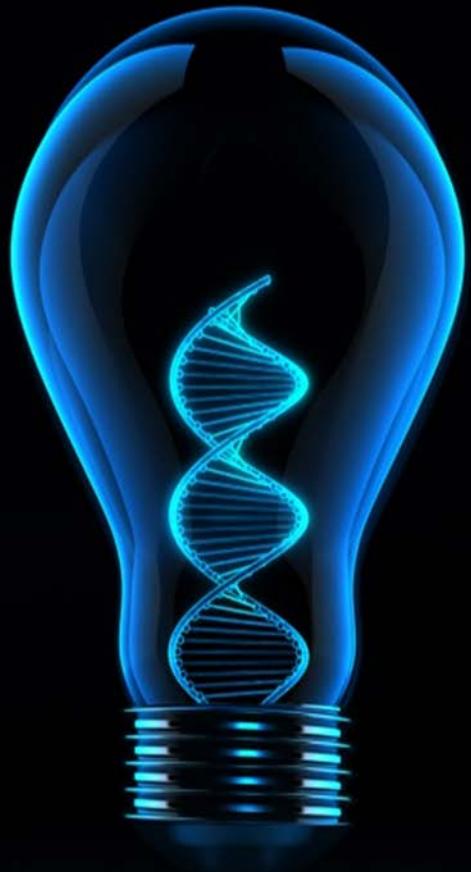
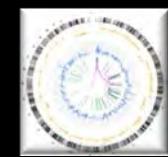


Beyond 2020



'Hot Areas' in Genomic Medicine

Cancer Genomics



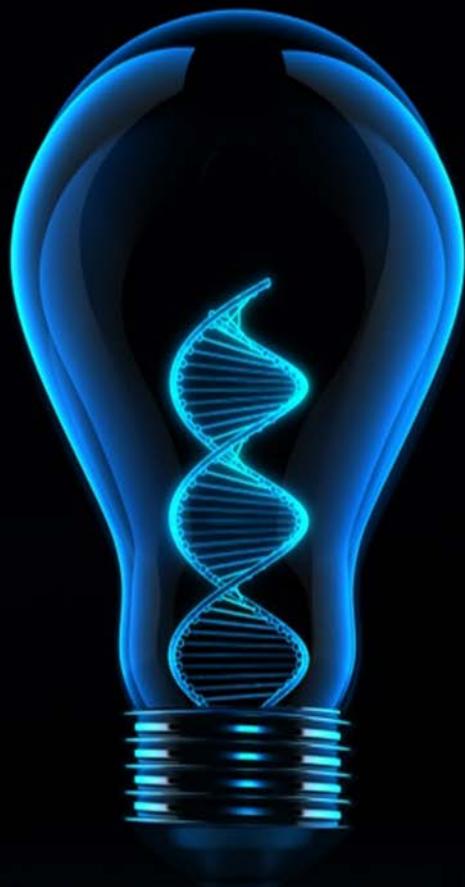
Genome Sequencing: Cancer



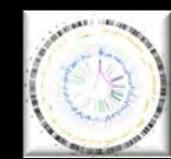
The screenshot shows the homepage of The Cancer Genome Atlas (TCGA) website. At the top, there is a red curved arrow pointing from the left towards the page. The header includes the TCGA logo and the tagline "Understanding genomics to improve cancer care". A search bar is located in the top right corner. Below the header, a navigation menu offers links to Home, About Cancer Genomics, Cancers Selected for Study, Research Highlights, and Publications. The main content area features a photograph of two researchers examining a computer screen displaying genetic data. To the right, a section titled "News Releases and Announcements" provides information about completed characterization studies for acute myeloid leukemia (AML) and endometrial cancer, with a "Learn More" link. At the bottom, there are four buttons: "Two New TCGA Publications" (with a thumbnail of a publication cover), "Case Study" (with a thumbnail of a man's face), "Cancers Selected for Study" (with a thumbnail of a colorful circular diagram), and "About TCGA" (with a thumbnail of a circular diagram).

Cancer Genomics

'Hot Areas' in Genomic Medicine



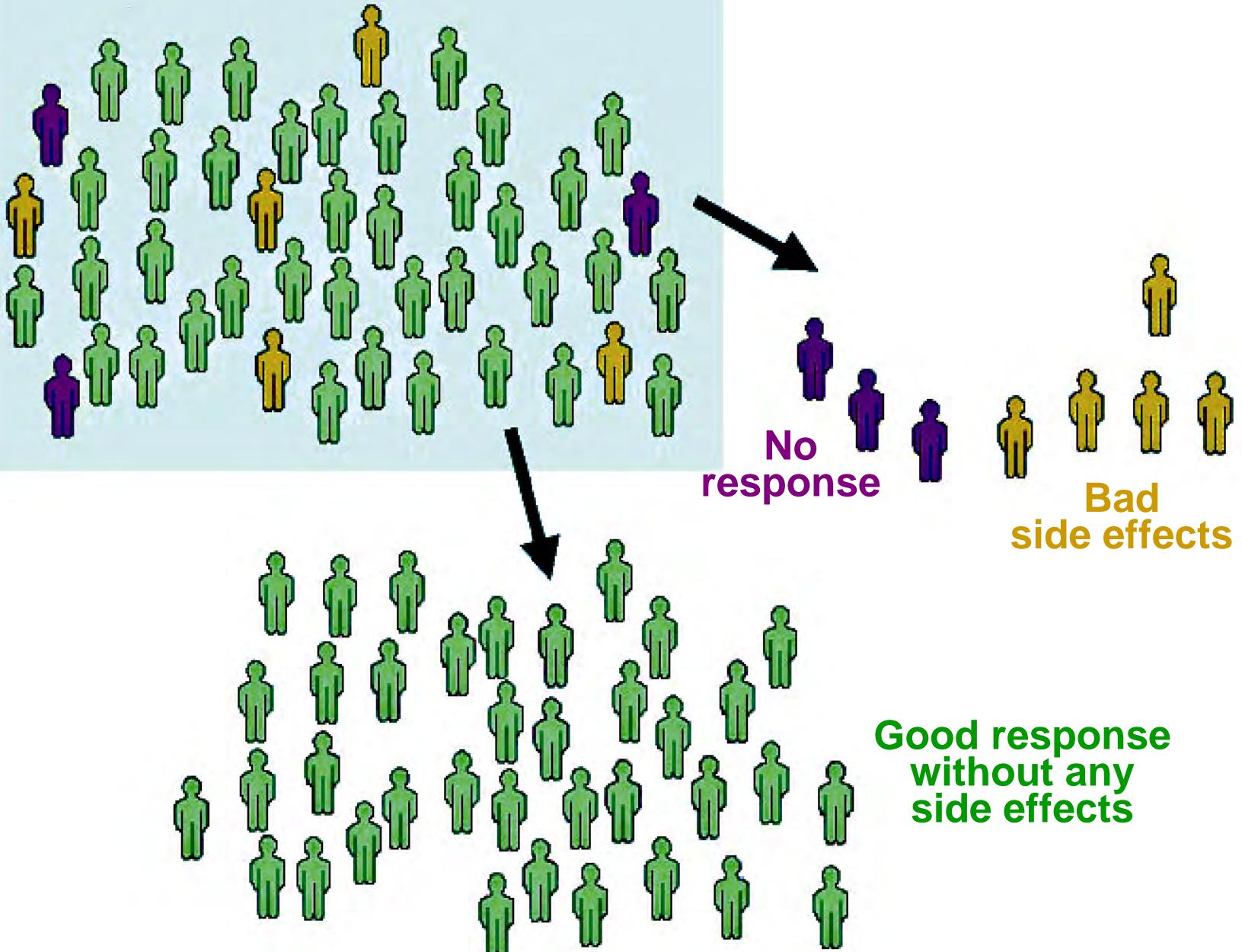
Cancer Genomics



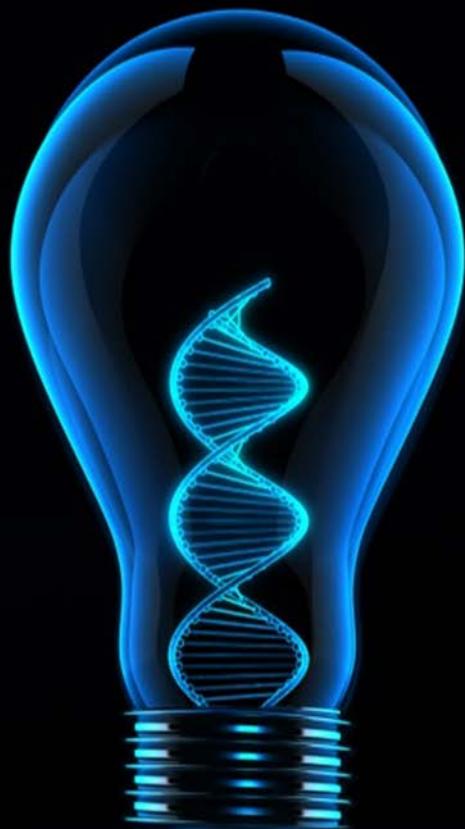
Pharmacogenomics



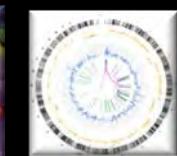
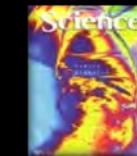
All patients with same disease



'Hot Areas' in Genomic Medicine



Cancer Genomics



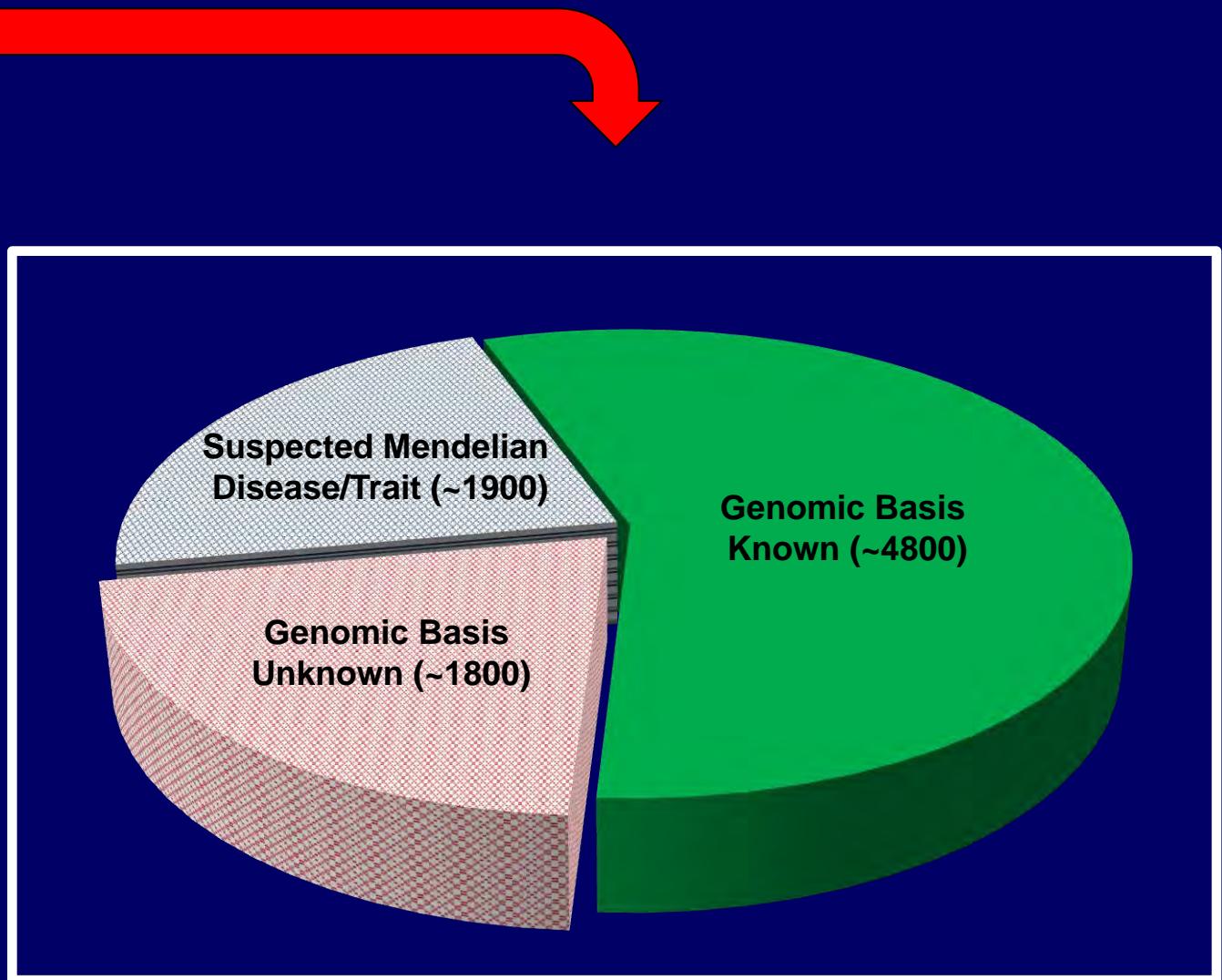
Pharmacogenomics



Rare & Common
Genetic Diseases



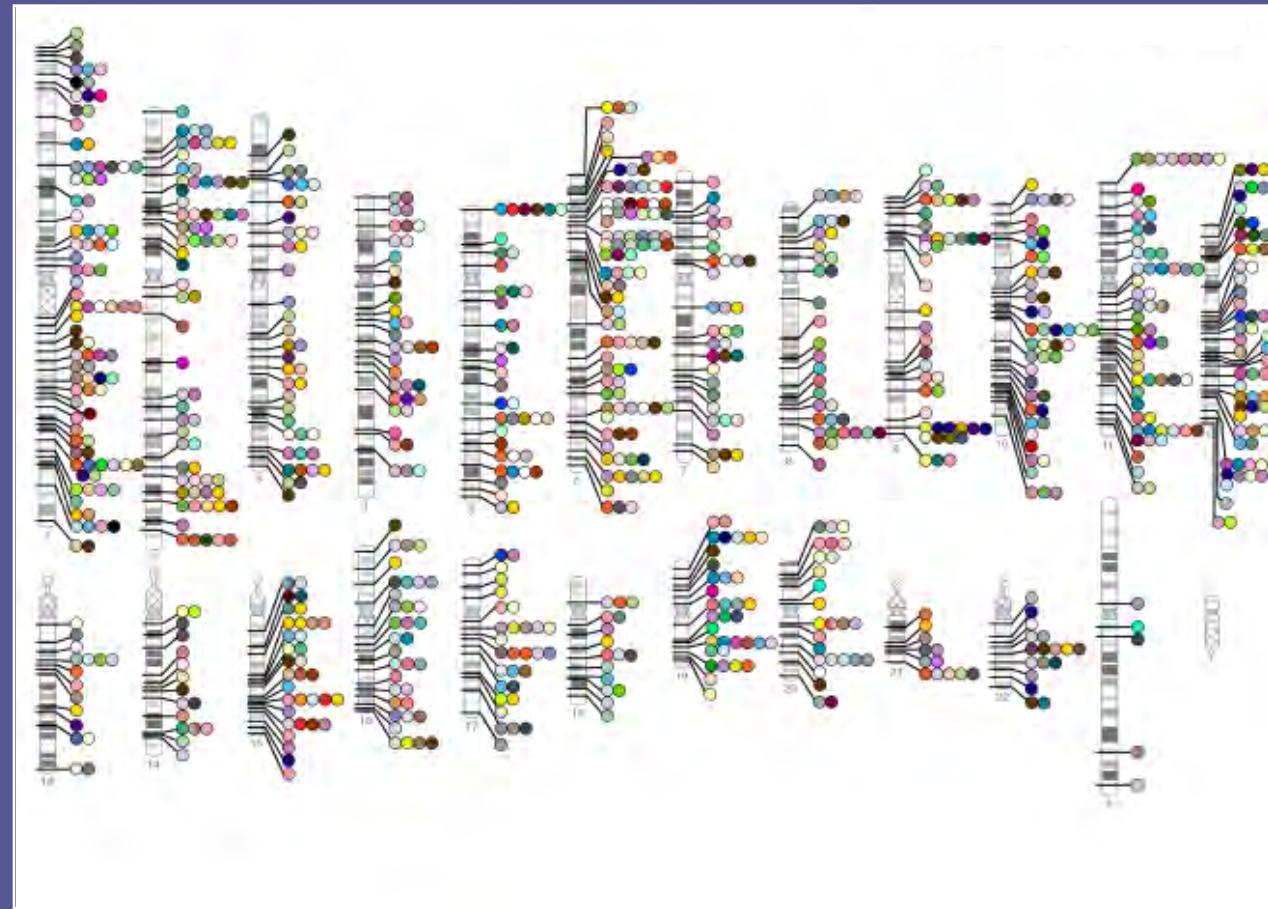
Genome Sequencing: Rare Diseases



Mendelian Diseases/Traits

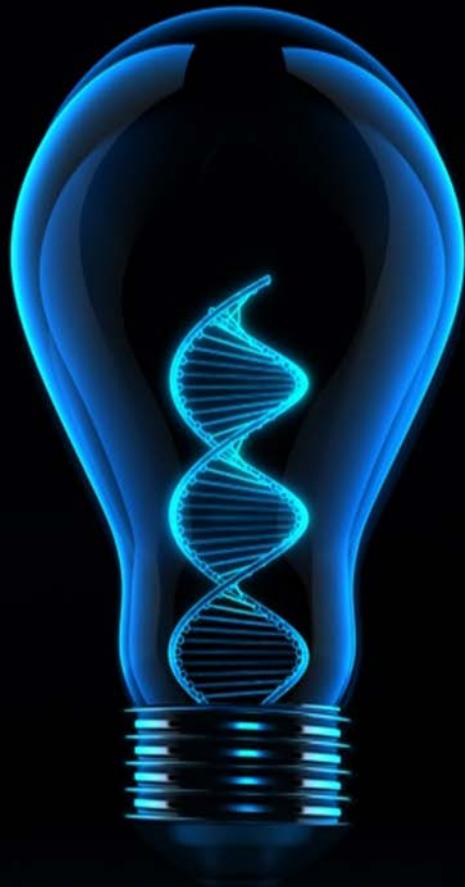


Genome Sequencing: Common Diseases

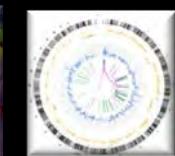


Complex Diseases/Traits

'Hot Areas' in Genomic Medicine



Cancer Genomics



Pharmacogenomics



Rare & Common
Genetic Diseases



Prenatal & Newborn
Genomic Analysis



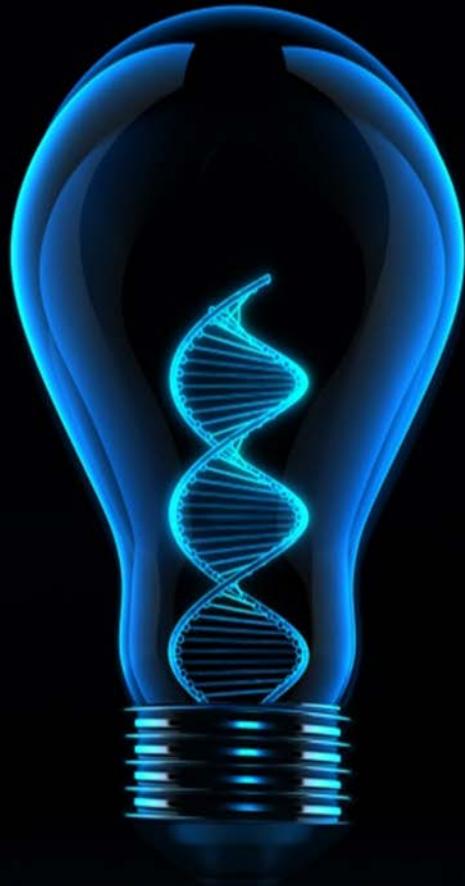
Clinical Genomics
Information Systems



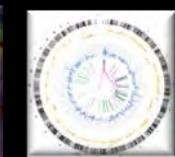
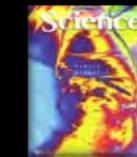
Clinical Genomic Information Systems



'Hot Areas' in Genomic Medicine



Cancer Genomics



Pharmacogenomics



Rare & Common
Genetic Diseases



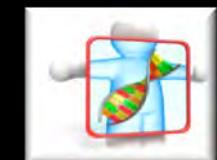
Prenatal & Newborn
Genomic Analysis



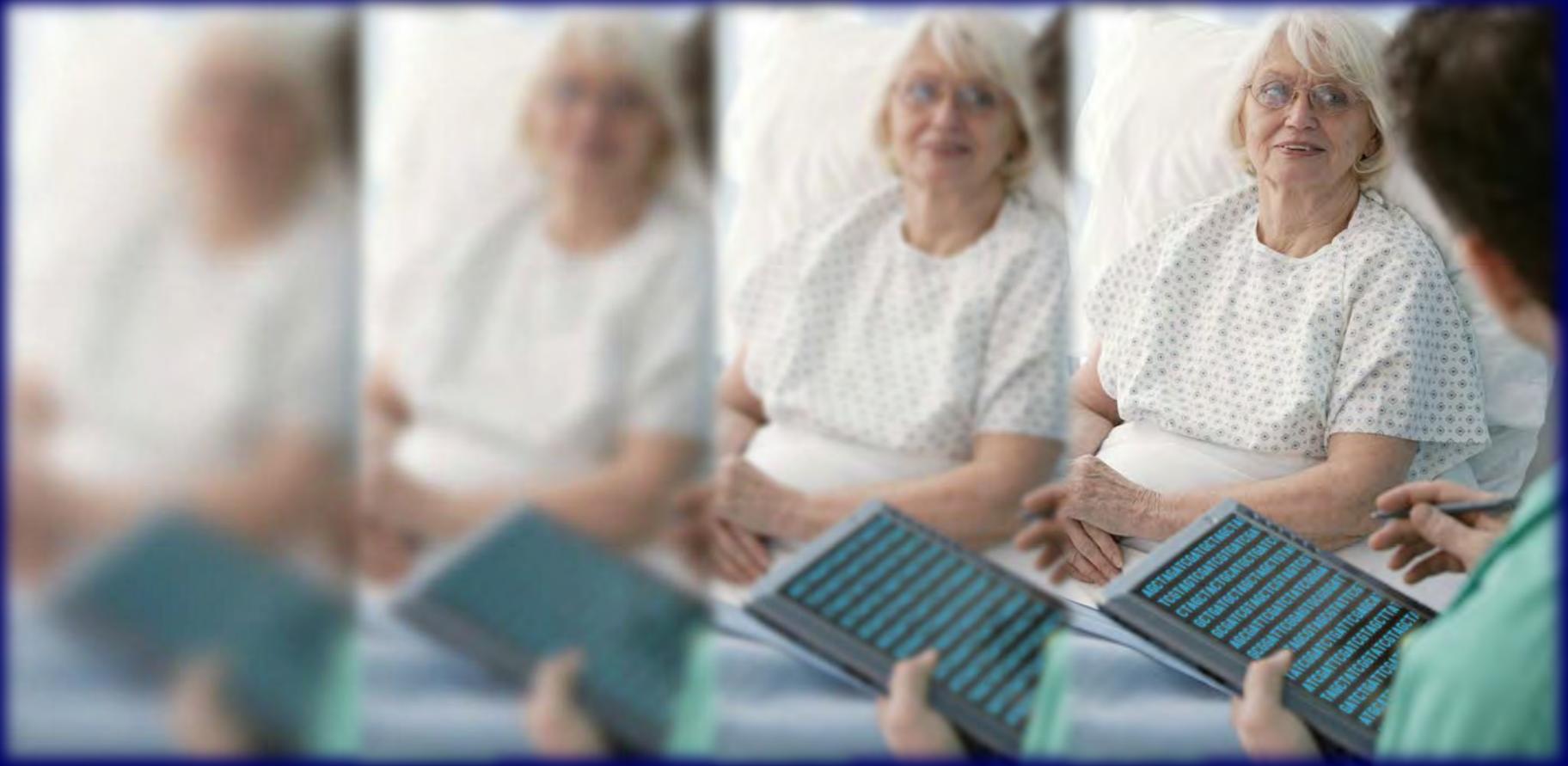
Clinical Genomics
Information Systems



Genomic Medicine
'Test Drive' Programs



Bringing Genomic Medicine into Focus



~1990

~2003

~2011

~2020

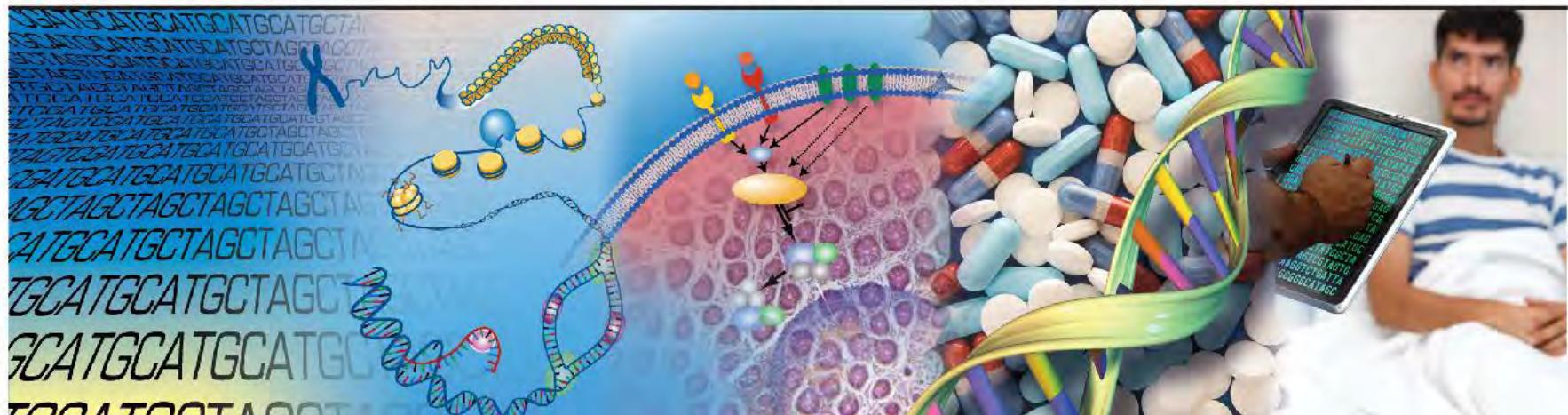
Understanding the Structure of Genomes

Understanding the Biology of Genomes

Understanding the Biology of Disease

Advancing the Science of Medicine

Improving the Effectiveness of Healthcare



**A pessimist sees the difficulty in every opportunity.
An optimist sees the opportunity in every difficulty.**

--Winston Churchill



NATIONAL HUMAN GENOME
RESEARCH INSTITUTE



*Advancing human health
through genomics research*