# The National Cancer Institute: Leading Cancer Research in 2021 and Beyond

Norman E. Sharpless, M.D.

94th Meeting of the National Advisory Council for Human Genome Research (Virtual)
September 13, 2021

@NCIDirector @TheNCI



# NATIONAL CANCER ACT MINISERIES

#### Cancer innovation

2021 marks the 50th anniversary of the National Cancer Act of 1971 — legislation that intended to end the "war on cancer" by expanding funding and authorities to the National Cancer Institute. Half a century since the act was signed into law, NCI and the biomedical research community hade leaps and bounds in cancer research, clinical trials, diagnostics and treatment. As NCI and the country continue to fight the war on cance today, GovernmentCIO Media & Research is collaborating with the institute to bring you a six-part HealthCast miniseries to commemorate the National Cancer Act's impact. We're releasing episodes every other month throughout 2021 leading to the act's official anniversary in December



### 50 Years of Cancer: Progress in Overcoming Health Disparities

Making the fight against cancer more equitable requires diversifying cancer research, clinical trials and treatments. Jun 30, 2021



#### 50 Years of Cancer: The Road to Better Trea Diagnostics

Clinical trials, innovation in research, and technology have improved cancel in public health.

Apr 29, 2021





How the National Cancer Act of 1971 Revolutionized Cancer Care and What Lies Ahead: Celebrating 50 Years of Cancer Progress: Conversations With Seven Cancer Care Experts ascopost.com/issues/may-25-... #cancercare #NothingWillStopUs

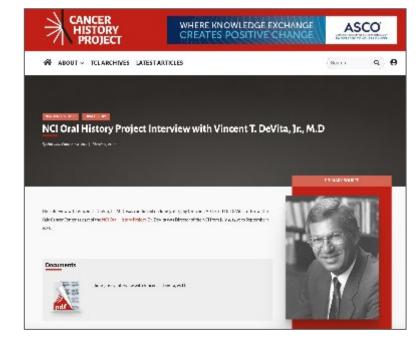




#### Ned Sharpless, MD 📀 @NCIDirector · Jun 16

Looking forward to participating in a dialogue on the great progress in #CancerResearch since the #NationalCancerAct of 1971, and the next 50 years, with @CNN's Andrew Kaczynski (@KFILE) tomorrow at 11 am ET on #WashingtonPostLive. #NothingWillStopUs





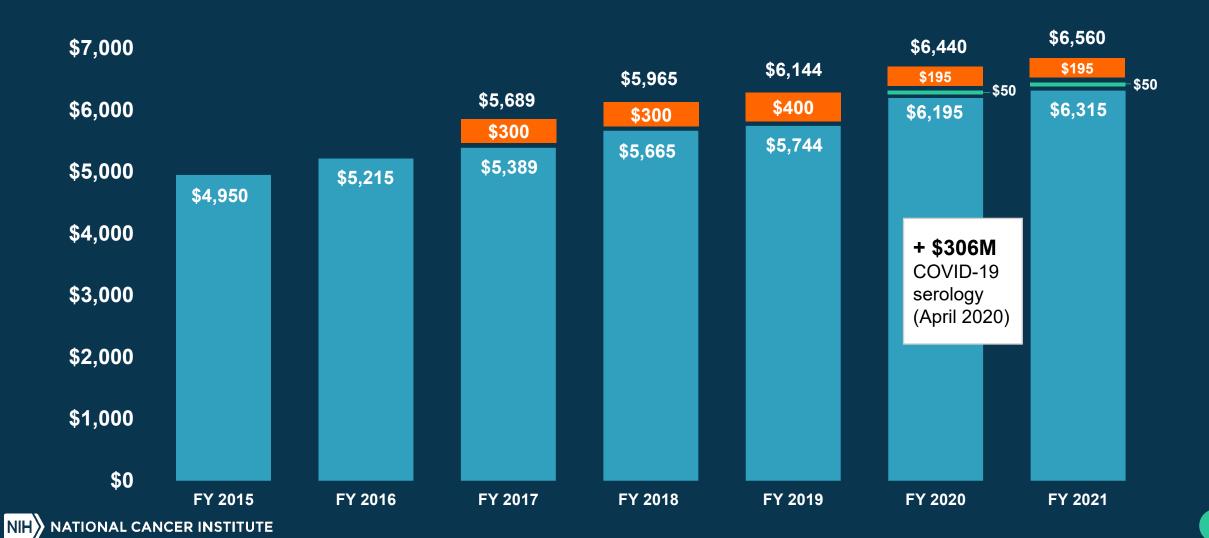
# Annual Plan & Budget Proposal for Fiscal Year 2023





# NCI Appropriations FY 2015 – 2021 (in millions)

### 21st Century Cures Act - orange Childhood Cancer Initiative - green

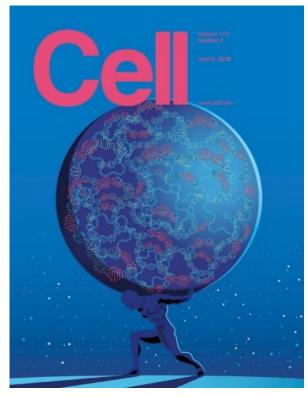


### **December 13, 2005**



TCGA Launch at the National Press Club

### **April 3, 2018**



Publication of the Pan-Cancer Atlas

# TCGA by the numbers



To put this into perspective, **1 petabyte** of data is equal to

212,000 DVDs

TCGA data describes

DIFFERENT RARE CANCERS

...based on paired tumor and normal tissue sets collected from

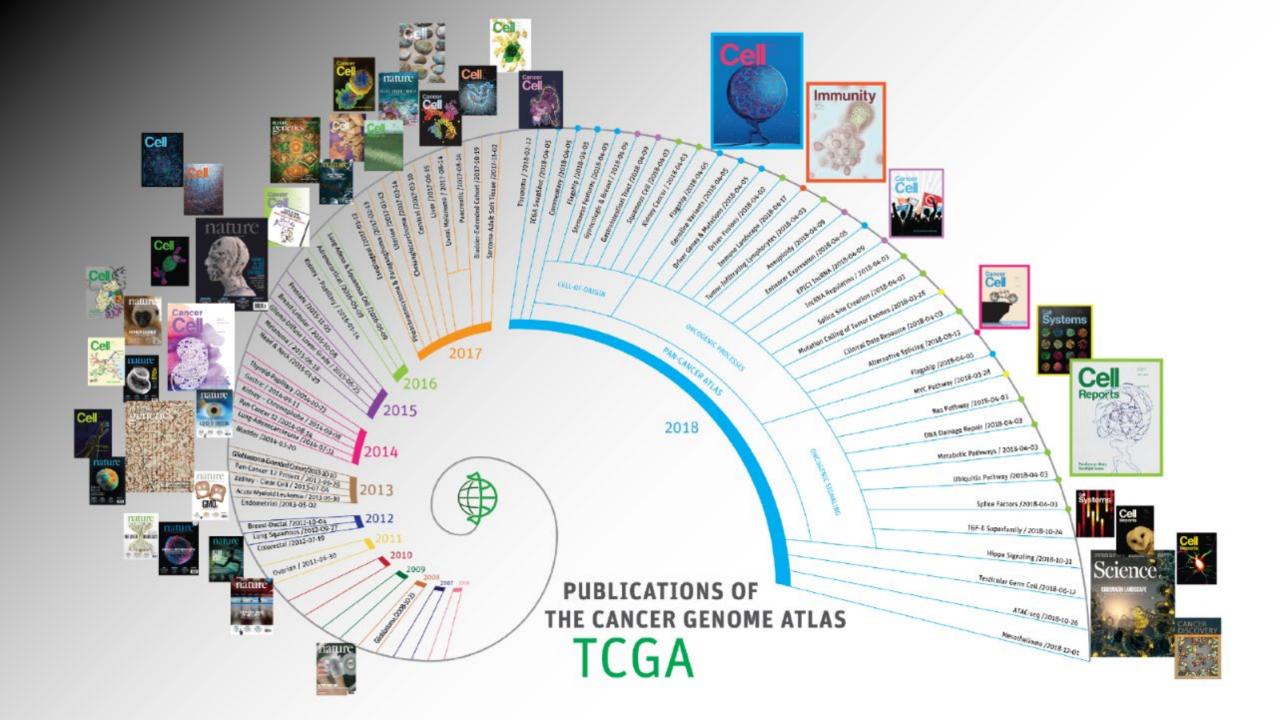


...using

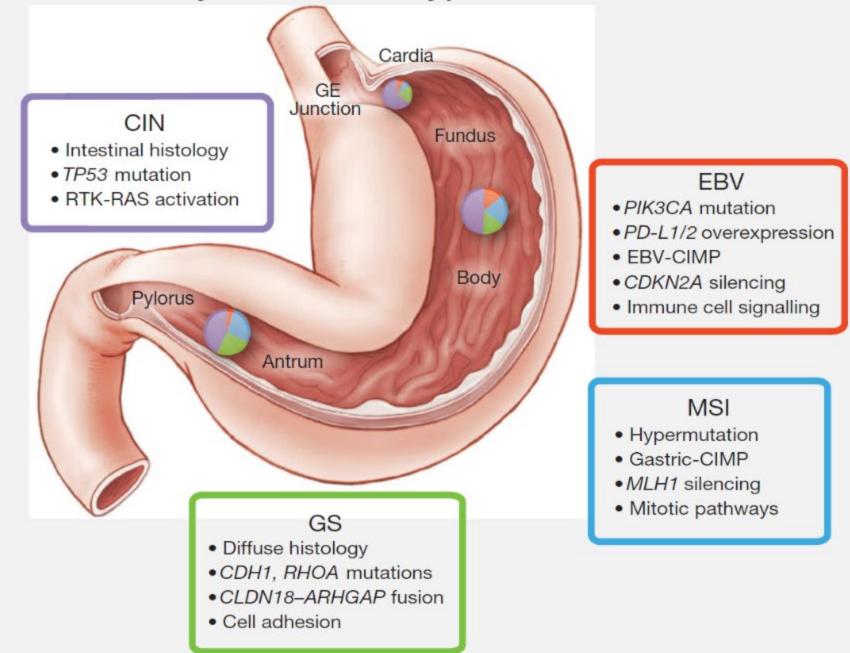
DIFFERENT DATA TYPES



...including



### Four Molecularly Distinct Subtypes of Gastric Cancer



TCGA. Nature. 2014 513:202-9.

# JEAN C. ZENKLUSEN, CAROLYN HUTTER and the Cancer Genome Atlas Team

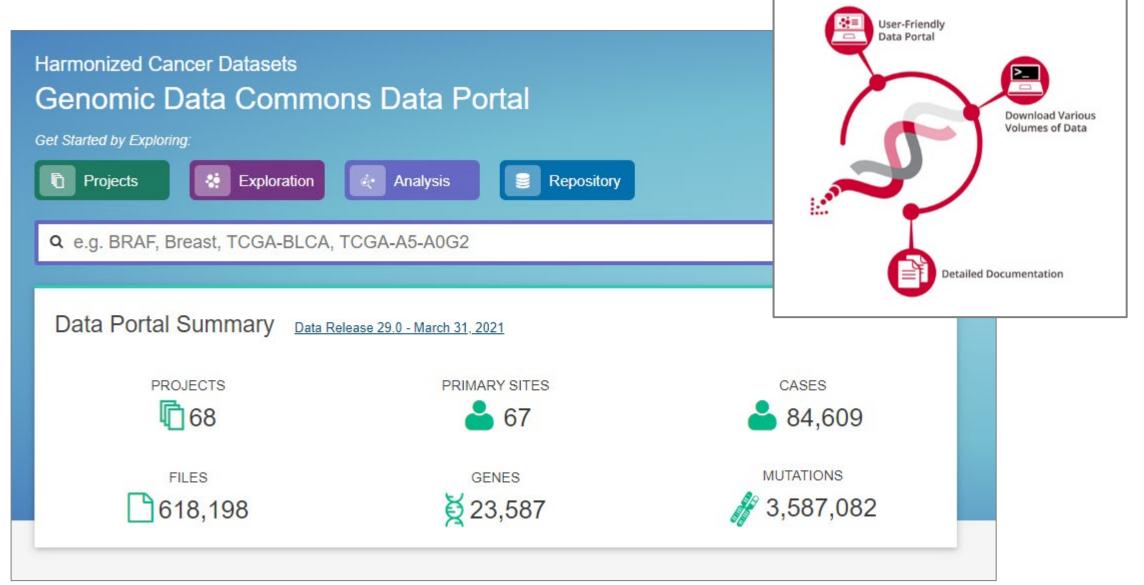
2015 WINNER
PEOPLE'S CHOICE AWARD

Mapped thousands of gene sequences for more than thirty types of cancer, advancing precision medicine in the diagnosis, treatment and prevention of these deadly diseases.



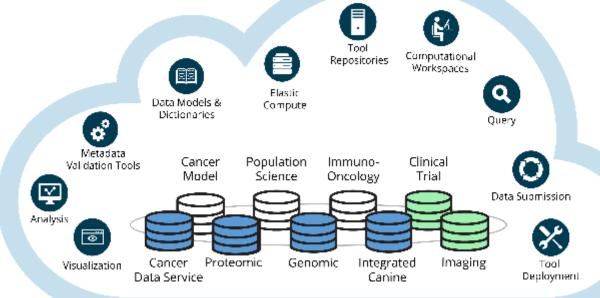


### **Genomic Data Commons Data Portal**



# NCI Cancer Research Data Commons

Cloud-based
data science infrastructure
that provides secure access to a
large, comprehensive, and
expanding collection of cancer
research data



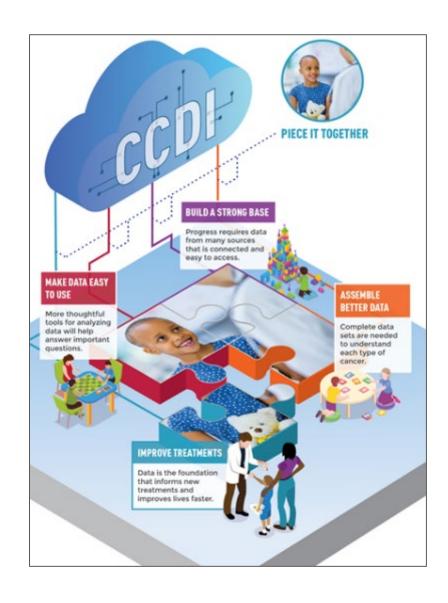
#### **Authentication & Authorization**



#### Data Contributors & Consumers







### **Childhood Cancer Data Initiative**

- Gather data from every child, adolescent, and young adult diagnosed with a pediatric cancer, regardless of where they receive their care
- 2 Create a national strategy of appropriate clinical and molecular characterization to speed diagnosis and inform treatment for all types of pediatric cancers
- Develop a platform and tools to bring together clinical care and research data that will improve preventive measures, treatment, quality of life, and survivorship for pediatric cancers

Childhood Cancer
Data Platform

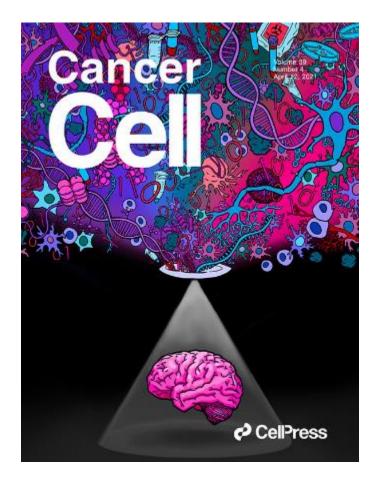
National Childhood
Cancer Cohort

Childhood
Molecular
Characterization
Protocol



# Proteogenomic and metabolomic characterization of human glioblastoma

- Phosphorylated PTPN11 and PLCG1 represent a signaling hub in RTK-altered tumors
- Four immune GBM subtypes exist, characterized by distinct immune cell populations
- Mesenchymal subtype EMT signature is specific to tumor cells, but not to stroma
- Histone H2B acetylation is enriched in classical GBMs with low macrophage content



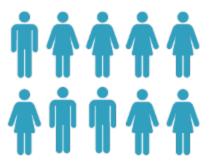


## SHERLOCK LUNG

Tracking Lung Cancer Mutational Processes in Never Smokers

### **232** Patients

General Population Never Smokers European Descent



### **Whole Genome Sequencing**

Tumor Tissue 85x Blood/Normal Tissue 32x



### **How Tumors Were Evaluated**

- Q Genomic Landscape
- Q Mutational Signatures
- Q Evolutionary History
- Molecular Subtypes
- Clinical Outcome



### SHERLOCK LUNG

### **Findings**

Burden of Genomic Alterations Impacting Survival

### Identification of Genomic Subtypes

f

Forte (20%)

Smoker Like Features (WGD)

mf

*Mezzo-forte* (30%)

APOBEC Signature; SBS18

**EGFR** Mutations

p

Latency 1

Piano-Adenos (34%)

Stem Cell Like Features

p

*Piano*-Carcinoids (14%)

Alkylating Signatures; SBS8

Stem Cell Like Features

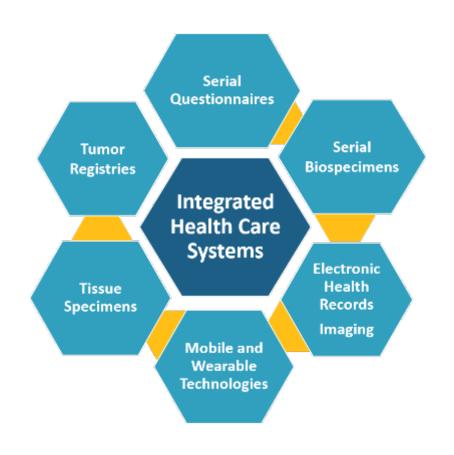
2 1

Germline AR; UBA1; NKX2-1

MYC; HLA LOH; MCL1; RBM10

EGFR; UBA1; ERBB2; ARID1A

### **Connect for Cancer Prevention Study**



- Prospective cohort of 200,000 adults in the United States
- Designed to further investigate the etiology of cancer and its outcomes, which may inform new approaches in precision prevention and early detection
- Will capitalize on research innovations to advance the field of cancer epidemiology and prevention including:
  - New technologies for exposure assessment
  - Large-scale analyses of the genome, epigenome, transcriptome, proteome, metabolome, microbiome
  - Molecular profiling of tumors and precursor lesions





>300,000 Breast Cancer **Patients** 



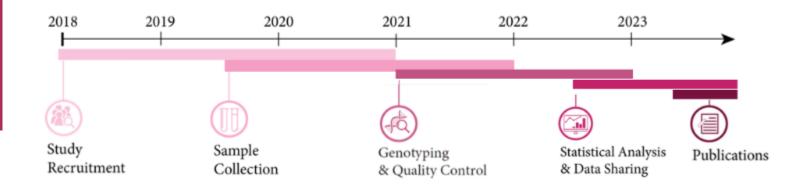
>300,000 Controls

International and multiancestry



### Multi-ancestry genome-wide association study (GWAS) to:

- Discover variants for breast cancer risk overall and by subtype
- Develop multi-ancestry polygenic risk scores for personalized risk assessment
- Discover variants for breast cancer survival, pharmacogenomics, and second cancers



### **Novel Uses of Sequencing**

Mutational signatures to determine etiology





### NCI Symposium on

### **MUTATION SIGNATURES AND CANCER**

December 2-3, 2021

### https://events.cancer.gov/nci/mutationsignatures

### *Invited Speakers:*

Ludmil Alexandrov, Ph.D., University of California, San Diego Maria Teresa Landi, M.D., Ph.D., National Cancer Institute Peter Campbell, Ph.D., Wellcome Trust Sanger Institute Phil Jones, M.D., Ph.D., University of Cambridge Elli Papaemmanuil, Ph.D., Memorial Sloan Kettering Gad Getz, Ph.D., Broad Institute Steven Rozen, Ph.D., Duke-NUS Allan Balmain, Ph.D., University of California, San Francisco Serena Nik-Zainal, M.D., Ph.D., University of Cambridge Reuben Harris, Ph.D., University of Minnesota

Adam Shlien, Ph.D., University of Toronto
Teresa Przytycka, Ph.D., National Center for Biotechnology
Information
David Wedge, Ph.D., University of Manchester
Peter Park, Ph.D., Harvard University
Joshua Campbell, Ph.D., Boston University
Ludmila Prokunina-Olsson, Ph.D., National Cancer Institute
Hannah Carter, Ph.D., University of California, San Diego
Nuria Lopez-Bigas, Ph.D., IRB Barcelona

### Division of Cancer Control and Population Sciences Leadership



Robert T. Croyle, Ph.D.

Director

NCI Division of Cancer Control and
Population Sciences

Retiring December 2021



Katrina Goddard, Ph.D.
Kaiser Permanente Center
for Health Research
Incoming DCCPS Director
pending final reviews.

# Discussion