



Advancing Innovation and Convergence in Cancer Research

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Health Sciences Director

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National Institutes of Health (NIH)



Northern California Blood Cancer Conference, Hyatt Regency- San Francisco
Feb 4, 2017



- **WHY** am i here?
- **WHO** i am & **WHAT** is cssi?
- **WHY data quality & sharing are IMPORTANT**
- **2016** at a glance
- **2017** moving the needle

04/09/2013

+

**Systems Biology and
the Impact on
Understanding
Cancer**

Systems Fiveology

Fredricka Hunter
Sanford Jeames
Kim Alexandre Powell
Lauren Wilkerson

Patient Mentor: Peg Ford

Scientific Mentor: Dr. Jerry Lee

AACR American Association
for Cancer Research



04/21/2015

CANCER COMPLEXITY



Work Group 4: Mary Kay Dauria, Leila Evangelista, Jeri Francoeur,
Michael Jones, Yoshiyuki Majima, Jeannine Walston
Mentor: Cynthia Ryan, Ph.D.
Scientific Mentor: Jerry Lee, Ph.D.



Work Group 4: Mary Kay Dauria, Leila Evangelista, Jeri Francoeur,
Michael Jones, Yoshiyuki Majima, Jeannine Walston
Mentor Cynthia Ryan
Scientific Mentor: Jerry Lee

04/20/2016



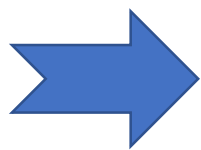
Mission

- Make a decade's worth of progress in **cancer prevention, diagnosis, treatment, and care** – ultimately to end cancer as we know it.

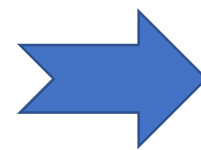




2001



2010



1 million healthy genomes

2015



2016

1,685,210

new cases of cancer in the U.S.

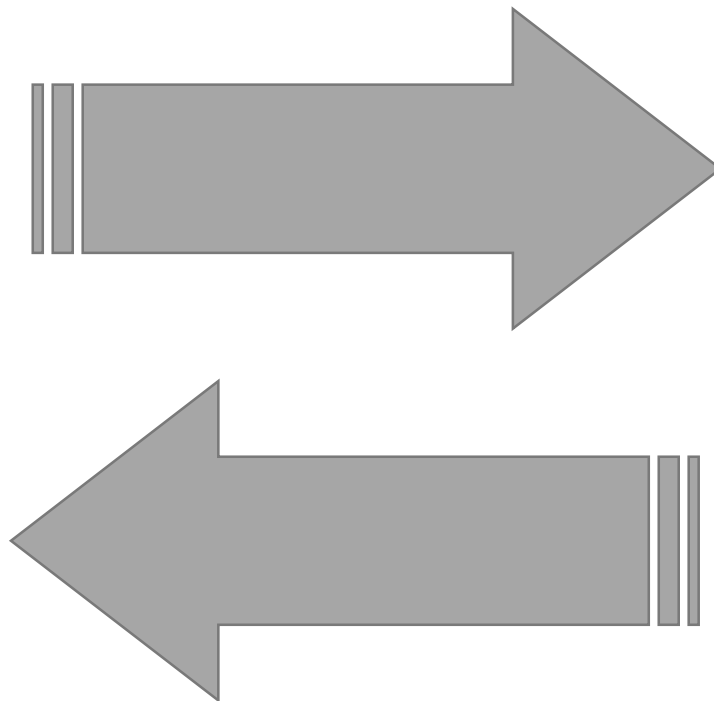
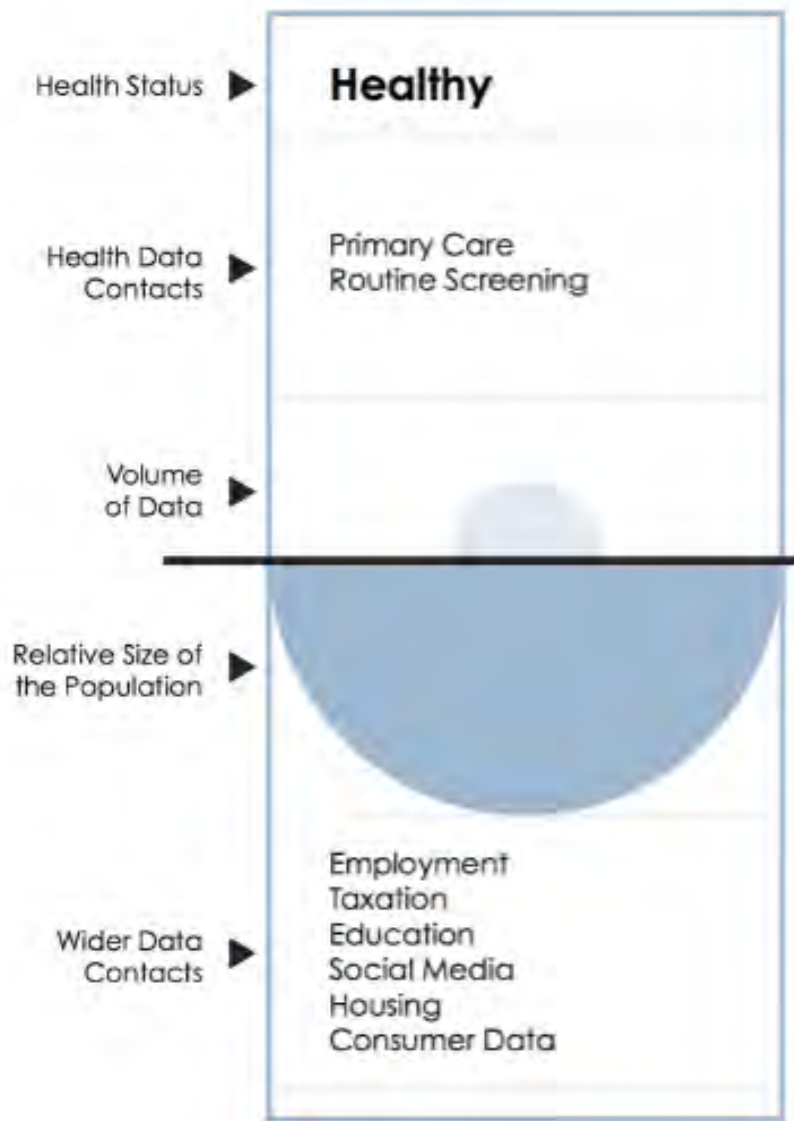
595,690

projected deaths due to cancer in the U.S.

2016

15,533,220

cancer survivors in the U.S.



Big Hopes for Big Data: Digital Information Focus Underpins Cancer Moonshot Goals

Andrew D. Smith | January 13, 2017



Daniel F. Hayes, MD

The passage of the 21st Century Cures Act¹ sparked unusually robust holiday celebrations throughout the cancer research community, which harbors high hopes for both the newly funded initiatives and future budget increases.

The act will provide the National Cancer Institute (NCI) an extra \$1.8 billion to fund the Cancer Moonshot, an effort to make 10 years of research gains in just 5 years by directing new money to 10 potentially transformative research areas

(**Chart, Table**). It also gives the FDA the ability to consider real-world clinical data in determining drug indications.

“...asked what tied the various aspects of the Moonshot together, he [Daniel Hayes] answered with a single word: **data**. Most of the specific research recommendations envision the **aggregation** of huge amounts of data and the **extraction** of important discoveries from those collection points...”

'Automated dermatologist' detects skin cancer with expert accuracy

By Susan Scutti, CNN

Updated 6:37 PM ET, Thu January 26, 2017



LETTER

nature

doi:10.1038/nature21056

Dermatologist-level classification of skin cancer with deep neural networks

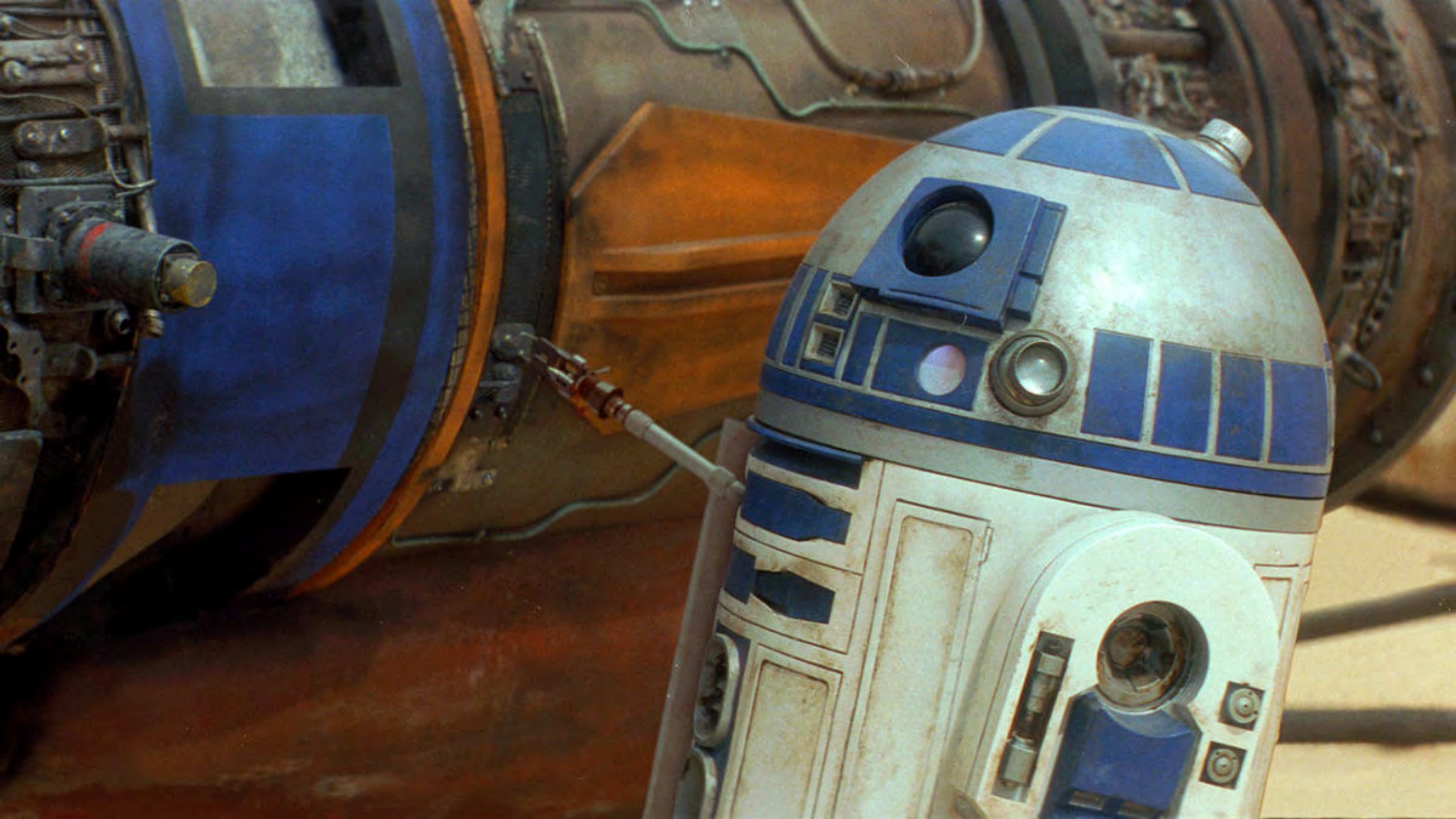
Andre Esteva^{1*}, Brett Kuprel^{1*}, Roberto A. Novoa^{2,3}, Justin Ko², Susan M. Swetter^{2,4}, Helen M. Blau⁵ & Sebastian Thrun⁶

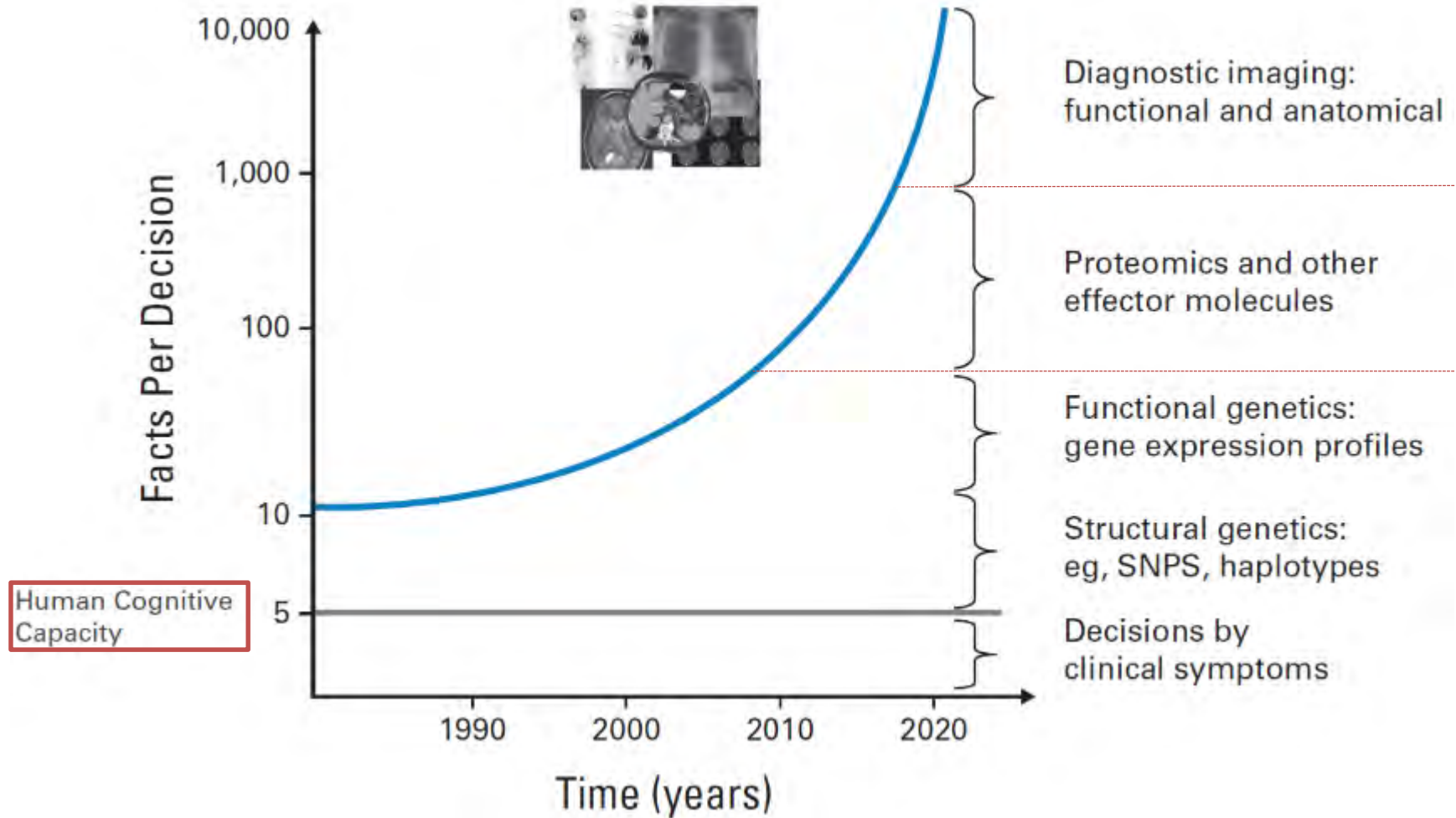
Skin cancer, the most common human malignancy¹⁻³, is primarily diagnosed visually, beginning with an initial clinical screening and followed potentially by dermoscopic analysis, a biopsy and histopathological examination. Automated classification of skin lesions using images is a challenging task owing to the fine-grained variability in the appearance of skin lesions. Deep convolutional neural networks (CNNs)^{4,5} show potential for general and highly variable tasks across many fine-grained object categories⁶⁻¹¹. Here we demonstrate classification of skin lesions using a single CNN, trained end-to-end from images directly, using only pixels and disease labels as inputs. We train a CNN using a dataset of 129,450 clinical images—two orders of magnitude larger than previous datasets¹²—consisting of 2,032 different diseases. We test its performance against 21 board-certified dermatologists on biopsy-proven clinical images with two critical binary classification

images (for example, smartphone images) exhibit variability in factors such as zoom, angle and lighting, making classification substantially more challenging^{23,24}. We overcome this challenge by using a data-driven approach—1.41 million pre-training and training images make classification robust to photographic variability. Many previous techniques require extensive preprocessing, lesion segmentation and extraction of domain-specific visual features before classification. By contrast, our system requires no hand-crafted features; it is trained end-to-end directly from image labels and raw pixels, with a single network for both photographic and dermoscopic images. The existing body of work uses small datasets of typically less than a thousand images of skin lesions^{16,18,19}, which, as a result, do not generalize well to new images. We demonstrate generalizable classification with a new dermatologist-labelled dataset of 129,450 clinical images, including 3,374 dermoscopy images.

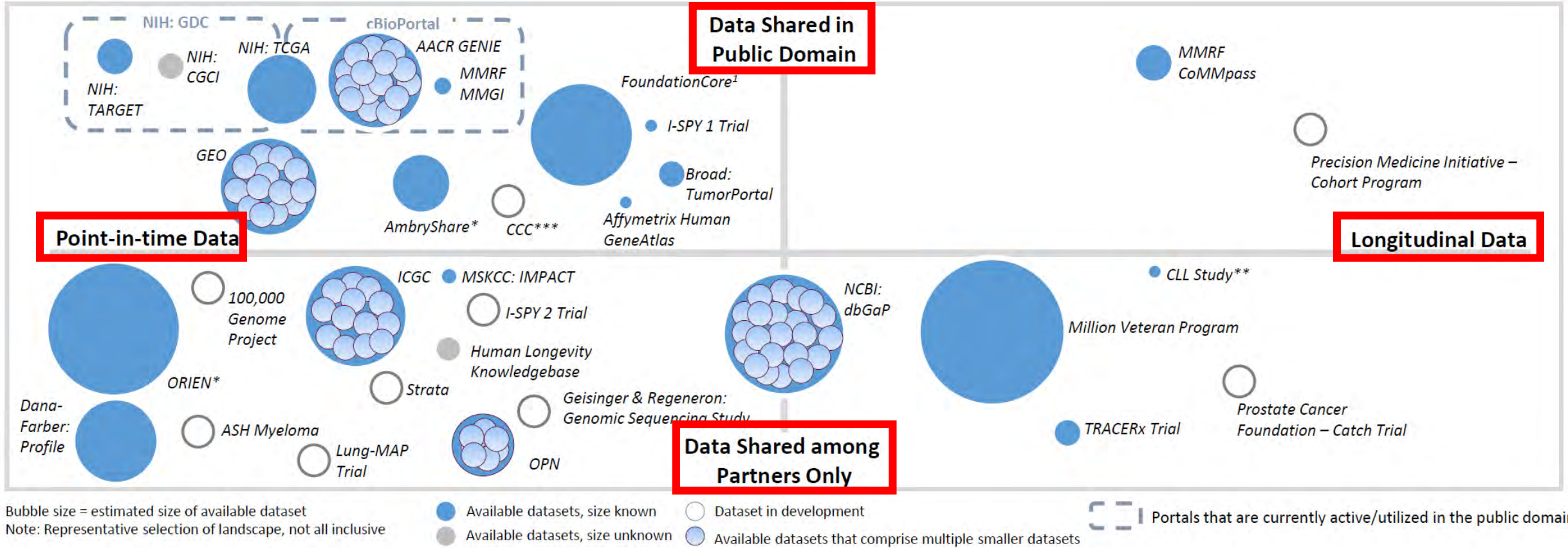
“...an advantage of machine learning is that it can be used even in cases where it is **infeasible** or **difficult** to write down explicit rules to **solve** a problem...”







“...to apply machine learning, a practitioner starts with a historical data set, which the practitioner divides into a *training set* and a *test set*...”



Opportunity exists to generate publicly available longitudinal data to drive understanding of genetic mutations and find Precision Medicine cures

*Datasets have potential to include longitudinal data in the future

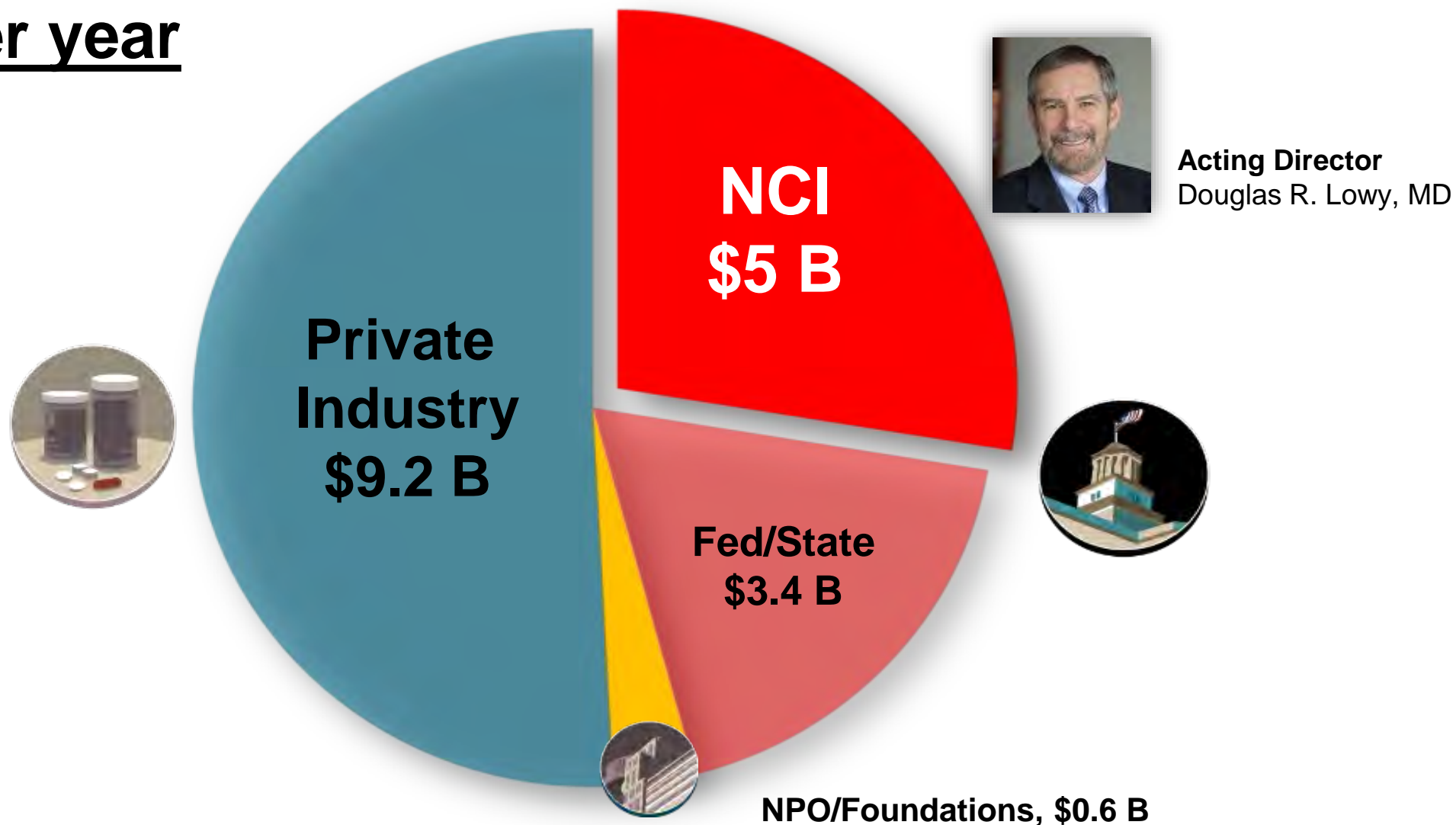
**Public/private information not available

***Serves as a portal also, has potential to include longitudinal data in the future

1. FoundationCore's pediatric cancer data has been made public

U.S. National Cancer Program: Stakeholders

~\$18 B per year



NCI Center for Strategic Scientific Initiatives (CSSI): Concept Shop



Director
Douglas R. Lowy, MD



Deputy Director
Jerry S.H. Lee, PhD

Mission

“...to create and uniquely implement exploratory programs focused on the development and integration of advanced technologies, trans-disciplinary approaches, infrastructures, and standards, to accelerate the creation and broad deployment of data, knowledge, and tools to empower the entire cancer research continuum in better understanding and leveraging knowledge of the cancer biology space for patient benefit...”



2003, 2007, 2011, 2013, 2014



2005, 2010, 2015



2008, 2013*



2011, 2014



2004, 2008, 2014



2005, 2008



2010

2006-2015: Building a historical data set

A Decade of Illuminating the Underlying Causes of
Primary Untreated Tumors



(12,000+ patient tumors and increasing)



THE LANCET

2002

MECHANISMS OF DISEASE

Mechanisms of disease

Use of proteomic patterns in serum to identify ovarian cancer

Emmanuel C Petricoin III, An M Aravamudan, Detlev A Holt, Peter A Levine, Willem A Grefenst, Seth M Steinberg, Gordon D Mills, Charles Slaughter, David A Fitzmaurice, Elise C Kohn, Denise A Liotta

Summary

Background New technologies for the detection of early-stage ovarian cancer are a priority need. Pathological changes within an organ might be reflected in molecules secreted in serum. We developed a peptide-based tool and used it to identify markers in patterns of serum that distinguish asymptomatic non-neoplastic disease within the ovary.

Methods Proteomic spectra were generated by mass spectrometry (surface-enhanced laser desorption and ionisation). A preliminary training set of spectra derived from analysis of serum from 50 unaffected women and 50 patients with ovarian cancer were analysed by an iterative searching algorithm that identified a proteomic pattern that consistently distinguished cancer from non-cancer. The discovered pattern was then used to classify 50 or independent set of 110 masked serum samples: 50 from women with ovarian cancer and 60 from unaffected women or those with non-malignant disorders.

Findings The algorithm identified a marker set that, used in the training set, completely segregated cancer from non-cancer. The discovered marker pattern correctly classified all 50 ovarian cancer cases in the masked set, including all 15 stage I cases. Of the 60 cases of non-malignant diseases, 83 were recognised as not cancer. This was a false-positive sensitivity of 100% (95% CI 99–100), specificity of 95% (87–100), and positive predictive value of 54% (34–84).

Introduction

Application of new technologies for detection of ovarian cancer could have an important effect on public health, but to achieve this goal, specific and sensitive molecular markers are essential. This goal is especially urgent in women who have a high risk of ovarian cancer due to family or personal history of cancer, and for women with a genetic predisposition to cancer that is abnormal alleles in predisposition genes such as *BRC1A1* and *BRC1A2*. There are no effective screening options for this population.

Ovarian cancer presents at a late clinical stage in more than 80% of patients, and is associated with a 5-year survival of 25% in this population. By contrast, the 5-year survival for patients with stage I ovarian cancer exceeds 90%, and most patients are cured of their disease by surgery alone.¹ Therefore, increasing the number of women diagnosed with stage I disease should have a direct effect on the mortality and economics of this cancer without the need to change surgical or chemotherapy approaches.

Cancer antigen 125 (CA125) is the most widely used biomarker for ovarian cancer.^{2,3} Although concentrations of CA125 are abnormal in about 50% of patients with advanced-stage disease, they are increased in only 50–60% of patients with stage I ovarian cancer.⁴ CA125 has a positive predictive value of less than 10% as a single marker, but the addition of ultrasound screening to CA125 measurement has improved the positive predictive value to about 20%.⁵

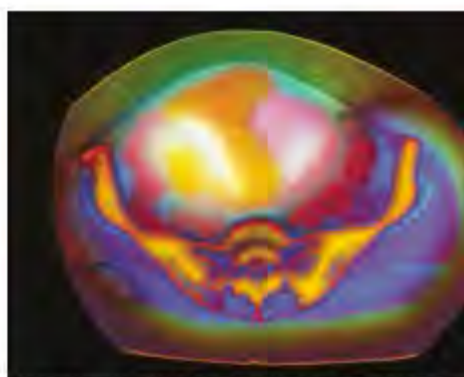
nature

2004

news feature

Running before we can walk?

Two years ago, a new proteomic test was heralded as the future of cancer diagnostics. But since then, doubts about its effectiveness have begun to grow. Erika Check reports.



Proteomic patterns in the blood serum of ovarian cancer patients (pink/yellow) before they reach the clinic.

Science does a single piece of research (through the US Congress to pass a resolution urging continued funding) to drive a new diagnostic test towards the clinic. But that's what happened in 2002, when *The Lancet* published a paper¹ claiming a breakthrough in the diagnosis of ovarian cancer.

The paper described the use of a peptide microarray to analyse the pattern of proteins present in samples of blood serum. On the basis of these patterns, the test detected all the patients with ovarian cancer in a set of 50 samples, and falsely identified just three healthy patients as suffering from the disease from a total of 65 control samples.

Most encouragingly, the technique seemed to work well on patients with early-stage disease — although the prospect of earlier diagnosis, which improves the chances of successful treatment, is the best current blood test, which relies on the detection of a single protein called CA125, misses at least half of patients in the earliest stages of the disease.

Early warning

critics warn that the grounds for optimism are shaky. The grounds for optimism are shaky. The grounds for optimism are shaky. The grounds for optimism are shaky.

Whether or not

Overalls work, we will learn from this experience what rules of evidence we might apply in the future to find useful results (page 497c).

was reported. They had recruited a set that Liotta and Petricoin's team published in August 2002. Soaje and Zou's study found numerous differences in protein patterns that distinguished between the cancer patients and the healthy ones.

The trouble, according to Soaje and Zou, is that these tools have little or no predictive value in real-world situations. The proteomic test relies on complex and intricate fields to separate the proteins given samples. Each protein is then given a number that represents the ratio of each and mass — called its *m/z* value. The

identifies proteins at 1 numbers to give a diagnostic and 2000 ions a second because the marked differences between cancer patients and healthy controls occurred for just with *m/z* values of less than 2000. Soaje and Zou's study

The New York Times

2004

February 3, 2004

New Cancer Test Stirs Hope and Concern

By ANDREW POLLACK

Jill Doimer's mother died in 2002 from ovarian cancer, detected too late to be effectively treated.

So Ms. Doimer is eagerly awaiting the introduction of a new test that holds the promise of detecting early-stage ovarian cancer far more accurately than any test available now, using only blood from a finger prick.

Not only does she plan to be tested, but an advocacy group she helped found, Ovarian Awareness of Kentucky, also intends to spread the word to women and doctors.

"If it's going to happen to me or anyone I know, I want it to be caught at an early stage," said Ms. Doimer, who lives in Louisville.

The new test, expected to be available in the next few months, could have a big effect on public health if it works as advertised. That is because when ovarian cancer is caught early, when it is treatable by surgery, more than 90 percent of women live five years or longer. But right now, about three-quarters of cases are detected after the cancer has advanced, and then only 35 percent of women survive five years.

The test is also the first to use a new technology that some believers say could revolutionize diagnostics. It looks not for a single telltale protein — like the prostate-specific antigen, or P.S.A., used to diagnose prostate cancer — but rather for a complex fingerprint formed by all the proteins in the blood. Similar tests are being developed for prostate, pancreatic, breast and other cancers. The technique may work for other diseases as well.

“What is Water?”: Measurements → Insights



Color (clear, yellow, brown)
Taste (none, metallic, awful)



Measurements
Taken

Pressure (kg/cm ²)	Temp (°C)	Saturated steam		Superheated steam	
		Vapour enthalpy (kcal/kg)	Specific volume (m ³ /kg)	Density (kg/m ³)	Specific volume (m ³ /kg) at 250°C at 300°C
1	99.1	638.8	1.725	0.580	2.454 2.691
2	119.6	646.2	0.902	1.109	1.223 1.342
3	132.9	650.6	0.617	1.621	0.812 0.893
4	142.9	653.7	0.471	2.123	0.607 0.668
5	151.1	656.0	0.382	2.618	0.484 0.533
6	158.1	657.0	0.321	3.115	0.402 0.443
7	164.2	659.5	0.278	3.597	0.343 0.379
8	169.6	660.8	0.245	4.082	0.299 0.331
9	174.5	661.9	0.219	4.566	0.265 0.293
10	179.1	662.9	0.198	5.051	0.238 0.263
12	187.1	664.5	0.166	6.024	0.196 0.218
14	194.1	665.7	0.143	6.993	0.167 0.186
16	200.4	666.7	0.126	7.937	0.145 0.162
18	206.1	667.4	0.112	8.929	0.128 0.143
20	211.4	668.0	0.101	9.901	0.114 0.128
22	216.2	668.4	0.092	10.870	0.103 0.116
24	220.7	668.7	0.085	11.765	0.093 0.106
26	225.0	669.0	0.078	12.821	0.085 0.097
28	229.0	669.1	0.073	13.699	0.078 0.089
30	232.7	669.2	0.068	14.706	0.072 0.083

**LOTS of
Quantitative
“Data”**

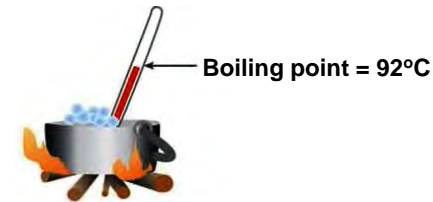


Phase (liquid, gas, solid)
Phase change (boil, melt, freeze)



Qualitative Descriptions

**But also LOTS of
disagreements...**

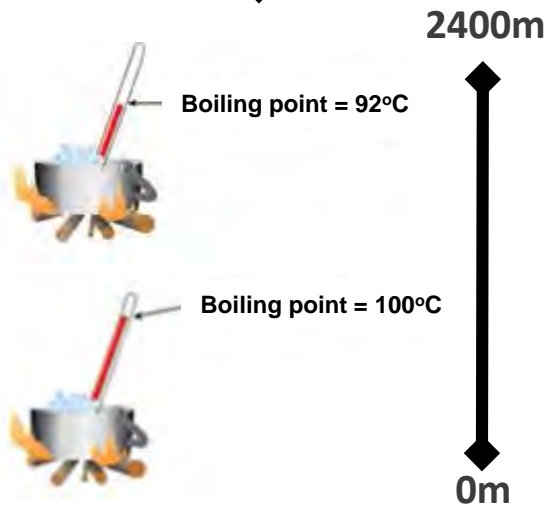
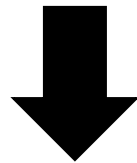


Boiling point = 100°C



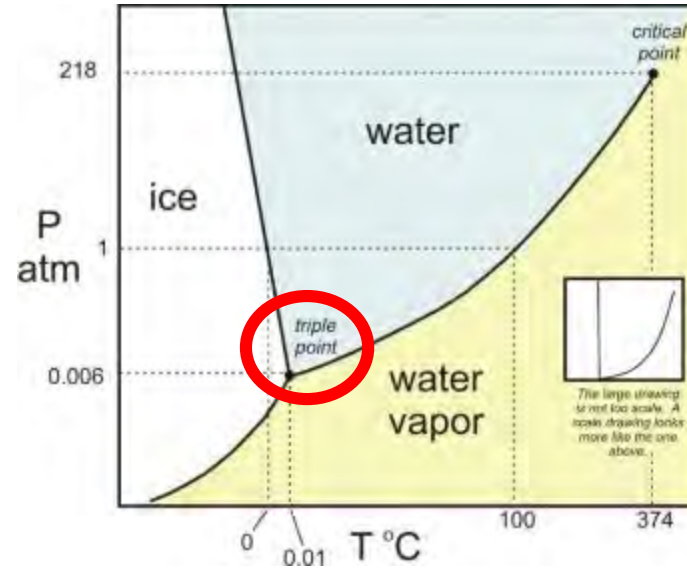
“What is Water?”: Standards and Sharing of Data → New Insights and Understanding

- Define samples and protocols
- Share collected data



New Parameter

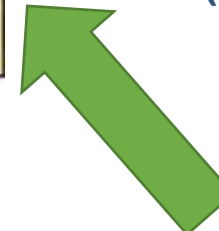
“Pressure”



New Understanding

- Phase boundaries
 - V/L equilibrium
- Triple Point

(Phase Diagram)

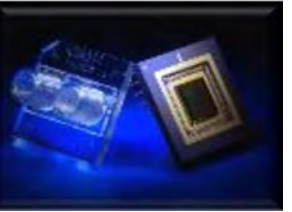
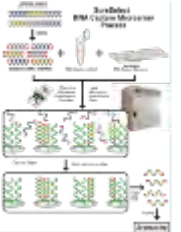


Pressure (kg/cm ²)	Temp (°C)	Saturated steam		Superheated steam		
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3	132.9	650.6	0.617	1.621	0.812	0.893
4	142.9	653.7	0.471	2.123	0.607	0.668
5	151.1	656.0	0.382	2.618	0.484	0.533
6	158.1	657.0	0.321	3.115	0.402	0.443
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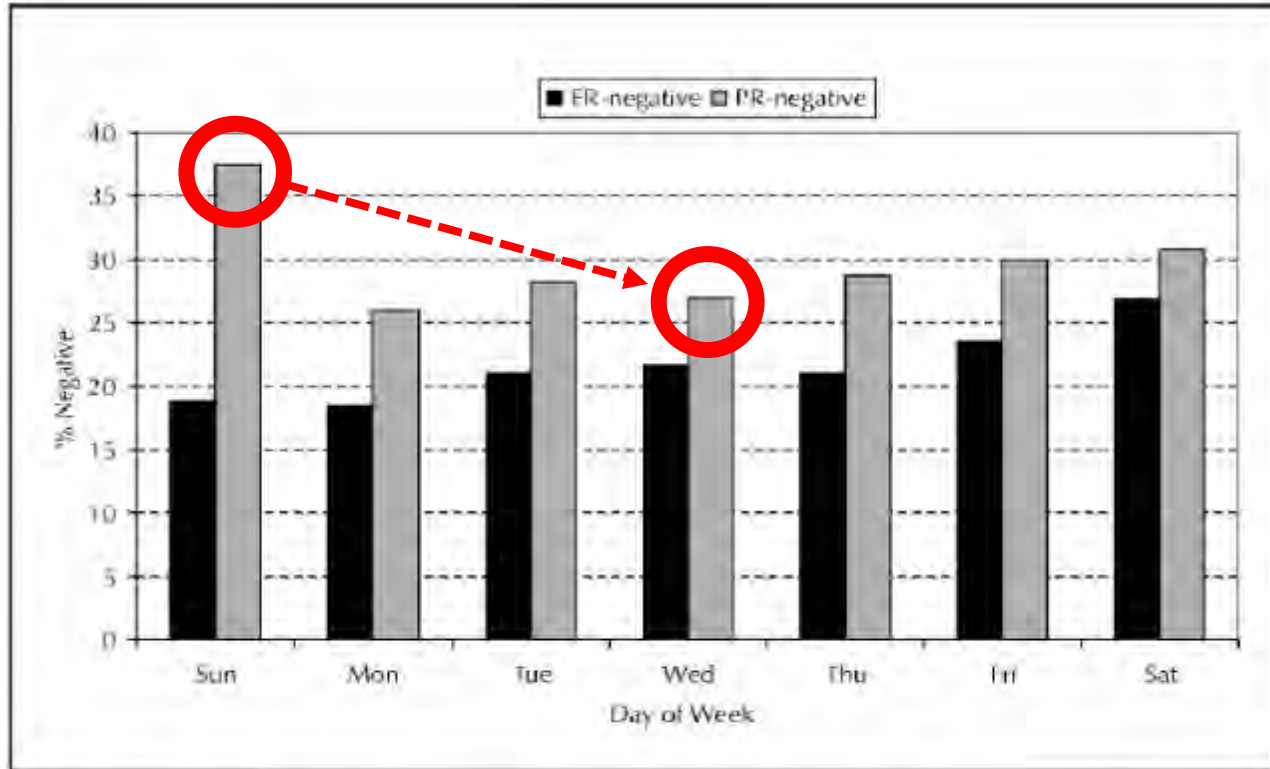
LOTS of
Quantitative
and
Reproducible
Data

(Steam Table)

Many "Thermometers" (Genomics and Proteomics)



Samples AND Handling Matter!

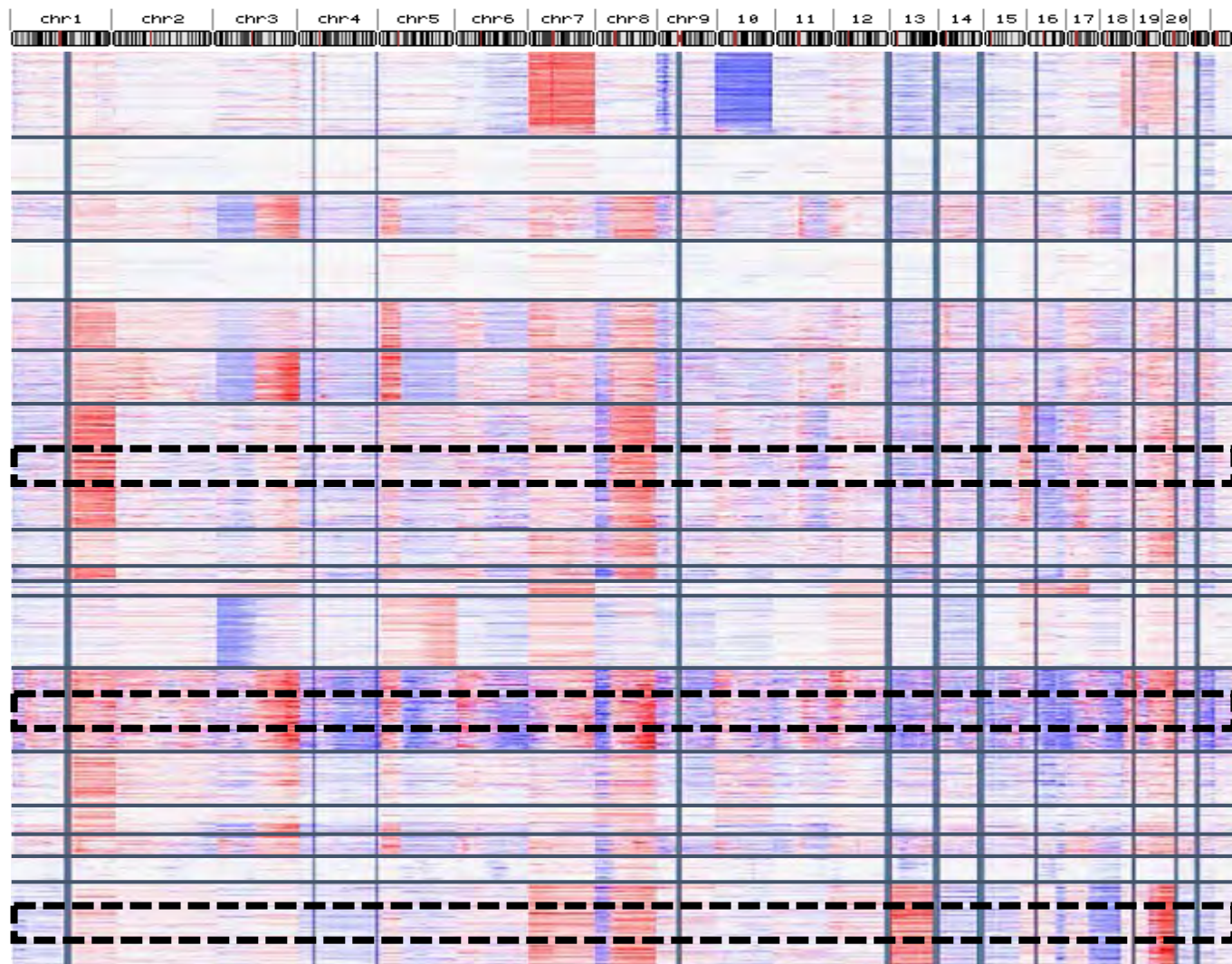


“...We found that specimens **obtained late in the week** are **more likely to be ER/PR negative** than specimens **obtained on other weekdays...**”

Day	Cases	ER-Negative	PR-Negative
Sunday	16	3	6
Monday	1252	230	325
Tuesday	1176	248	332
Wednesday	784	170	212
Thursday	904	191	259
Friday	919	216	276
Saturday	26	7	8
System	5077	1065	1418

Abbreviations: ER, estrogen receptor; PR, progesterone receptor.

The Cancer Genome Atlas



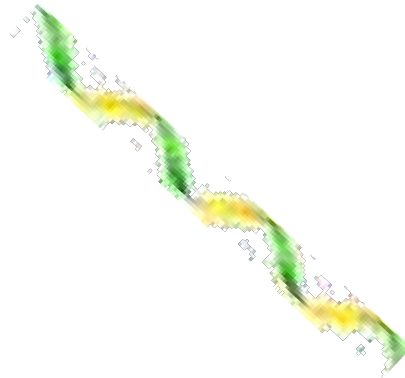
Glioblastoma:	563
Brain lower grade glioma:	180
Head & neck:	306
Thyroid carc:	401
Lung adeno:	356
Lung squamous:	343
Breast carc:	866
Stomach adeno:	237
Liver hep. carc:	97
Kidney pap. cell carc:	103
Kidney clear cell carc:	493
Ovarian serous:	559
Uterine corpus end. carc:	492
Cervical carc:	102
Bladder carc:	135
Prostate adeno:	171
Colon/rectum adeno:	575

Total: 5,979

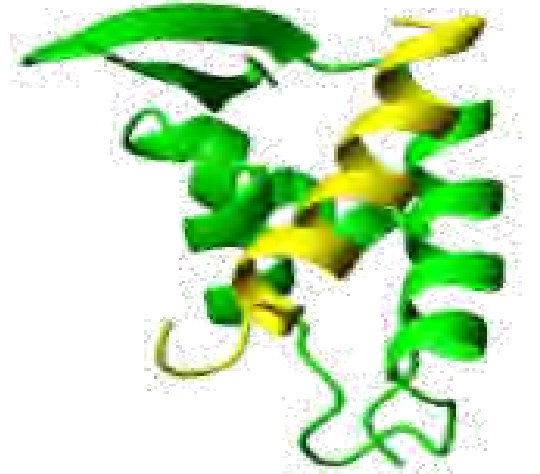
Central Dogma of Biology



DNA



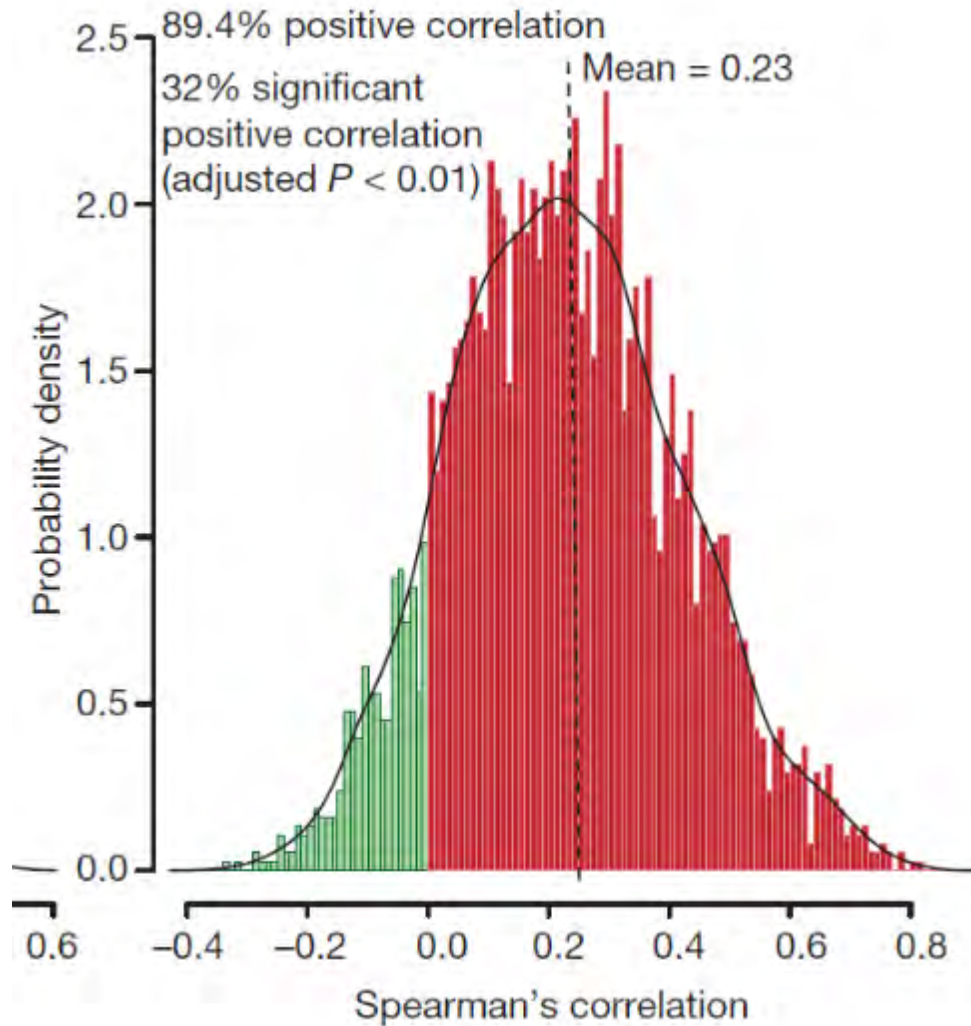
RNA



Protein

The Cancer Genome Atlas 

Re-writing Central Dogma “Rule”



On average across 375 tumor samples, **ONLY 33%** of DNA/RNA predicted cancer protein abundance

"...there is great potential for **new insights** to come from the **combined analysis** of cancer proteomic and genomic data, as proteomic data can now **reproducibly** provide information about protein levels and activities that are **difficult or impossible to infer from genomic data alone**..."

Douglas R. Lowy, MD

Acting Director of the National Cancer Institute, National Institutes of Health

5/25/2016

2016



Mission

- Make a decade's worth of progress in **cancer prevention, diagnosis, treatment, and care** – ultimately to end cancer as we know it.



STRATEGIC GOALS

Catalyze New Scientific Breakthroughs

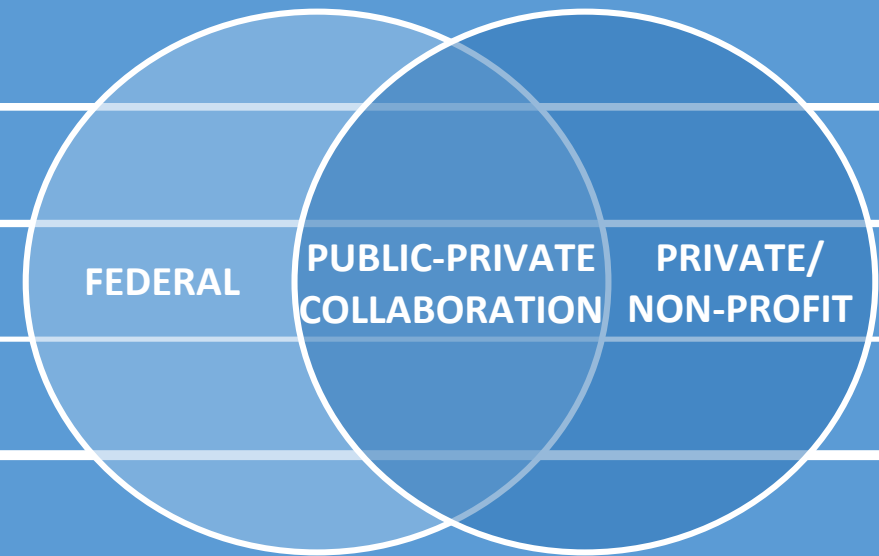
Unleash the Power of Data

Accelerate Bringing New Therapies to Patients

Strengthen Prevention and Diagnosis

Improve Patient Access and Care

IMPLEMENTATION PATH



2/1/2016

10/17/2016

Cancer Moonshot Data & Technology Team

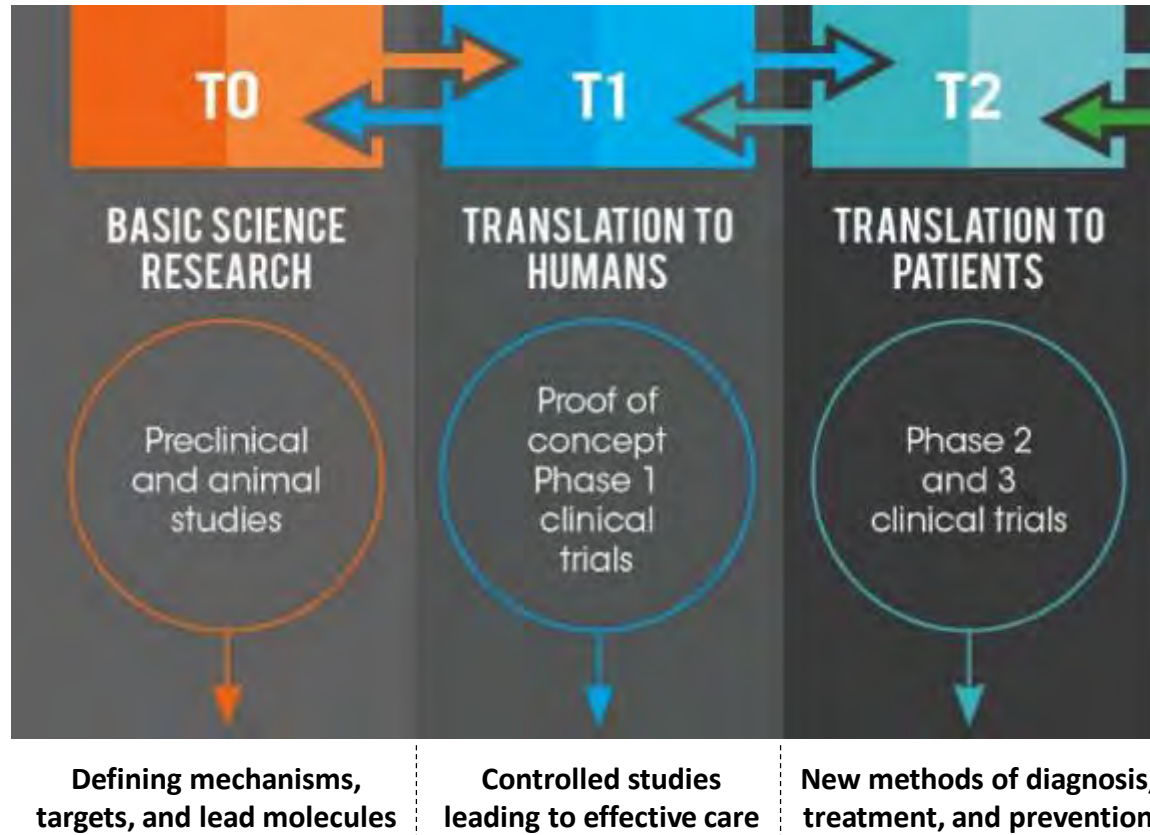
Co-Chairs: Dimitri Kusnezov (DOE), DJ Patil (OSTP), and Jerry Lee (OVP)



Members:

- John Scott (DoD)
- Craig Shriver (DoD)
- Cheryll Thomas (CDC)
- Frances Babcock (CDC)
- Teeb Al-Samarrai (DOE)
- Sean Khozin (FDA)
- Alexandra Pelletier (PIF)
- Maya Mechenbier (OMB)
- Henry Rodriguez (NCI)
- Karen Cone (NSF)
- Michael Kelley (VA)
- Louis Fiore (VA)
- Warren Kibbe (NCI)
- Betsy Hsu (NCI)
- Niall Brennan (CMS)
- Thomas Beach (USPTO)
- Claudia Williams (OSTP)
- Vikrum Aiyer (USPTO)
- Tom Kalil (OSTP)
- Kathy Hudson (NIH)
- Dina Paltoo (NIH)
- Al Bonnema (DoD)
- Michael Balint (PIF)
- Kara DeFrias (OVP)
- Greg Pappas (FDA)
- Erin Szulman (OSTP)
- Paula Jacobs (NCI)

Translational from basic science to human studies



TCGA

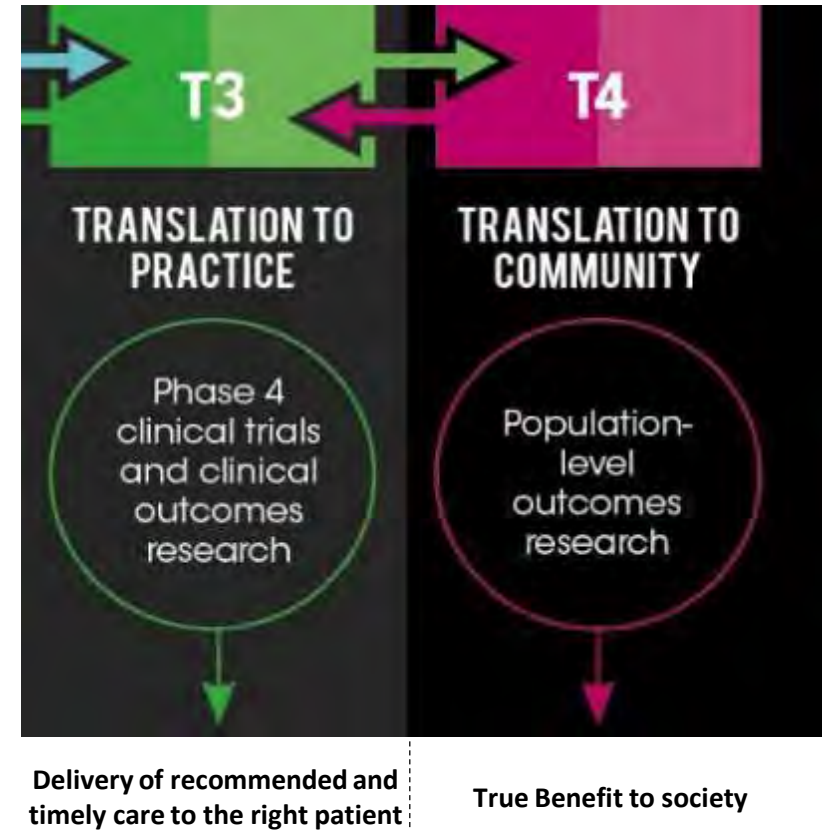
MPACT
LungMAP
ALCHEMIST

MATCH

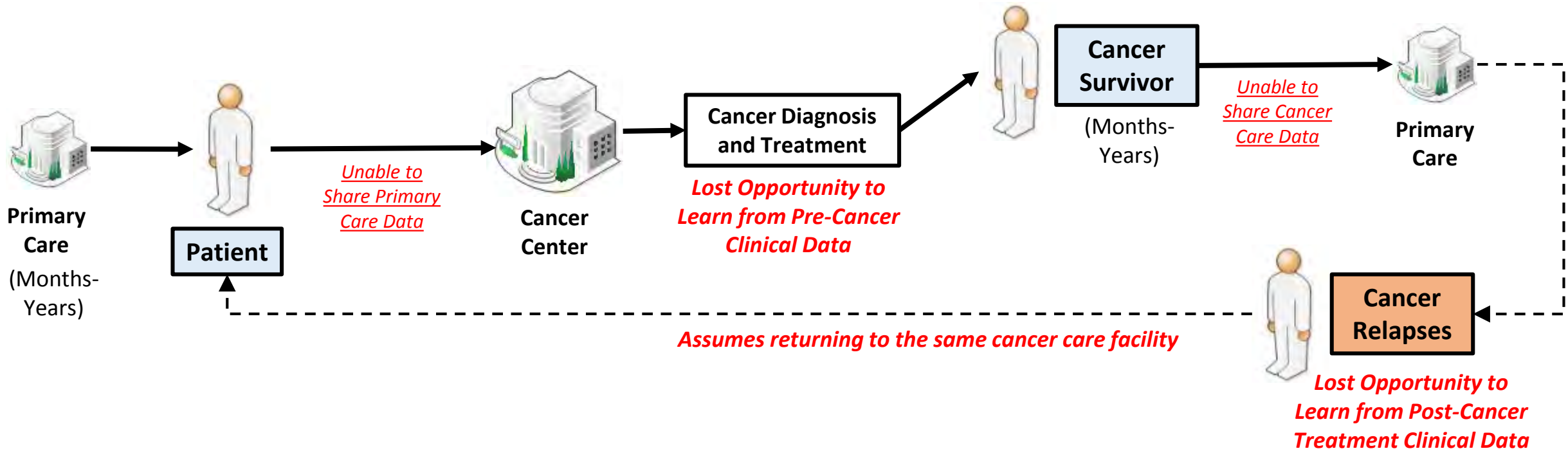
2004

2016

Translational of new data into the clinic and health decision making



Without a National Learning Healthcare System for Cancer



Vision:

Enable the creation of a *Learning Healthcare System for Cancer*, where as a nation we learn from the **contributed knowledge** and experience of **every cancer patient**. As part of the Cancer Moonshot, we want to *unleash the power of data* to **enhance, improve,** and **inform** the **journey of every cancer patient** from the *point of diagnosis through survivorship*.

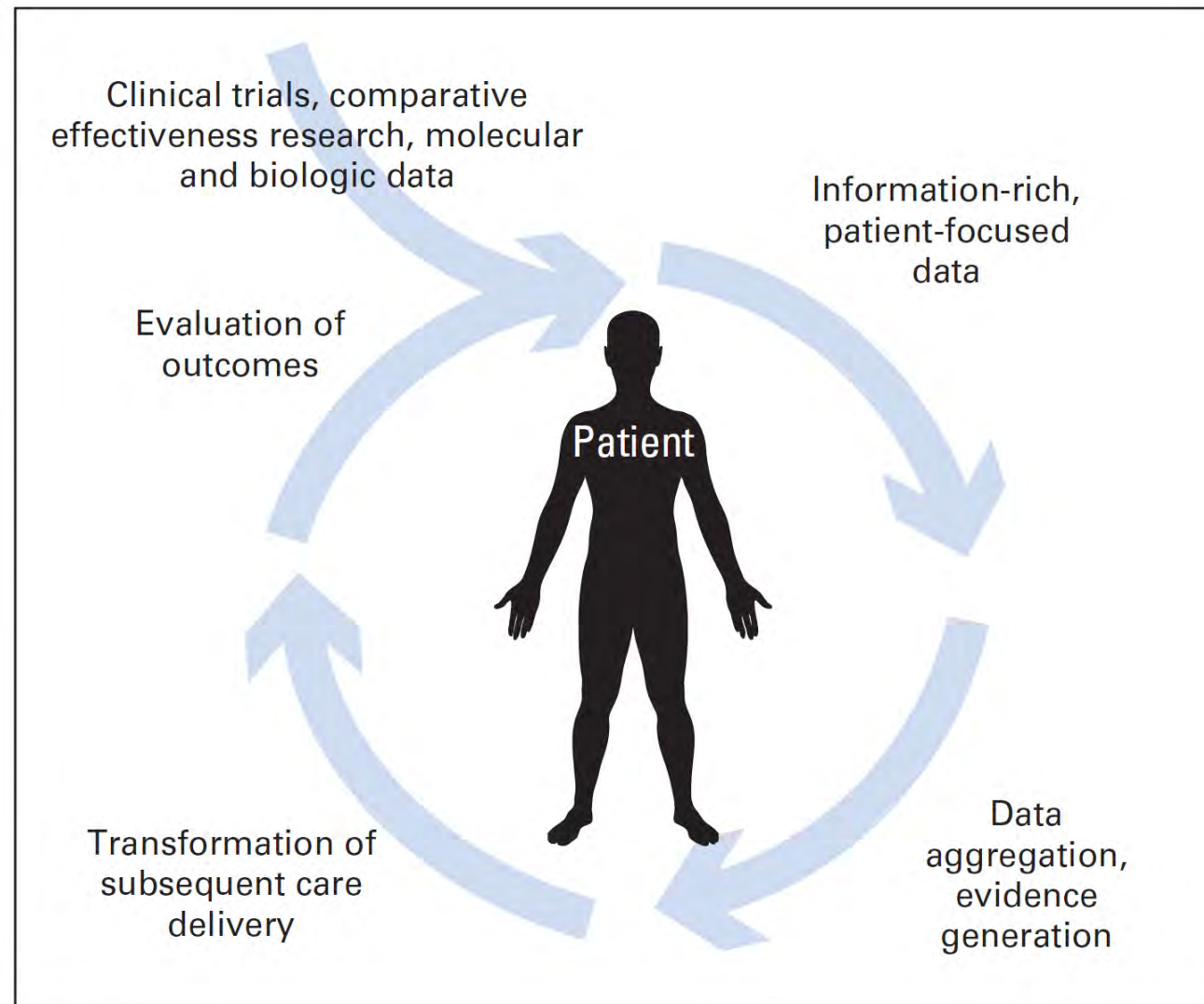


Fig 1. Cycle of evidence in rapid-learning health care. In a patient-centered system of rapid-learning health care, patient-level data are aggregated to achieve population-based change, and results are applied to care of individual patients to achieve meaningful patient-level practice change.

Priorities Areas: Cancer Moonshot Data & Technology

- Priority Area A: Enabling a seamless data environment **[If you build it...]**
- Priority Area B: Unlocking science through open computational and storage platforms **[Make it easy AND relevant to use...]**
- Priority Area C: Workforce development using open and connected data **[They will come...]**

NCI Blue Ribbon Panel Report Recommendations

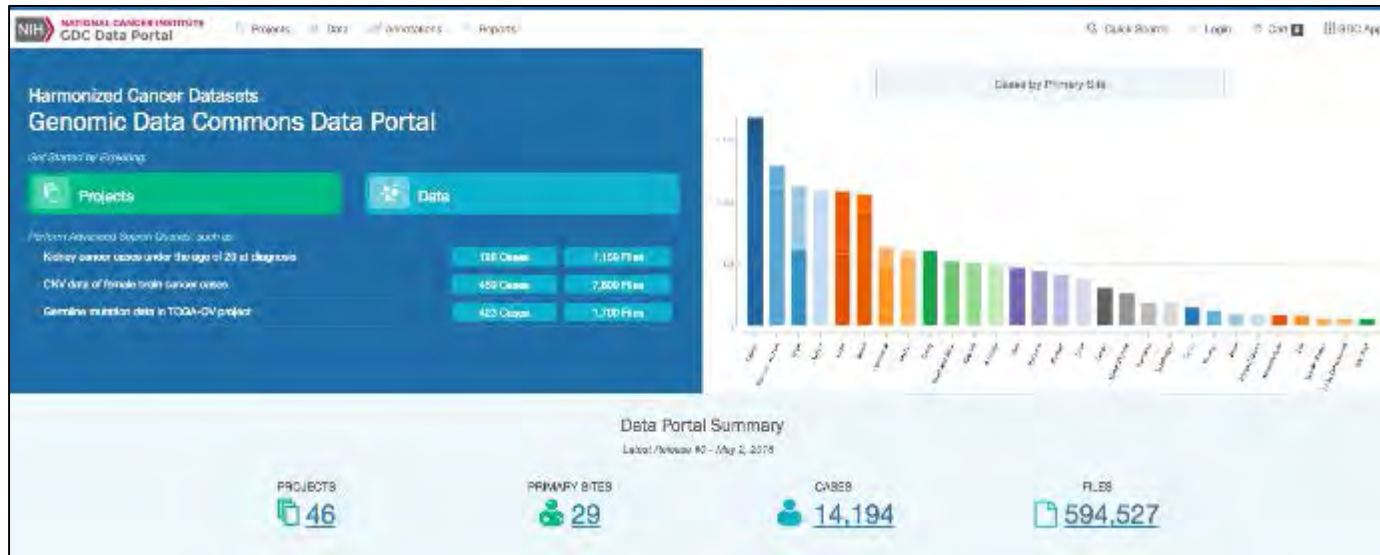


<http://www.cancer.gov/brp>

- A. Establish a network for direct patient involvement**
- B. Create a clinical trials network devoted exclusively to immunotherapy
- C. Develop ways to overcome cancer's resistance to therapy
- D. Build a national cancer data ecosystem**
- E. Intensify research on the major drivers of childhood cancers
- F. Minimize cancer treatment's debilitating side effects
- G. Expand use of proven cancer prevention and early detection strategies
- H. Mine past patient data to predict future patient outcomes**
- I. Develop a 3-D cancer atlas
- J. Develop new cancer technologies

NCI Genomic Data Commons

launched at ASCO on **June 6, 2016**



<https://gdc-portal.nci.nih.gov>

2.6 PB of legacy data and **1.5 PB** of harmonized data.

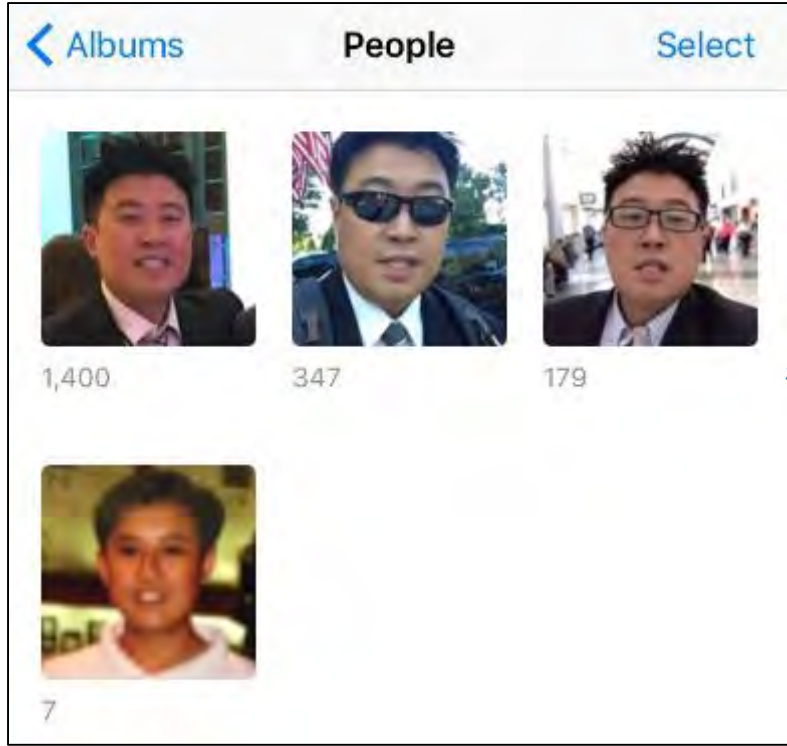
Making Data FAIR

Findable

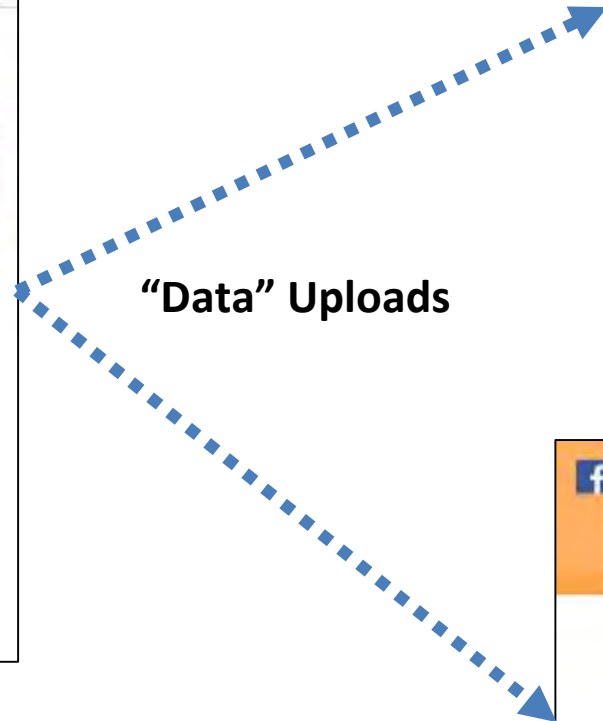
Accessible

Interoperable

Reusable



My Local "Data"
&
Reuse



"Data" Uploads



"Data" Uploads
Reuse A



"Data" Uploads
Reuse B



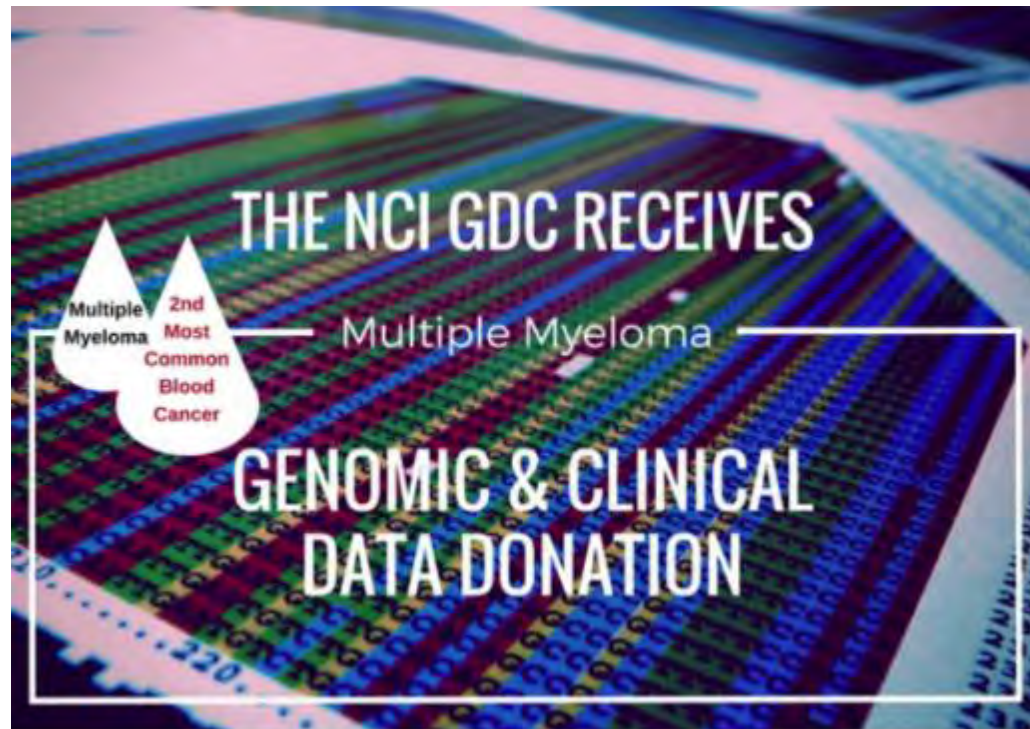
Foundation Medicine Shares Genomic Cancer Data with National Cancer Institute as Part of Cancer Moonshot and Precision Medicine Initiatives

June 29, 2016 Foundation Medicine

At the June 29th Cancer Moonshot Summit, *Foundation Medicine* announced the release of **18,000** genomic profiles to the NCI GDC

Multiple Myeloma Research Foundation to Contribute to NCI's Genomic Data Commons

Sep 29, 2016 | a [GenomeWeb](#) staff reporter



- MMRF is the **first non-profit organization** to upload information to the GDC
- Among its contributions will be data from relating **Clinical Outcomes** in **MM** to **Personal Assessment of Genetic Profile (CoMMpass)** study which began in 2011 and has thus far enrolled over **1,150** patients
- Over the next eight years, patients in CoMMpass will get a **repeat biopsy** and a new genomic analysis at **each six-month checkup** and/or at disease progression
- Tumor samples are being collected and analyzed when possible at the time of any relapse. **New data will be deposited every six months at a minimum**

01/05/2017



AACR Project GENIE: Data

Today, the first set of cancer genomic data aggregated through AACR Project Genomics Evidence Neoplasia Information Exchange (GENIE) is available to the global community. The data set includes nearly **19,000 de-identified** genomic records collected from patients who were treated at each of the consortium's participating institutions, making it among the largest fully public cancer genomic data sets released to date. These data will be continuously updated on a quarterly basis.

The release includes data for 59 major cancer types, including data on nearly 3,000 patients with lung cancer, more than 2,000 patients with breast cancer, and more than 2,000 patients with colorectal cancer. For more details about the data, and how to use it, consult the [data guide](#).

Users can access the data directly via [cbioportal](#), or download the data directly from [Sage Bionetworks](#). Users will need to create an account for either site and agree to the [terms of access](#).

“...while many of the important lessons learned from this first year of the project...Sawyers said that the data also speaks for itself in terms of the project's success **in tackling difficult harmonization challenges...**

...All the subjects in the GENIE collection had to have **level 1A clinical data**, Sawyers explained, and the group had to make sure that they could harmonize this information across sites. For example, **"you can imagine a name of a rare condition might be very different in Amsterdam versus Toronto or Nashville," Sawyers said...**”

GDC Content

Current

- ❖ TCGA 11,353 cases
- ❖ TARGET 3,178 cases

Coming soon

- ❖ Foundation Medicine 18,000 cases
- ❖ Cancer studies in dbGAP ~4,000 cases

Planned (1-3 years)

- ❖ NCI-MATCH ~3,000 cases
- ❖ Clinical Trial Sequencing Program ~3,000 cases
- ❖ Cancer Driver Discovery Program ~5,000 cases
- ❖ Human Cancer Model Initiative ~1,000 cases
- ❖ **APOLLO – VA-DoD ~8,000 cases**

~56,000 cases





Col. Craig Shriver, MD



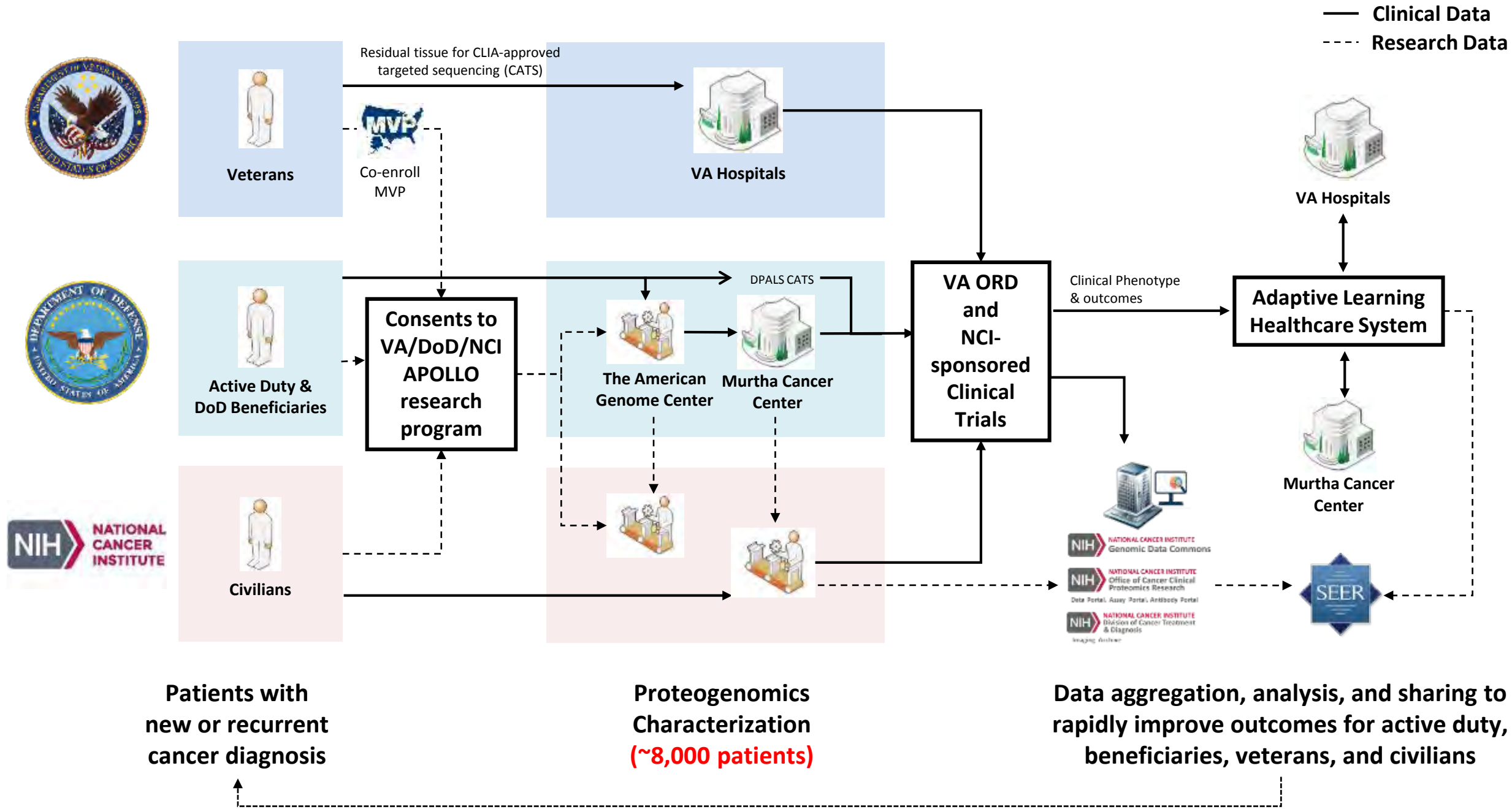
Jennifer Lee, MD



Henry Rodriguez,
PhD, MBA

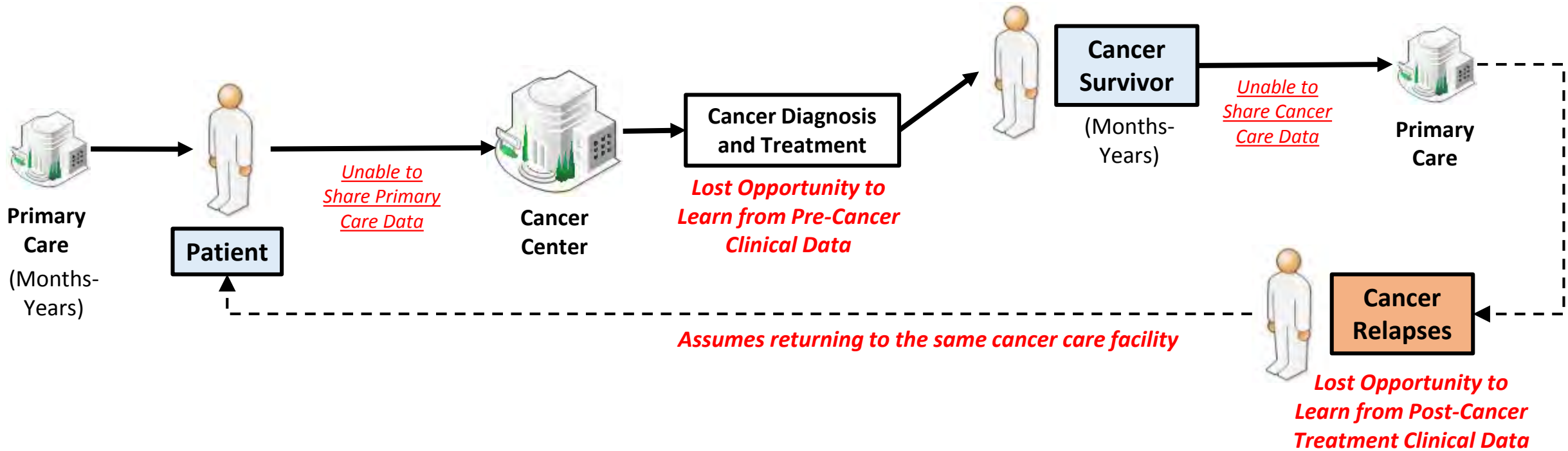


<https://medium.com/cancer-moonshot/>

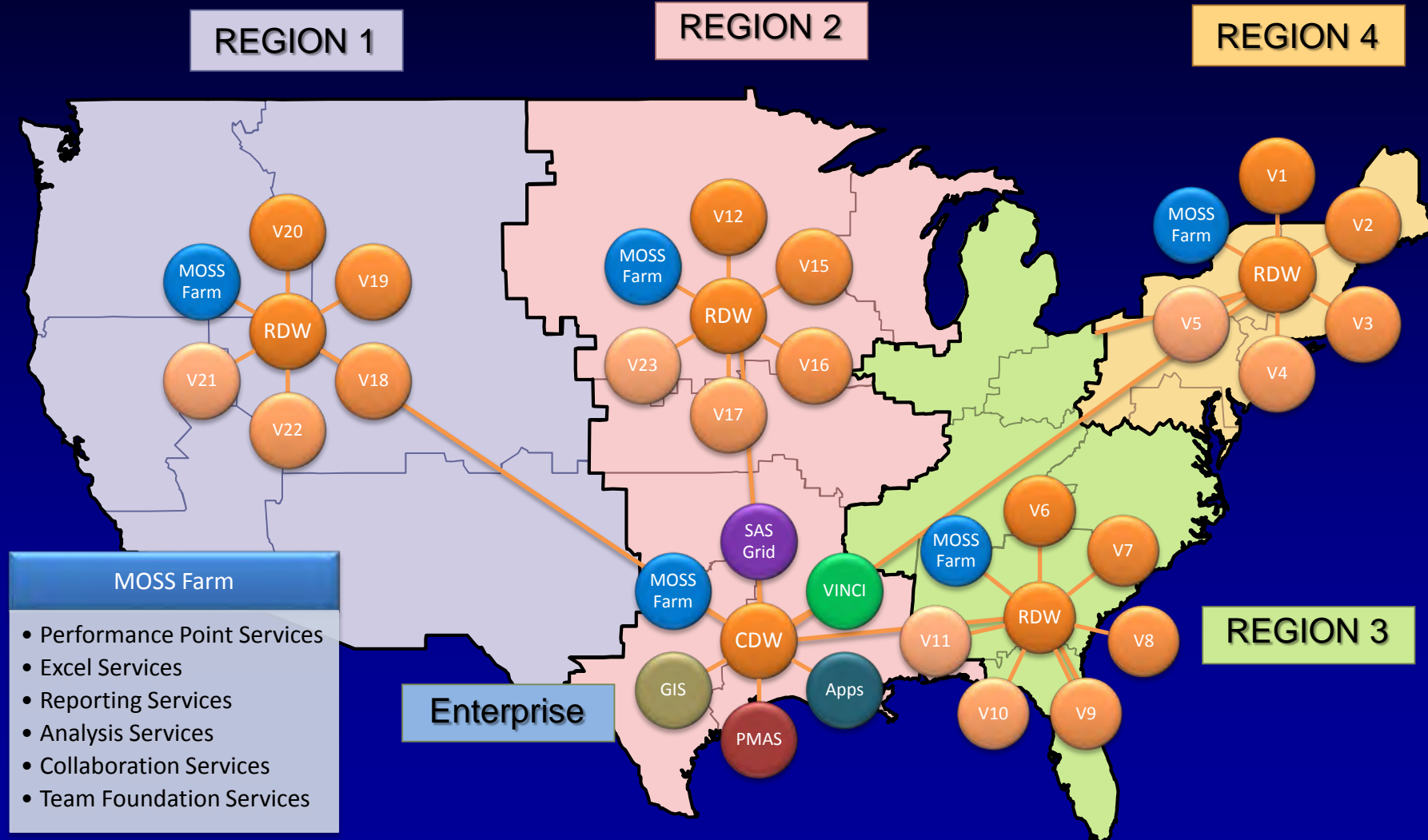


APOLLO – Applied Proteogenomics Organizational Learning and Outcomes consortium

Without a National Learning Healthcare System for Cancer



VA Medical Centers Regional / Corporate Data Warehousing and Analytical Environment

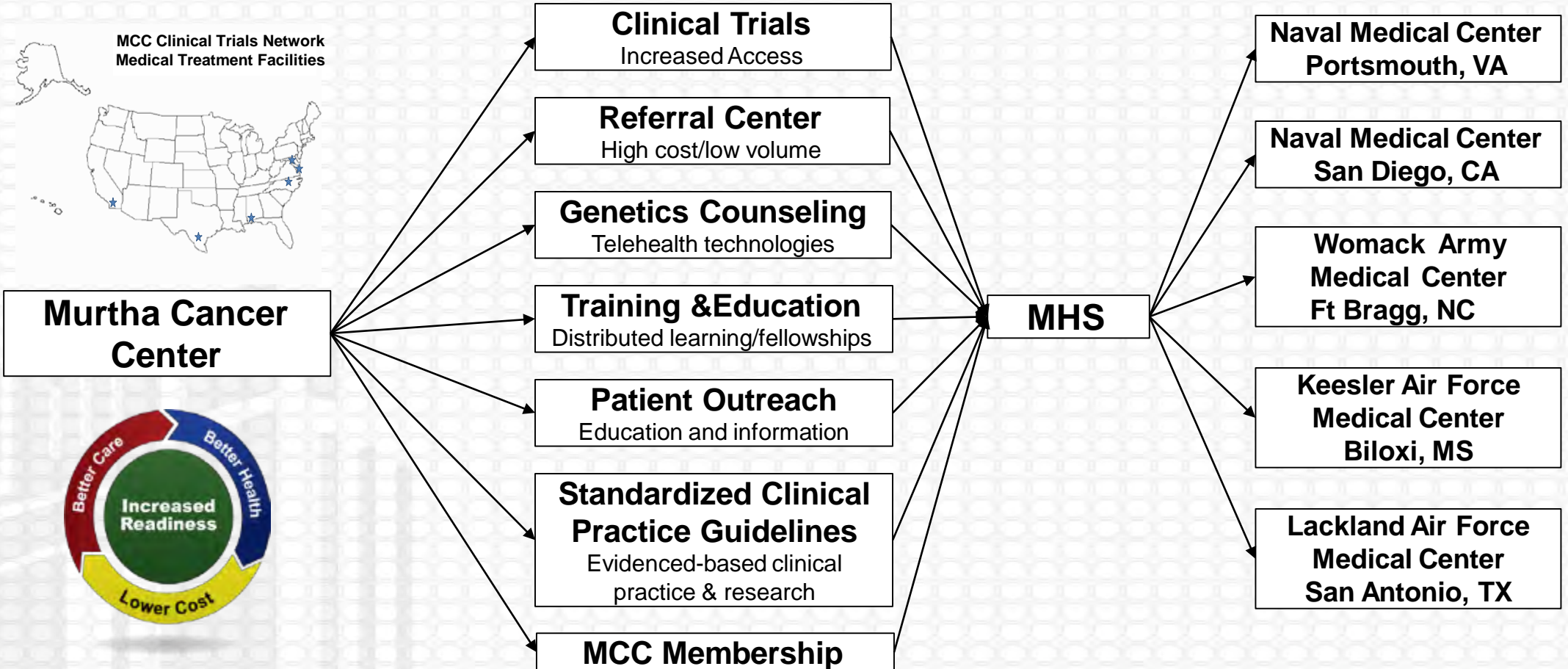




Walter Reed
National Military
Medical Center

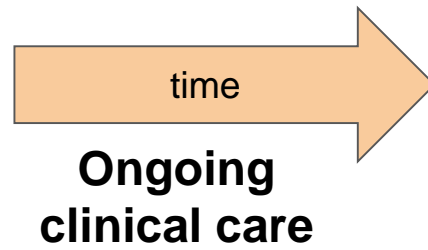


MCC Military Clinical Trials Network



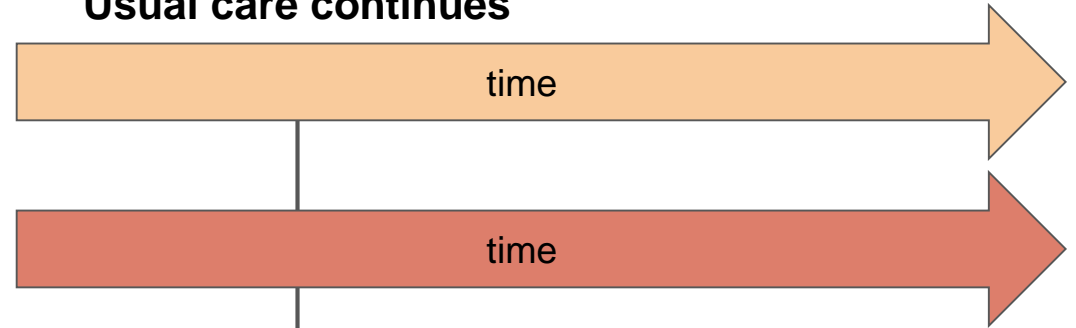


Patient and Providers



Initiate system for specific learning or research activities

Usual care continues



Research activities



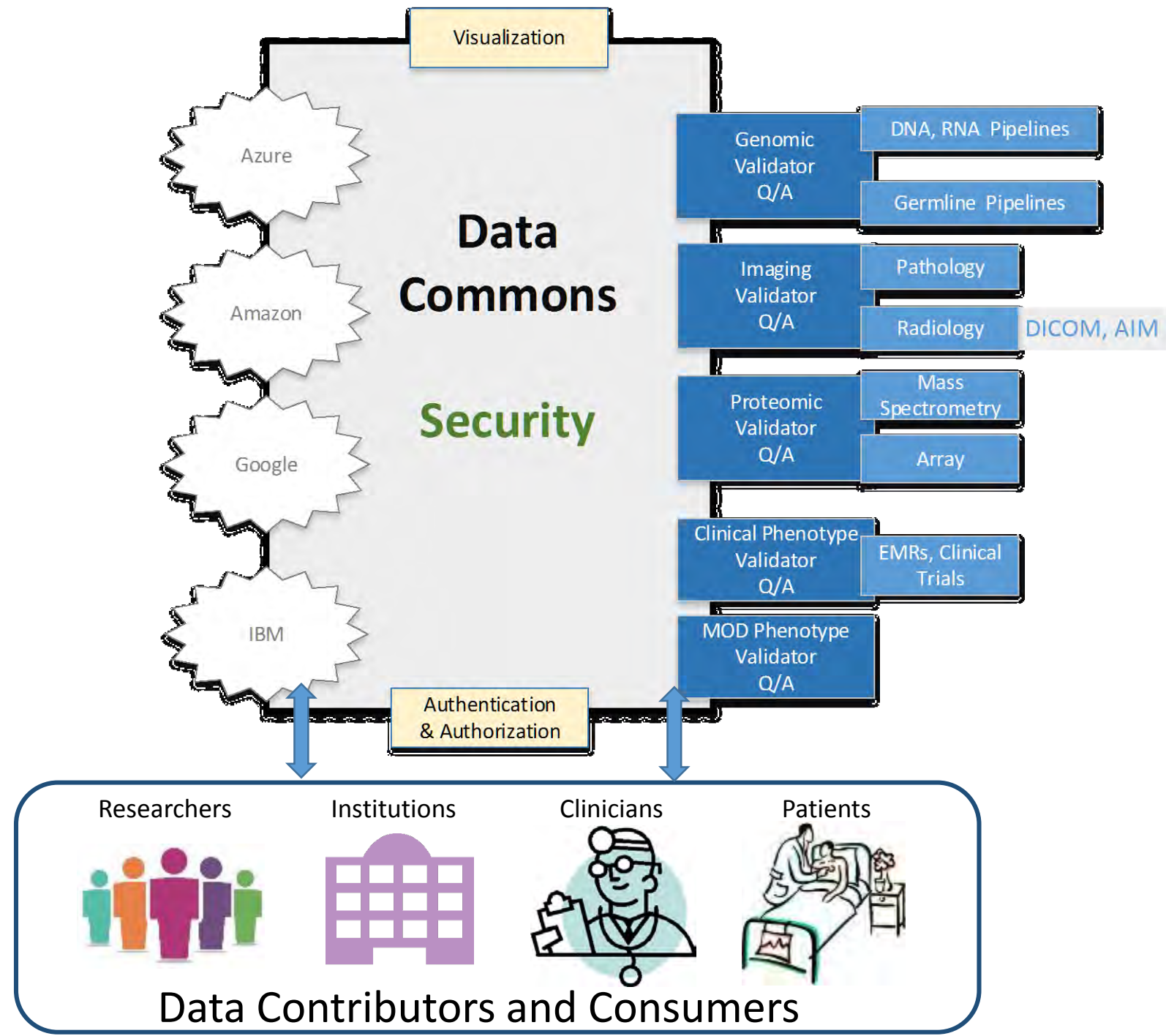
Ongoing collection of data [genomics, proteomics, medical imaging]



Applied Proteogenomics Organizational Learning and Outcomes

**APOLLO Leadership Meeting
August 29, 2016**

NCI Thesaurus
caDSR
NLM UMLS
RxNorm
LOINC
SNOMED



7/17/2016



<https://www.whitehouse.gov/the-press-office/2016/07/16/fact-sheet-victoria-comprehensive-cancer-center-vice-president-biden>

“...**proteogenomics**, which is -- as I used a metaphor -- it's like the **genes** are the **full roster** of a **basketball team**...but the winning strategy comes from finding out who their starting lineup is. The **proteins** are the **starters** you're going to play against -- the five you are going to **have to defend against**

I'm pleased to say, Mr. Prime Minister, that we've signed three memorandums of understanding between our two nations ...we're going to be able to **share patient histories, proteogenomics and clinical phenotypes data** -- data on various proteins and genetic characteristics of almost **60,000 patients** in Australia and the United States with full privacy protections...

And I predict that you're going to see this repeated around the world.”

- Vice President Biden, Australia

9/19/2016

Joe Biden announces major new steps in his fight for better cancer research

1.6k
SHARES

Share on Facebook

Share on Twitter



Vice President Joe Biden speaks at the 2016 Social Good Summit on Monday, Sept. 19, 2016

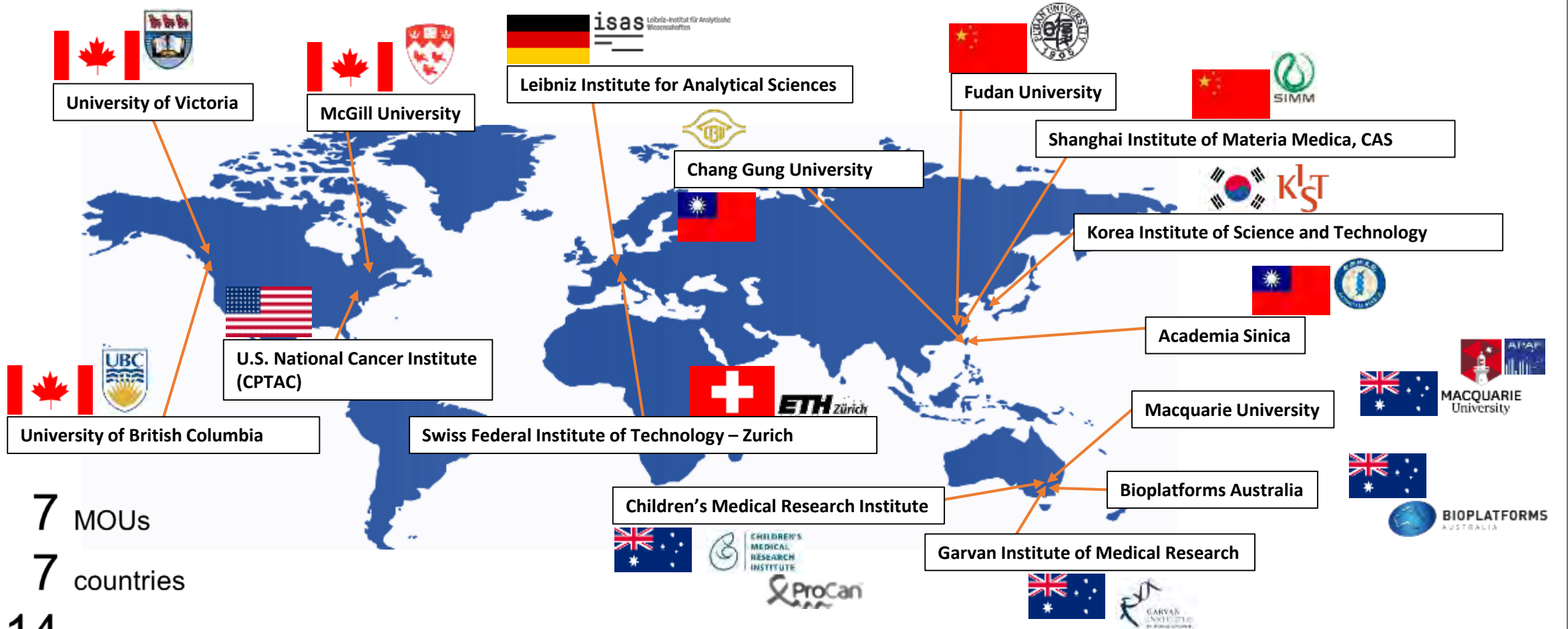
IMAGE: AP/WIDEWORLD

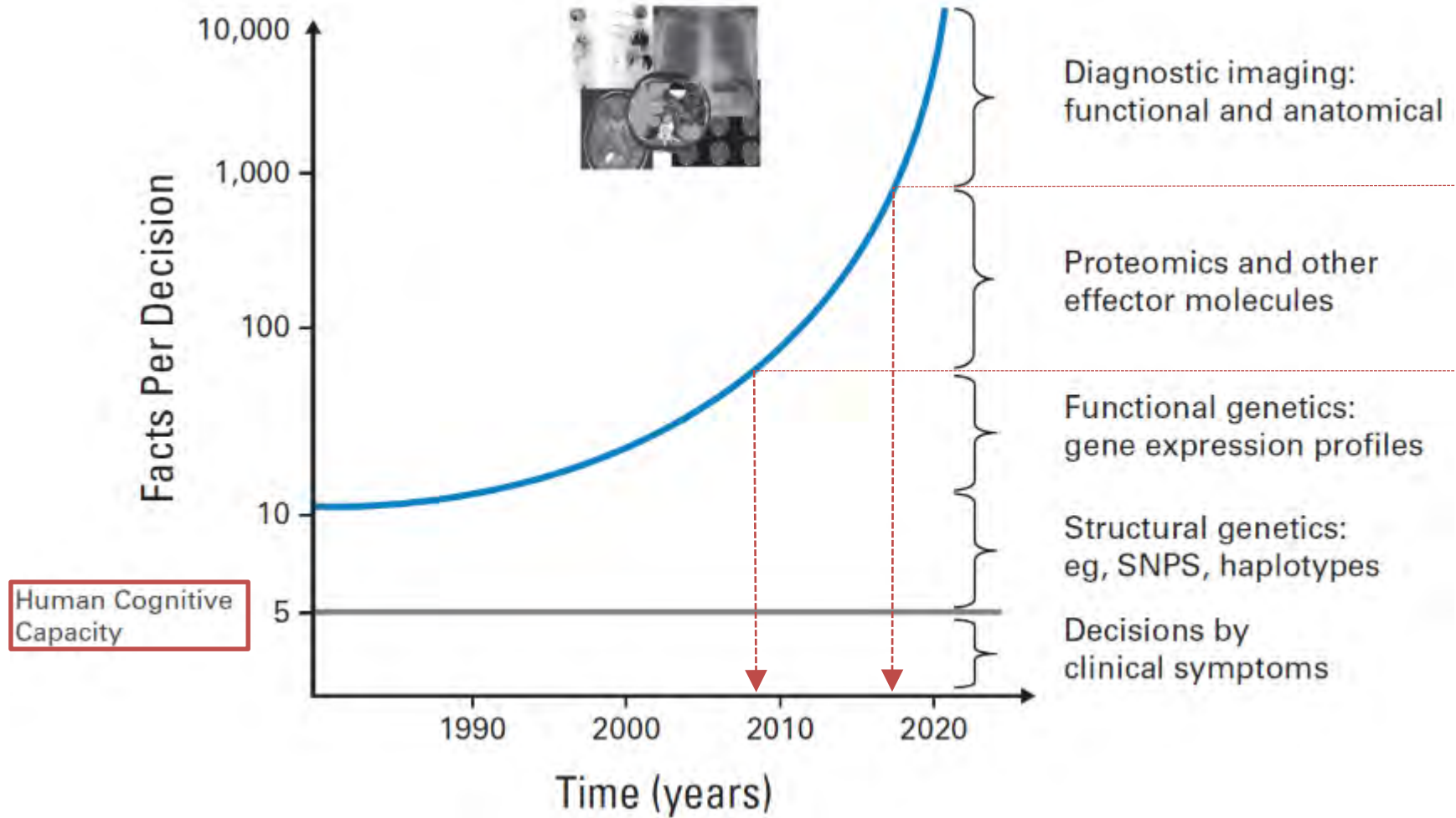
The first step is the announcement of 10 new commitments with nine nations to support better international cancer research and care. The U.S. will work with institutions in Canada, China, Germany, Switzerland, Taiwan, Japan and South Korea in the field of proteogenomics, and with Serbia, Sweden and Japan to open a discussion about better prevention, screenings, treatment and research collaborations. The U.S. Department of Energy will also work with Norway to share 1.7 million cervical screening results over the course of 25 years to seek out patterns in diagnoses and treatments.

Biden also announced the creation of regional hubs — collaborative centers that can help decrease disparities in cancer research around the world. These hubs will be funded by the National Cancer Institute, working with Japan, South Korea, New Zealand, the focus on areas of the world where spe



International Proteogenomic Moonshot Programs





What About Blood?

Eight Milestones Of 2016 In The War On Cancer



Arlene Weintraub, CONTRIBUTOR

I cover the science and business behind drug development and health. [FULL BIO](#)

Opinions expressed by Forbes Contributors are their own.

DEC 28, 2016 @ 08:35 AM

10,242 VIEWS

Only [22 novel drugs](#) were approved by the U.S. Food & Drug Administration in 2016—way down from the [45 approved in 2015](#)—and just six of the new entries are for treating or diagnosing cancer. Still, 2016 was far from a washout for oncology research. Here were some of the high points of the year in the war on cancer:



Between the Cancer Moonshot and the 21st Century Cures Act, the White House made a big commitment to cancer research in 2016. (Credit: MANDEL NGAN/AFP/Getty Images)

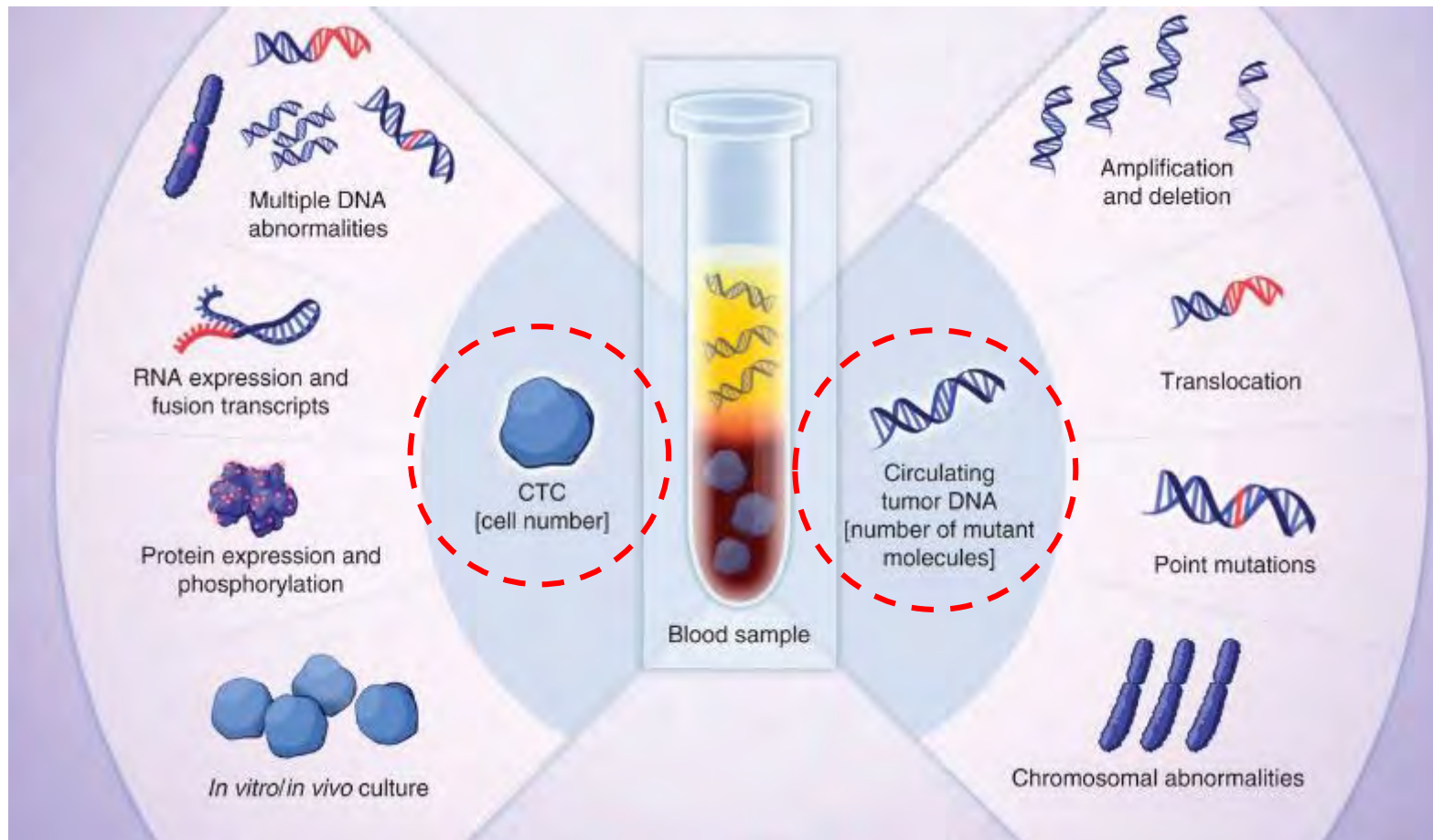
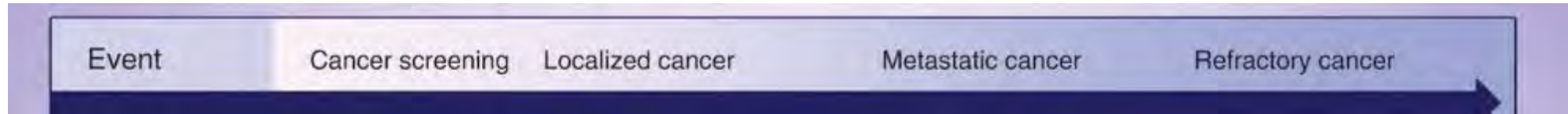
January 12: V.P. Biden is tapped to lead the new Cancer Moonshot. During the State of the Union address, President Barack Obama [launched the Cancer Moonshot](#) and appointed Vice President Joe Biden to lead the initiative. The goal is not just to speed new therapies to market, but also to improve access to treatment and come up with methods for detecting the disease early, when it's easier to treat and cure. Biden spent much of the year tapping academics, drug company executives, patients and physicians for advice, as well as [establishing five strategic goals](#) for the Moonshot.

In October, the veep announced a new project called the Blood Profiling Atlas, which will try to accelerate the development of “liquid biopsies,” tests designed to detect early-stage cancer by tracking tiny bits of tumor DNA cast off into the bloodstream. Much to the surprise of some executives, [he assembled an impressive list of organizations](#) that might normally balk at the idea of working together to help build the atlas, including

[Novartis](#) , [Pfizer](#) [PFE +0.50%](#) , Thermo Fisher and Foundation Medicine.

April 13: Napster billionaire Sean Parker launches a cancer institute. The glitzy star-studded gala that ushered in Sean Parker's new cancer research institute may have [drawn some ire](#), but the money the Napster founder and former Facebook president is devoting to the effort is nothing to sneeze at. The Parker Institute for Cancer Immunotherapy is backed

Finding the Right “Needle” at the Right “Time” of Disease



10/17/16

Biden announces USC participation in new Cancer Moonshot project

The project aims to accelerate the development of reliable blood tests for biologically based precision treatment and disease monitoring

Personal Genome Diagnostics Selected to Participate in U.S. Cancer Moonshot Effort to Jump Start Liquid Biopsy Database

-- PGDx Joins Consortium of Public, Private and Academic Innovators Developing Pilot for a Blood Profiling Atlas to Help Ensure the Validity, Utility and Accessibility of Liquid Biopsies for Cancer Research, Diagnosis and Treatment --

Seven Bridges Joins Cancer Moonshot Initiative with Cloud Platform to Speed Blood Profiling Research

Blood Profiling Atlas Project Aims to Advance the Development of Simple, Blood-Based Test for Early Cancer Diagnosis

U-M researchers will support new 'liquid biopsy' Cancer Moonshot initiative

Government, academia, pharma partnership will create open dataset to stimulate translational research

Lilly Partnering on Cancer Research

Posted: Oct 17, 2016 5:00 PM EDT

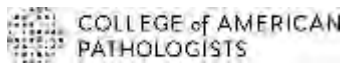
Updated: Oct 18, 2016 8:53 AM EDT



Guardant Health makes commitment to share expertise and data to advance Blood Profiling Atlas, part of White House Cancer Moonshot

College Of American Pathologists To Participate In The Cancer "Moonshot" Effort

Industry News: Thermo Fisher Scientific Joins the Cancer Moonshot Initiative to Help Advance Precision Medicine



10/18/16: Blood Profiling Atlas Face 2 Face

12/20/16



Blood Profiling Atlas Members

- Open Commons Consortium (OCC) in collaboration with the University of Chicago commits to organizing and operating an open Blood Profiling Atlas Commons. The Commons will be based upon the same open source software stack used by the NCI Genomic Data Commons so that the genomic, image and clinical data in the Commons can be shared with the appropriate security, privacy and compliance controls. To facilitate the rapid development of this critical resource, the OCC / University of Chicago team will contribute up to six months of engineering, bioinformatics and project management resources to the project and up to \$500,000 of compute and storage resources for building the commons and for the use of the commons by the research community.
- Seven Bridges will contribute its experience in accelerating pharmaceutical research and development and in building national-scale research systems by developing the Blood Profiling Atlas Analysis Cloud, specifically tailored to the needs of the liquid biopsy community. This environment – based on the company's work partnering with the NCI, Genomics England, the VA and others – will integrate with the Blood Profiling Atlas Commons, allowing molecular, clinical and imaging data to be easily, securely, and cost effectively analyzed by researchers across disciplines. In addition, Seven Bridges will share its expertise in cancer genomics and immunoinformatics analysis. To do so, the company will release algorithms for analyzing liquid biopsy data at scale, committing six months of engineering, bioinformatics and project management resources, and up to \$500,000 compute and storage resources to facilitate use of the analysis tools and data donated by the Blood Profiling Atlas community.
- In support of the Blood Profiling Atlas, AstraZeneca will provide standard operating procedures for ctDNA isolation and library construction for targeted and whole genome/exome sequencing of ctDNA. The AstraZeneca bioinformatics pipeline for variant calling in ctDNA is available for all interested parties. Furthermore, AstraZeneca will generate ctDNA for comparative studies of other bioinformatics pipelines with the goal to develop best practices in identifying variants in ctDNA after high depth sequencing that will standardize analyses of data acquisition for the Atlas. AstraZeneca will additionally provide data on method comparisons and can assist in data generation for samples provided to this project.
- Celgene will contribute advanced analytic capabilities to the effort and sponsor a crowdsourced Challenge for patient benefit in the blood profiling domain. Celgene has



Members

- Open Commons Consortium
- University of Chicago
- Seven Bridges
- AstraZeneca
- Celgene
- CytoLumina
- Eli Lilly and Company
- Epic Sciences
- Foundation Medicine
- Genentech
- Genomic Health, Inc
- Guardant Health
- Memorial Sloan Kettering Cancer Center
- Novartis
- Personal Genome Diagnostics
- Pfizer
- Thermo Fisher Scientific
- University of Michigan
- University of Southern California

<https://medium.com/cancer-moonshot/blood-profiling-atlas-in-cancer>

<https://www.bloodpac-data.org/>



2017



“Working together, we got the job done” - Medical Innovation Game-Changer Now Law

Dec 13, 2016 [Press Release](#)

Signing Ceremony Marks Culmination of Three-Year Journey on the #Path2Cures

Delivering #CuresNow: Speaker Ryan Signs Bipartisan Game-Changing Medical Innovation Bill

Dec 8, 2016 [Press Release](#)

Bill Officially Heads to The White House to be Signed into Law





Funding for NIH Innovative Research Initiatives under the Cures Act.*				
Fiscal Year	BRAIN	PMI	Cancer Moonshot	Regenerative Medicine
	<i>millions of \$</i>			
2017	10	40	300	2
2018	86	100	300	10
2019	115	186	400	10
2020	140	149	195	8
2021	100	109	195	
2022	152	150	194	
2023	450	419	216	
2024	172	235		
2025	91	36		
2026	195	31		
10-Yr total	1,511	1,455	1,800	30

* BRAIN denotes Brain Research through Advancing Innovative Neurotechnologies, and PMI Precision Medicine Initiative.

“***To support*** cancer research, such as the development of cancer vaccines, the development of more sensitive diagnostic tests for cancer, immunotherapy and the development of combination therapies, and research that has ***the potential to transform*** the scientific field, that has ***inherently higher risk***, and that seeks to ***address major challenges*** related to cancer...”

H.R.34—21st Century Cures Act

NCI Blue Ribbon Panel Report Recommendations



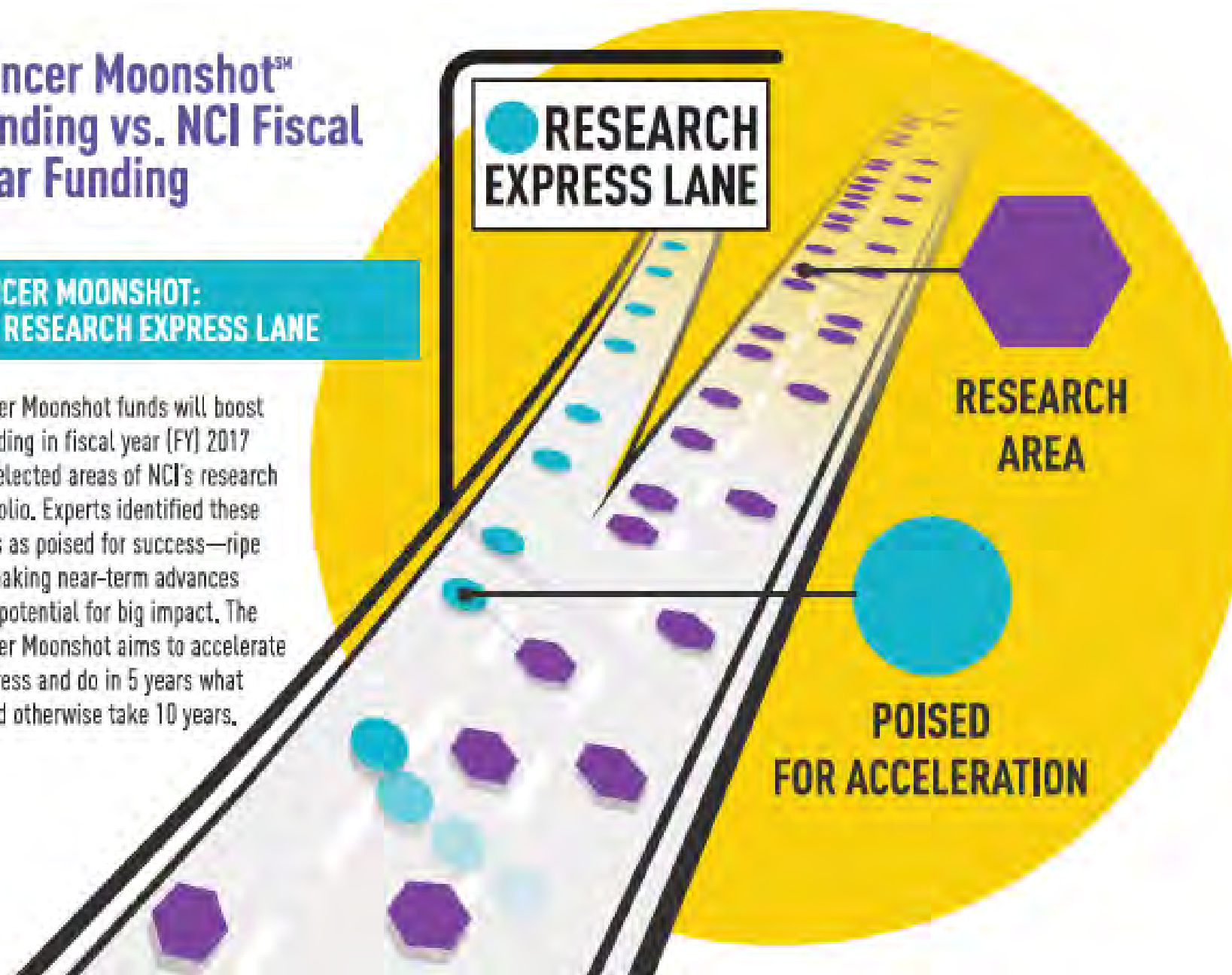
<http://www.cancer.gov/brp>

- A. Establish a network for direct patient involvement
- B. Create a clinical trials network devoted exclusively to immunotherapy
- C. Develop ways to overcome cancer's resistance to therapy
- D. Build a national cancer data ecosystem
- E. Intensify research on the major drivers of childhood cancers
- F. Minimize cancer treatment's debilitating side effects
- G. Expand use of proven cancer prevention and early detection strategies
- H. Mine past patient data to predict future patient outcomes
- I. Develop a 3-D cancer atlas
- J. Develop new cancer technologies

Cancer MoonshotSM Funding vs. NCI Fiscal Year Funding

CANCER MOONSHOT: THE RESEARCH EXPRESS LANE

Cancer Moonshot funds will boost spending in fiscal year (FY) 2017 for selected areas of NCI's research portfolio. Experts identified these areas as poised for success—ripe for making near-term advances with potential for big impact. The Cancer Moonshot aims to accelerate progress and do in 5 years what would otherwise take 10 years.



The screenshot shows the National Cancer Institute (NCI) website. At the top left is the NIH logo and the text 'NATIONAL CANCER INSTITUTE'. A navigation bar includes links for '1-800-4-CANCER', 'Live Chat', 'Publications', and 'Dictionary'. Below this is a main menu with 'ABOUT CANCER', 'CANCER TYPES', 'RESEARCH', 'GRANTS & TRAINING', 'NEWS & EVENTS', and 'ABOUT NCI'. A search bar is on the right. A light blue banner at the top right says 'Get email updates from NCI on the Cancer Moonshot'. The breadcrumb trail reads 'Home > Research > Key Initiatives > Cancer Moonshot® > Milestones'. A sidebar on the left has 'CANCER MOONSHOT®' and three items: 'Blue Ribbon Panel Report', 'Milestones', and 'Related Activities'. The main content area has the title 'NCI-Related Cancer Moonshot Activities' and a paragraph: 'Through the Cancer Moonshot™, many new collaborations and activities have been launched that address barriers and opportunities in cancer research. NCI is involved in a number of these activities.' Below this is a section header 'NCI-Related Activities Underway' followed by 'National Cancer Institute Formulary'. The text describes leveraging lessons from the NCI-MATCH Trial to create a formulary for testing agents. A second section header is 'Proposed Activities with NCI Involvement', followed by the sub-header 'Create a high-quality performance status tracking system for cancer patients during therapy and long-term follow-up'. The text describes a joint effort between NCI and DoD to improve patient and military personnel outcomes.

01/11/2017

NCI Press Release

New Drug Formulary Will Help Expedite Use of Agents in Clinical Trials

Posted: January 11, 2017

Contact: NCI Press Office
301-496-6641

The National Cancer Institute (NCI) today launched a new drug formulary (the “NCI Formulary”) that will enable investigators at NCI-designated Cancer Centers to have quicker access to approved and investigational agents for use in preclinical studies and cancer clinical trials. The NCI Formulary could ultimately translate into speeding the availability of more-effective treatment options to patients with cancer.

The NCI Formulary is a public-private partnership between NCI, part of the National Institutes of Health, and pharmaceutical and biotechnology companies. It is also one of NCI’s efforts in support of the Cancer Moonshot, answering Vice President Biden’s call for greater collaboration and faster development of new therapies for patients. The availability of agents through the NCI Formulary will expedite the start of clinical trials by alleviating the lengthy negotiation process—sometimes up to 18 months—that has been required for investigators to access such agents on their own.

“The NCI Formulary will help researchers begin testing promising drug combinations more quickly, potentially helping patients much sooner,” said NCI Acting Director Douglas Lowy, M.D. “Rather than spending time negotiating agreements, investigators will be able to focus on the important research that can ultimately lead to improved cancer care.”



Credit: iStock

<https://nciformulary.cancer.gov/>

The screenshot shows the NCI Formulary website interface. At the top, there is the NIH logo and the text "NATIONAL CANCER INSTITUTE DCTD Division of Cancer Treatment & Diagnosis". A search bar is located on the right. Below the header, there is a navigation menu with options like "Home", "Participating Companies", "Available Agents", "Participation Eligibility", "Information for Investigators", and "Information for Companies". The main content area is titled "Available Agents" and contains a table with columns for Agent Name, NSC Number, Company, Agent Class, and Agent Target/Molecular Target(s). The table lists several agents including Alectinib, Alectinib, Brevinertib, Cobimetinib, Erlotinib, Ipilimumab, Larotrectinib, LY3009478, Mogamulizumab, Nivolumab, and Ramipril.

Agent Name	NSC Number	Company	Agent Class	Agent Target/Molecular Target(s)
Alectinib	754811	Genentech	ALK inhibitor, tyrosine kinase inhibitor	ALK, KR1
Alectinib	753808	Genentech	PD-1 blocking monoclonal antibody	PD-1
Brevinertib	751605	Genentech	Anti-angiogenesis inhibitor, monoclonal antibody	VEGF
Cobimetinib	751257	Genentech	MEK1/2 inhibitor	MEK1/2
Erlotinib	754723	Xcovery Holding Company LLC	ALK inhibitor	ALK, TRK, TRK, ROS, EphA2, cMET
Ipilimumab	732442	Bristol-Myers Squibb	anti-CTLA-4 monoclonal antibody	CTLA-4
Larotrectinib	755607	Loxo Oncology	Tyrosine kinase inhibitor	NTRK1, NTRK2, NTRK3 fusion proteins, TRKA/B/C proteins
LY3009478		Eli Lilly and Company	Notch inhibitor	Notch
Mogamulizumab	751094	Kyowa Hakko Kirin Co., Ltd.	anti-CCR4 monoclonal antibody	CC chemokine receptor 4 (CCR4)-expressing cells
Nivolumab	746726	Bristol-Myers Squibb	PD-1 blocking monoclonal antibody	PD-1

Participating Companies (as of 01/11/2017)

- Bristol-Myers Squibb
- Eli Lilly and Company
- Genentech
- Kyowa Hakko Kirin Co., Ltd.
- Loxo Oncology
- Xcovery Holding Company LLC



BCRF NEWS
SCIENCE NEWS

PRESS RELEASE: Breast Cancer Research Foundation and Prostate Cancer Foundation Announce Their Commitment to Support the Blood Profiling Atlas in Cancer

In alignment with VP Biden's Cancer Moonshot initiative, the Blood PAC formed in October 2016 to support progress towards patient benefit through research


By BCRF | December 20, 2016



BCRF has awarded a team science grant to **Drs. Shriver and Kuhn from the Department of Defense's Murtha Cancer Center and the University of Southern California**, while PCF is supporting **Dr. Howard I. Scher of Memorial Sloan Kettering Cancer Center (MSKCC) and the Prostate Cancer Clinical Trials Consortium (PCCTC)**.

The funds have been awarded to recognized leaders in biomarker assay validation and are intended to **support pilot projects that will utilize multiple technologies for analyzing rare events in the blood of cancer patients and subsequently deposit the data and associated protocols into the Blood PAC commons.**

INFO@BLOODPAC-DATA.ORG



about members data group tech group sample group

ABOUT

BloodPAC

About

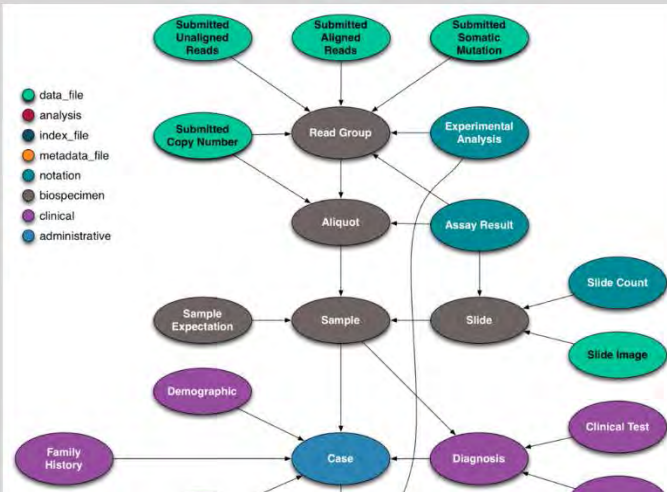
On October 17, 2016, the White House announced the [Blood Profiling Atlas Pilot](#), later renamed to BloodPAC. The original fact sheet can be found [here](#).

Overview

Data Matrices

Requires username and password.

- [Data Submission Counts Matrix](#)
- [Detailed Project Matrix](#)



Data Model

To facilitate a common set of standards and the use of tools for analysis, a data model has been developed for this project. We will work to make the data model as close to the Genomic Data Commons (gdc.cancer.gov) data model and as compliant to the GA4GH standards as practical. <https://github.com/occc-data/bpdictionary>

- [Overview of draft BPA data model v0.5](#)
- [Overview of draft BPA data model v0.4](#)
- [Overview of draft BPA data model v0.3](#)
- [Overview of draft BPA data model v0.2](#)
- [Overview of draft BPA data model v0.1](#)
- [Browse the GDC data model](#)
- [Examine the GDC data model in detail](#)
- [BPA Model in Github](#)
- [Clinical Data Harmonization](#)
- [Suggested template for brief experimental descriptions](#)



Lauren Leiman
 Executive Director
lauren@bloodpac.org

Updated: December 12, 2016

The screenshot shows the NIH National Cancer Institute website. The header includes the NIH logo and the text "NATIONAL CANCER INSTITUTE". Below the header is a navigation bar with links for "1-800-4-CANCER", "Live Chat", "Publications", and "Dictionary". A secondary navigation bar contains "ABOUT CANCER", "CANCER TYPES", "RESEARCH", "GRANTS & TRAINING", "NEWS & EVENTS", "ABOUT NCI", and a search box. The breadcrumb trail reads "Home > Research > Key Initiatives > Cancer Moonshot® > Blue Ribbon Panel Report". The main content area features a sidebar on the left with a "CANCER MOONSHOT®" section containing links for "Blue Ribbon Panel Report", "BRP Recommendations", "About the Panel", "Working Groups", "Funding Opportunities", and "Milestones". The main heading is "Blue Ribbon Panel Recommendations - Funding Opportunities". The text below the heading states: "At the end of 2016, NCI took the preliminary step of identifying current cancer research areas closely aligned with the BRP recommendations that could be leveraged with existing funds and with Cancer MoonshotSM funds. NCI identified six such areas, listed below, that are currently open for applications. Researchers with an interest in the areas identified in the BRP recommendations are encouraged to apply for these programs." A second paragraph follows: "These preliminary opportunities mark the beginning of the implementation phase of the Blue Ribbon Panel recommendations. These and future BRP funding opportunities will constitute part of the growing Blue Ribbon Panel portfolio." To the right of the text is a dark blue graphic with the text "FUNDING OPPORTUNITIES" in large white letters, "BLUE RIBBON PANEL RECOMMENDATIONS" in smaller white letters, and the "CANCER MOONSHOT" logo at the bottom.

Currently **9** opportunities addressing **4** BRP recommendations

Please check back regularly as additional Funding Opportunity Announcements are posted.

<https://www.cancer.gov/research/key-initiatives/moonshot-cancer-initiative/blue-ribbon-panel/funding>

iPhone (EDGE, 16 GB max)



1/9/2007
(~10 yrs old)

iPhone5 (LTE, 128 GB max)



9/12/2012
(~15 yrs old)

iPod (10GB max)



10/23/2001
(~5 yrs old)

iPad (EDGE, 64 GB max)



4/3/2010
(~13 yrs old)

**Google
Baseline**
7/14/2014
(~17 yrs old)

WinAMP(mp3)



4/21/1997



2/7/2007



7/15/2006

facebook

9/26/2006
(~9 yrs old)

**iPhone 3G
(16 GB max)**



7/11/2008
(~11 yrs old)



**Google
Drive**



4/24/2012
(~15 yrs old)



3/9/2015
(~18 yrs old)

802.11b WiFi



9/16/1999
(~3 yrs old)



4/23/2005
(~8 yrs old)



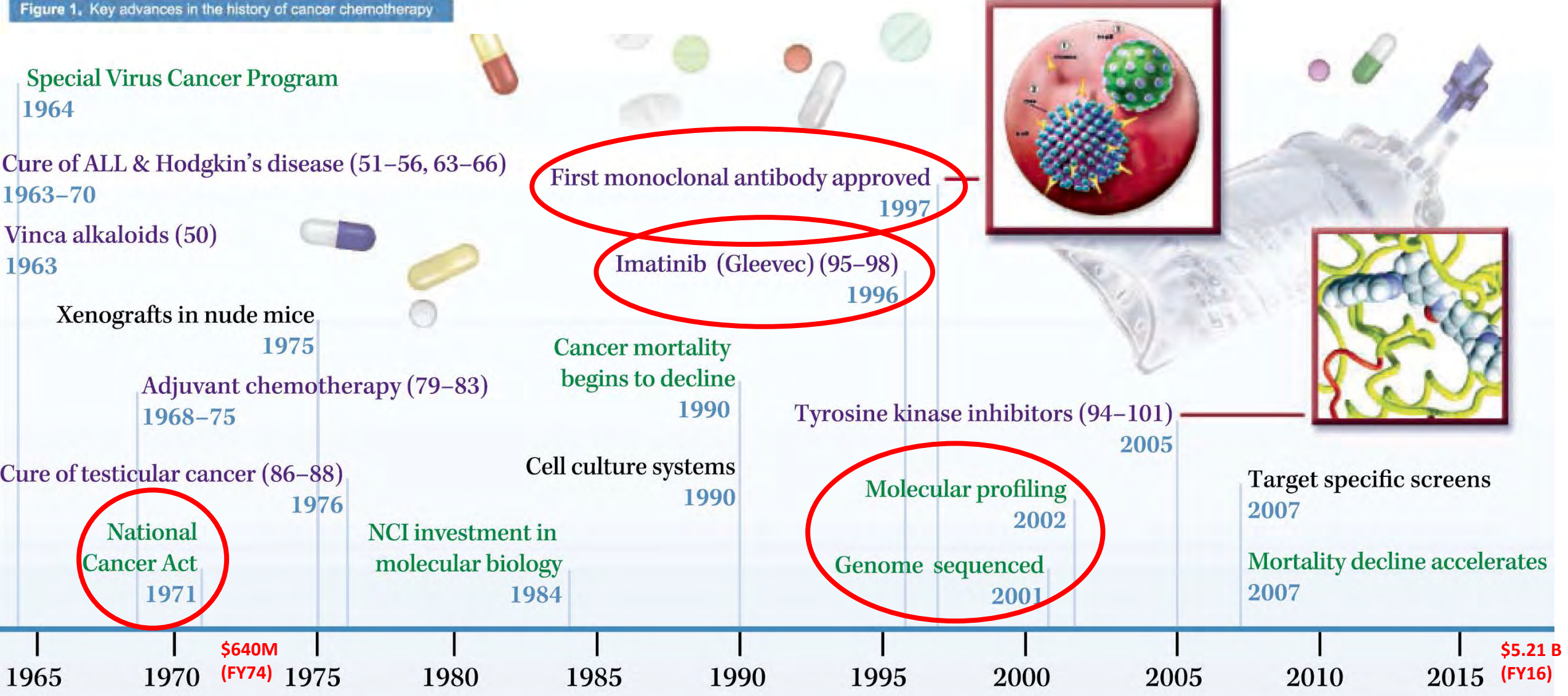
1997 1998 1999 2000 2001 2002 2003 2004 2005 2006 2007 2008 2009 2010 2011 2012 2013 2014 2015

A History of Cancer Chemotherapy

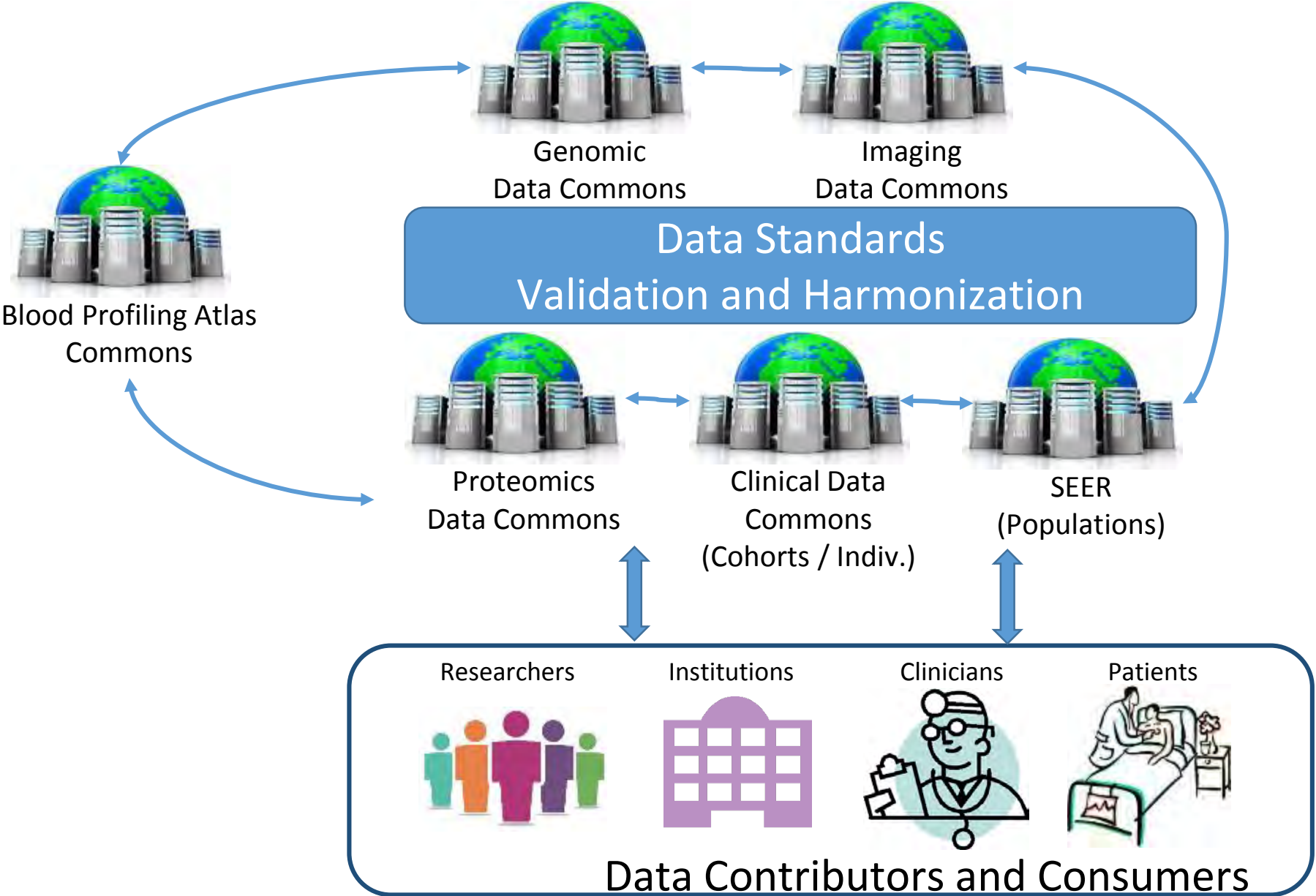
Vincent T. DeVita, Jr. and Edward Chu

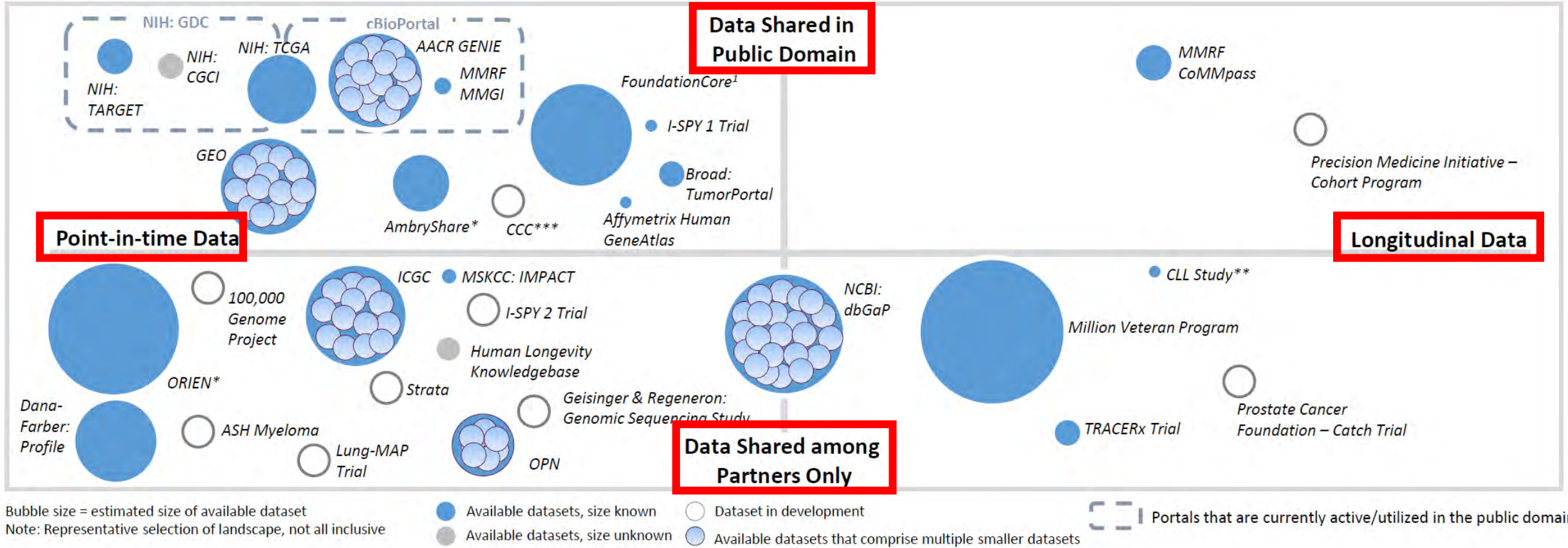
Yale Cancer Center, Yale University School of Medicine, New Haven Connecticut

Figure 1. Key advances in the history of cancer chemotherapy



Cancer Research Data Commons Ecosystem





Opportunity exists to generate publicly available longitudinal data to drive understanding of genetic mutations and find Precision Medicine cures

*Datasets have potential to include longitudinal data in the future

**Public/private information not available

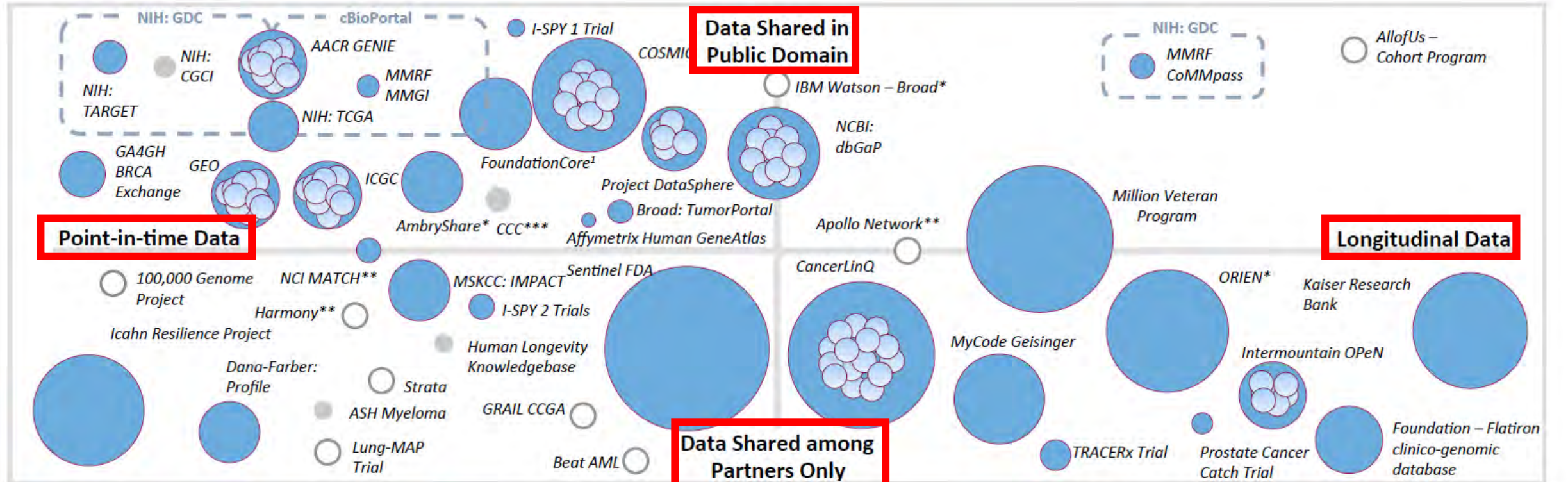
***Serves as a portal also, has potential to include longitudinal data in the future

1. FoundationCore's pediatric cancer data has been made public



Oncology Precision Medicine Data Landscape:

December 2016 Update



Bubble size = estimated size of current dataset
 Note: Representative selection of landscape, not all inclusive
 Note: Dataset selection focused on oncology specific and oncology-inclusive datasets

- Available datasets, size known
- Data initiative announced. Recruiting planned
- Available datasets, size unknown
- Independent initiatives as part of larger dataset

Portals that are currently active/used in the public domain

Opportunity exists to generate publicly available longitudinal data to drive understanding of genetic mutations and find Precision Medicine cures

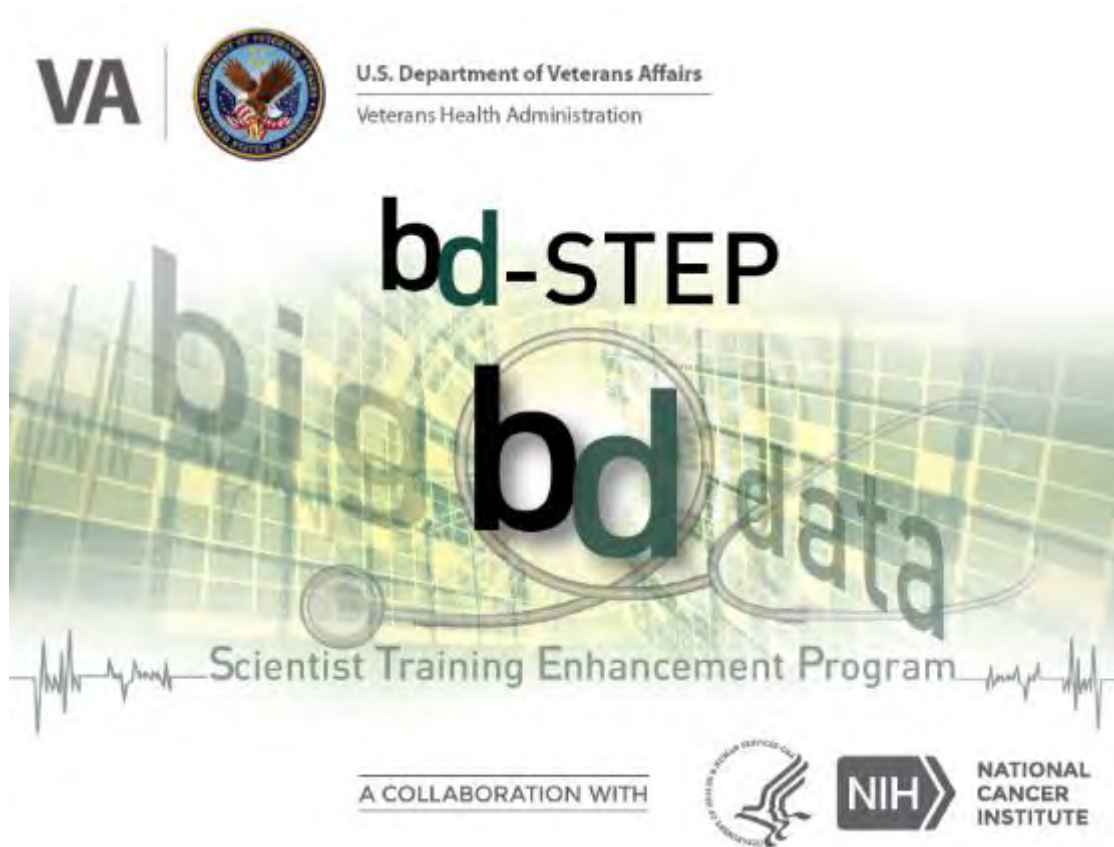
*Datasets have potential to include longitudinal data in the future

**Public/private information not available

† ***Serves as a portal also, has potential to include longitudinal data in the future

1. FoundationCore's pediatric cancer data has been made public

Big Data Scientist Training Enhancement Program (BD-STEP)



2017 Potential
Partners:



Graduates of BD-STEP would:

- have skillsets to perform next-generation **patient-centered outcomes research** by manipulating and analyzing large-scale, multi-element, patient data sets to **develop novel disease signatures or unique performance-based clinical benchmarks**
- have an understanding of real-time, performance-driven health care delivery in the VA systems

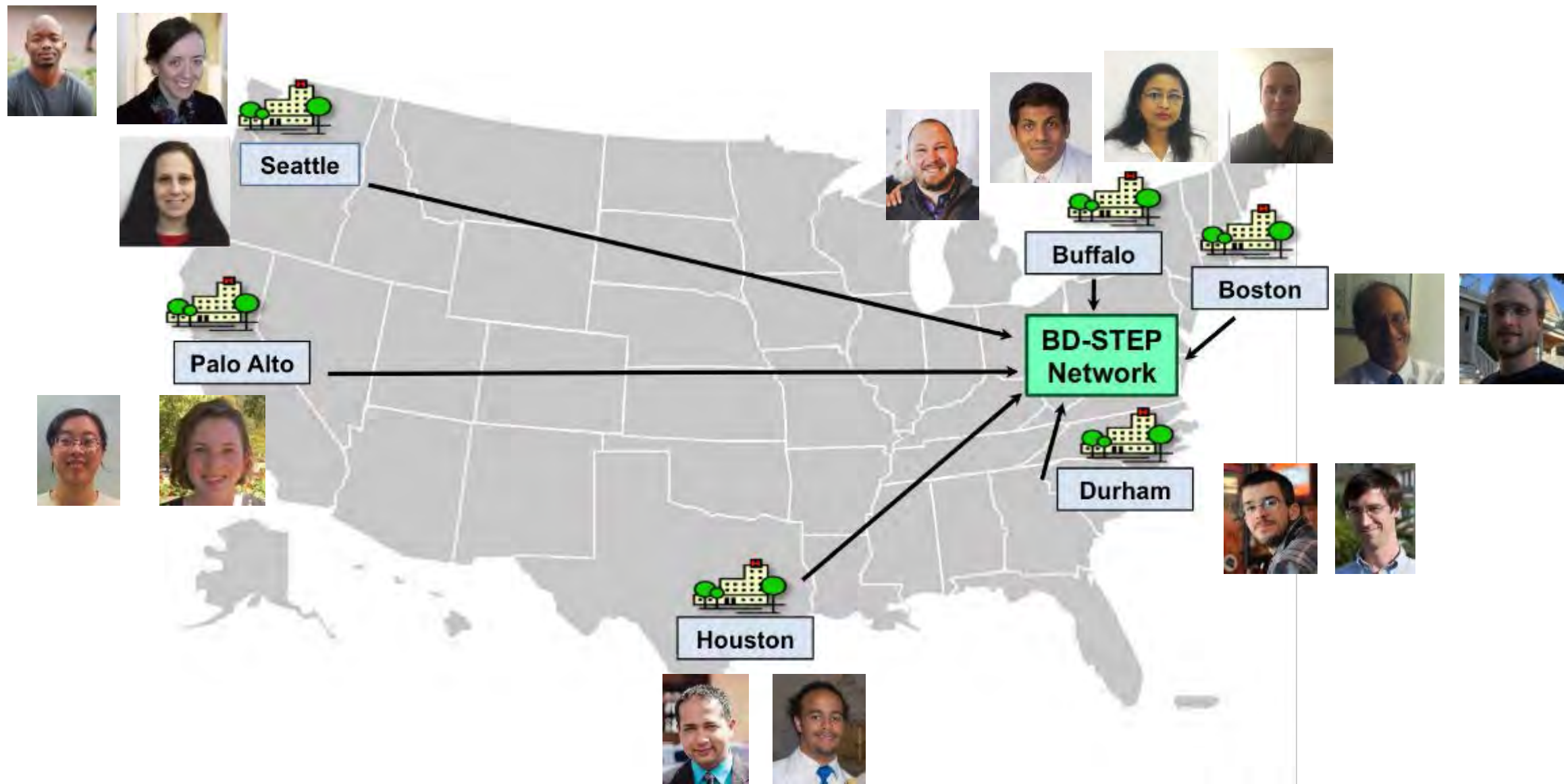


Connie Lee, VHA/EES

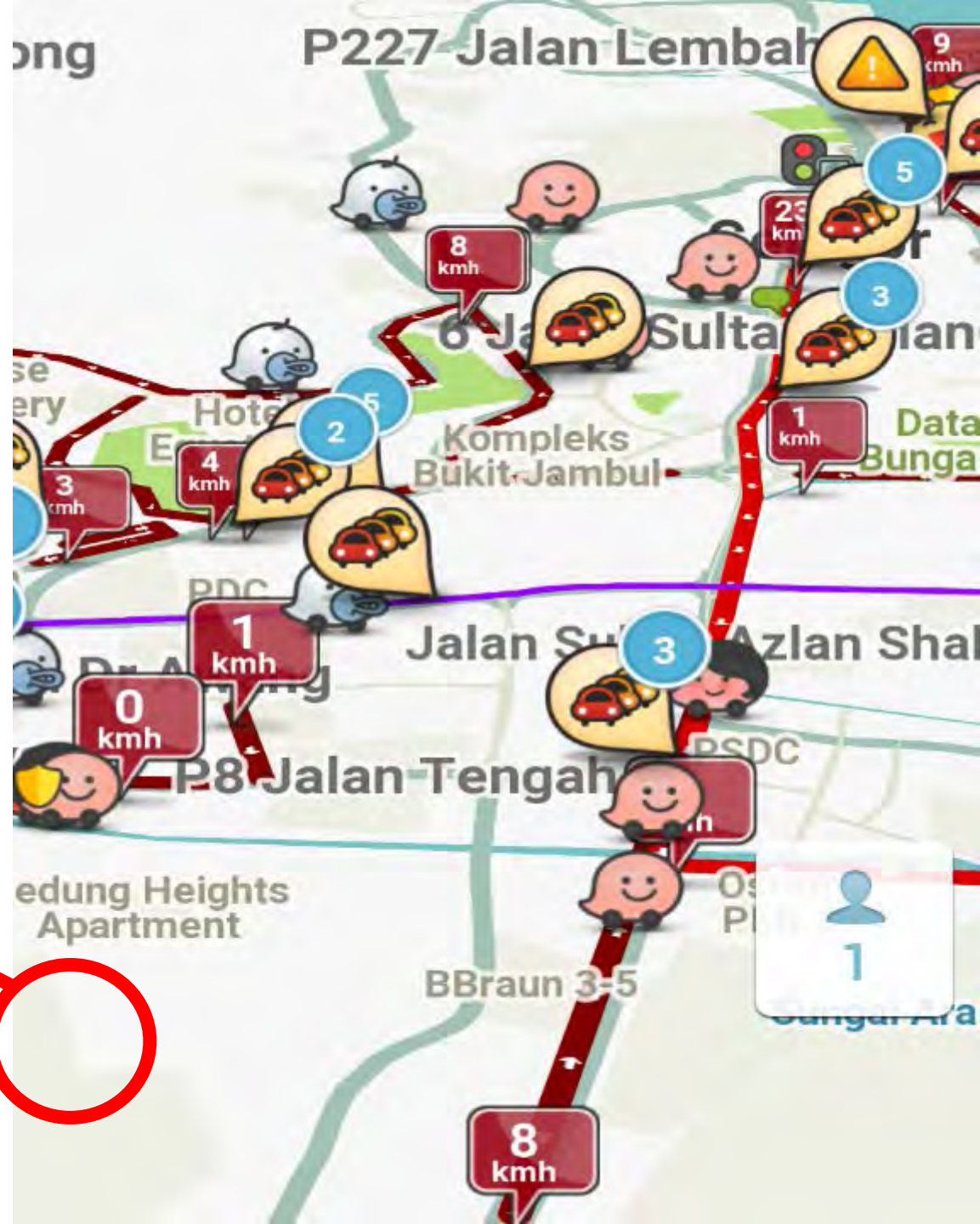
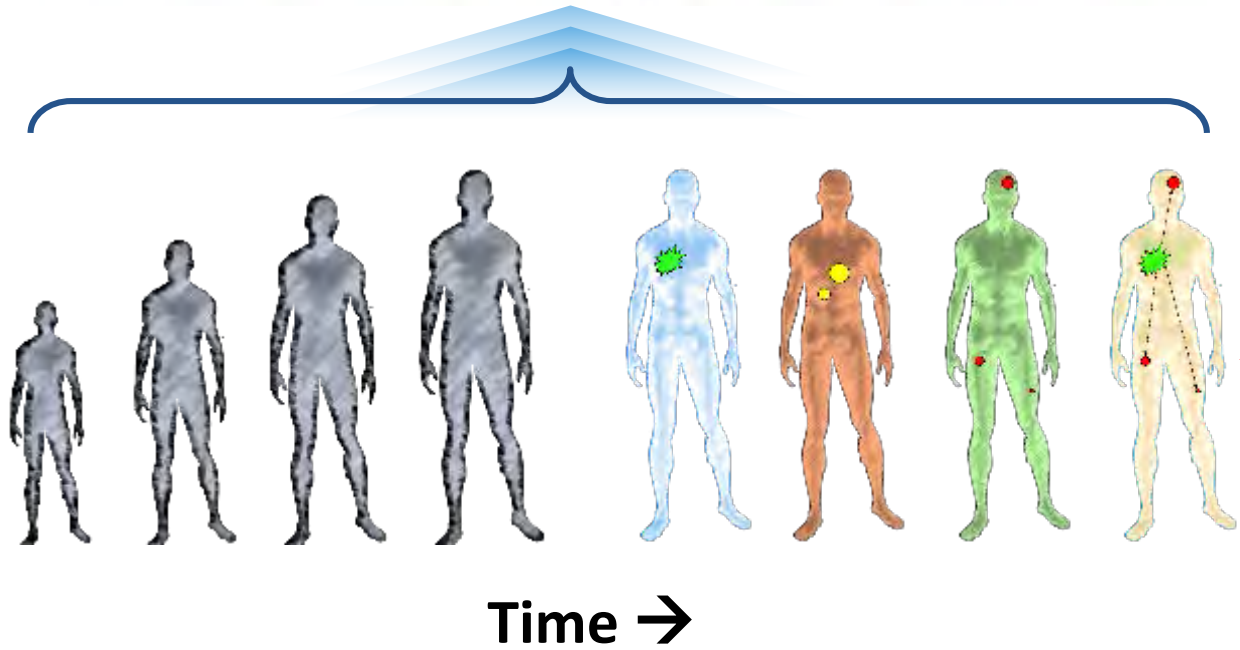
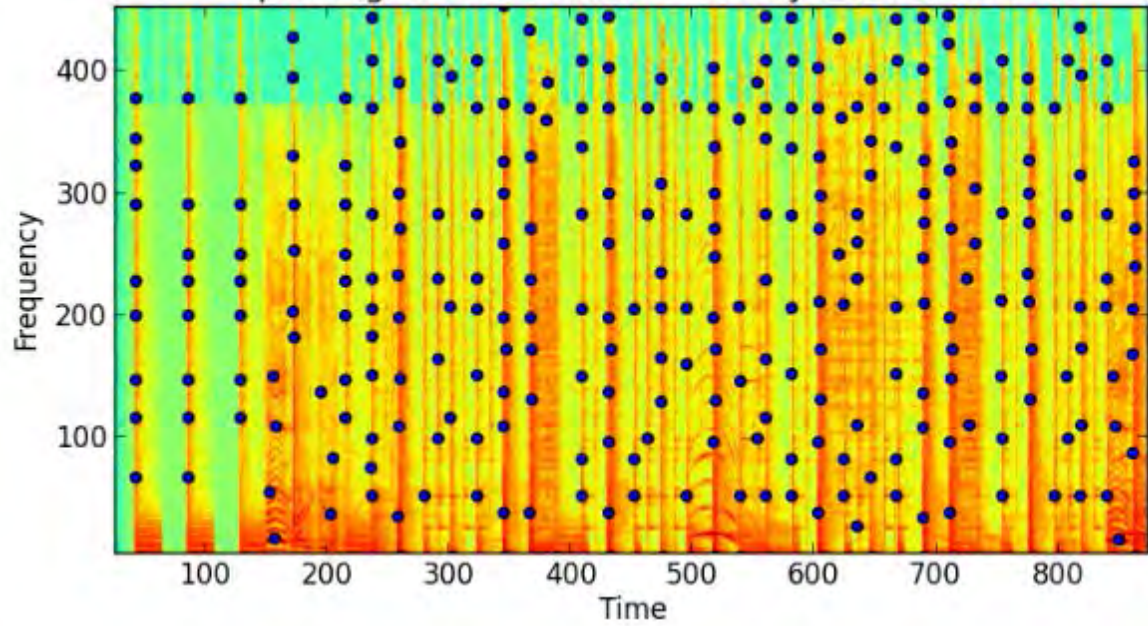


Michelle Berny-Lang, NCI

BD-STEP Sites and Fellows: 2016-2017



Spectrogram of "Blurred Lines" by Robin Thicke





CancerBase

Share, Discover & Change

Join / Login with



Or with email

Inviting those with cancer to join a global effort to make our data count.
When patients share real-time data with patients, the world changes.

<http://www.cancerbase.org>

Our worldwide map of real-time information.

Each dot is one of us. The clickable tabs for the different cancer types will light up when enough of us with that particular cancer have joined.



Acknowledgements/Thanks to the “Secret Ingredients”



Clinical Sciences

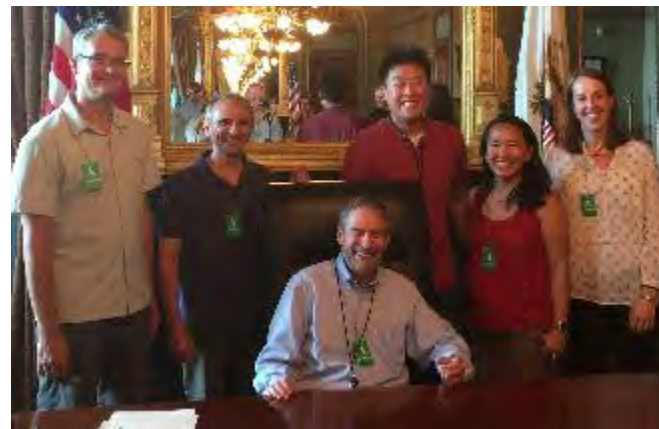


Life Sciences



Physical Sciences





Learn More About Us...

<http://cssi.cancer.gov>

The screenshot shows the top of the website with the National Cancer Institute logo and the text "National Cancer Institute" and "U.S. National Institutes of Health | www.cancer.gov". Below this is the "CENTER for STRATEGIC SCIENTIFIC INITIATIVES" logo and a navigation menu with "HOME", "ABOUT CSSI", "CSSI OFFICES", and "CONTACT CSSI". The main content area features a large image of a DNA double helix and the text: "ENABLING PROGRESS IN CANCER RESEARCH THROUGH ADVANCED TECHNOLOGIES, TRANS-DISCIPLINARY PROGRAMS AND RESOURCES". A "LEARN MORE" button is visible at the bottom of this section.

Jerry S.H. Lee, PhD
jerry.lee@nih.gov

The screenshot shows the "Timeline" section of the website. It includes a legend for "Offices" with icons for various departments like NCI, NCI/NIH, etc. Below the legend is a "Zoom Level" selector with options for "1 Year", "5 Years", and "All Time". The main part of the screenshot is a circular diagram with various icons representing different research areas or projects, arranged in a circular pattern.

The screenshot shows the "RESOURCES" section of the website. It features a "Current Funding Opportunities and Resources" table. The table has columns for "Research, Programs & Centers" and "Specialized Opportunities". The "Research, Programs & Centers" column is highlighted in red. Below the table, there are links for "Research, Programs & Centers" and "Specialized Opportunities".

Research, Programs & Centers	Specialized Opportunities
Research, Programs & Centers	Specialized Opportunities