

Biomedical data sharing to enable Learning Health Systems

Jonathan C. Silverstein, MD, MS, FACS, FACMI

Chief Research Informatics Officer,

Health Sciences and Institute of Precision Medicine

Visiting Professor, Department of Biomedical Informatics

Affiliate Scholar, Pitt Cyber

University of Pittsburgh

U.S. healthcare challenges in a slide?

- People are dying of preventable causes.
- Cost is out of control.
- Quality can't be measured.
- Variability is local and widespread.
- New technology is exponentiating.
- Decision-making is maximally distributed.
- Data is not available routinely for learning.



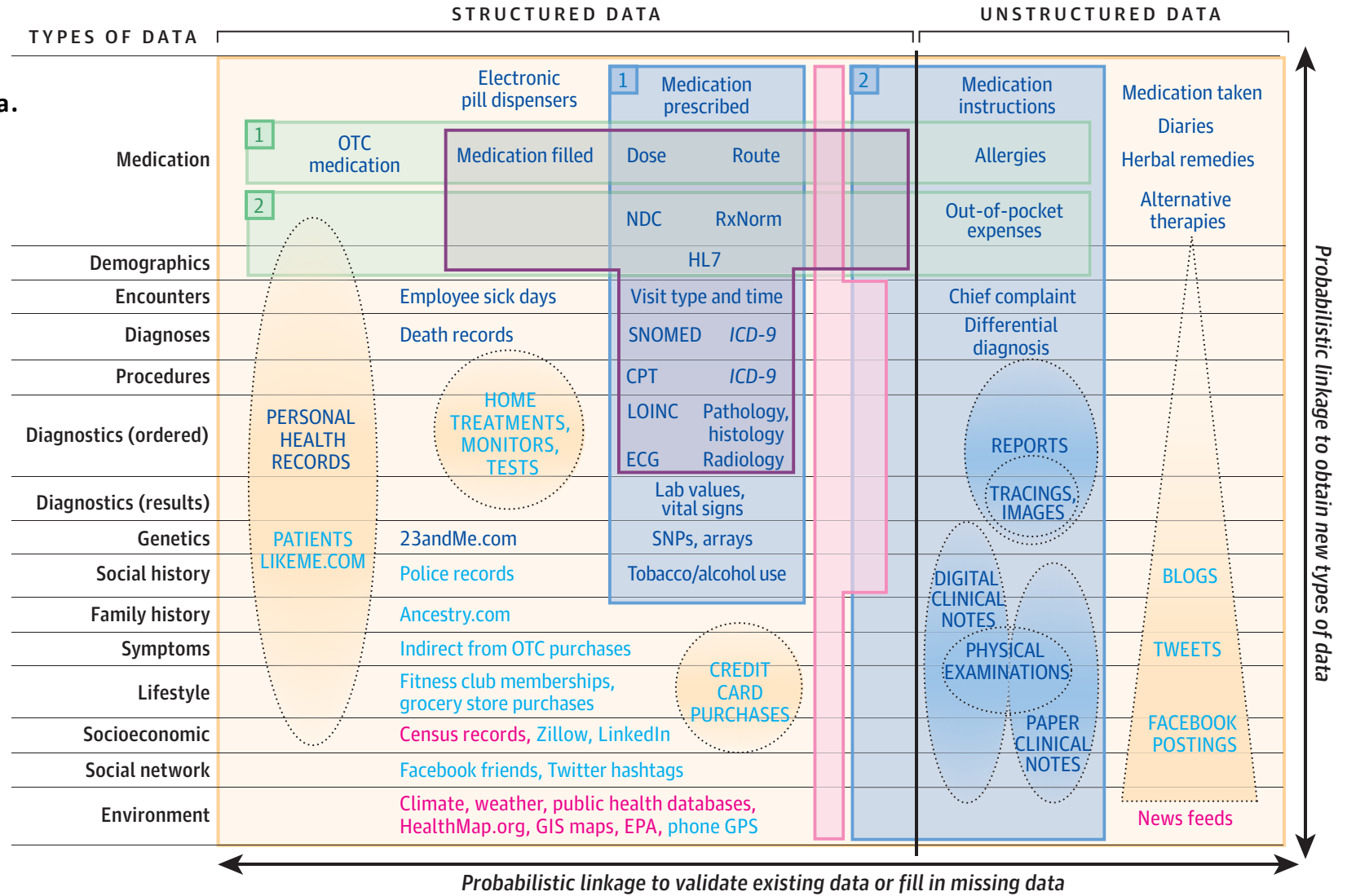
Figure. The Tapestry of Potentially High-Value Information Sources That May be Linked to an Individual for Use in Health Care











Weber GM, Mandl KD, Kohane IS.

Finding the Missing Link for Big Biomedical Data.

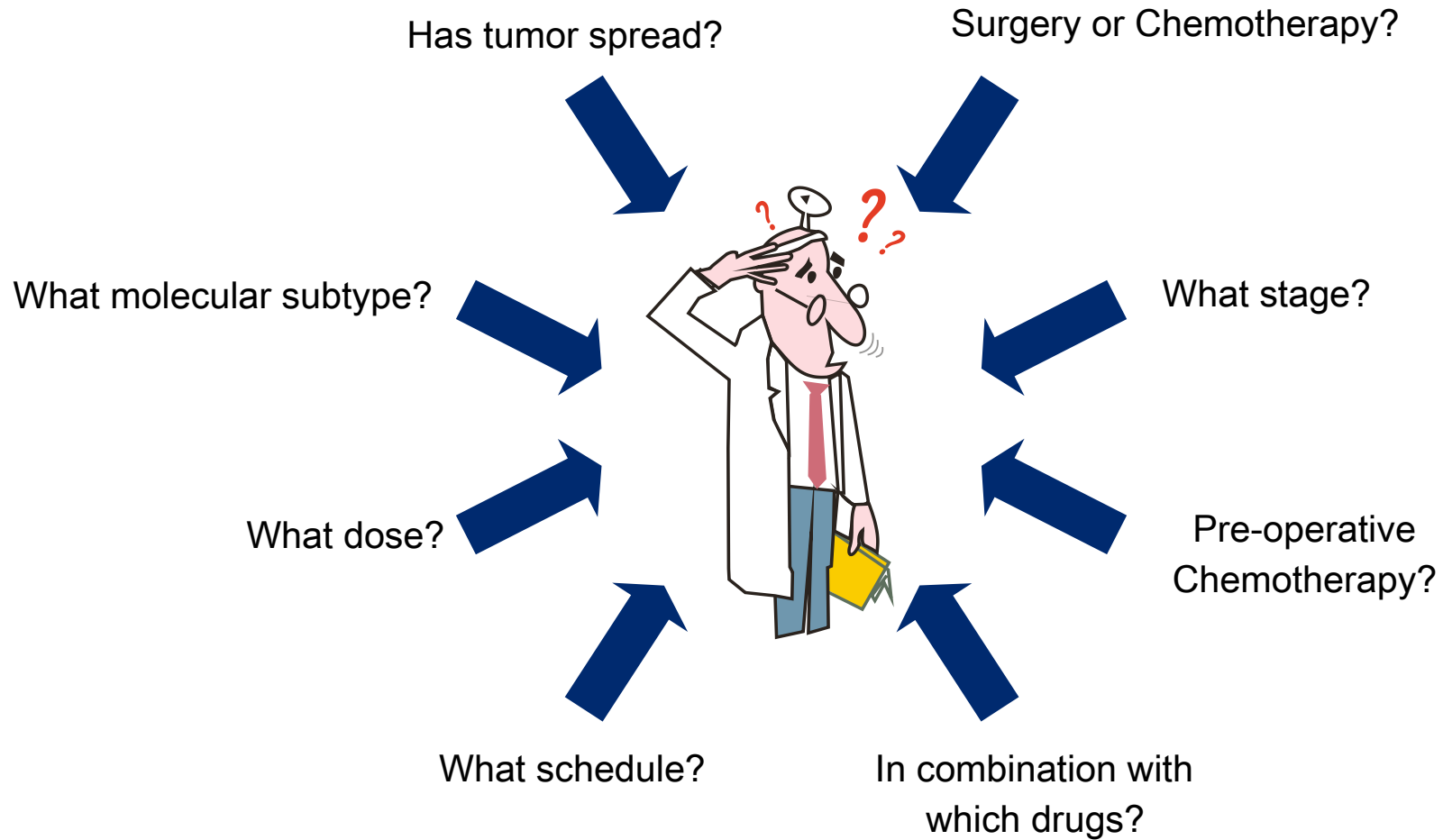
JAMA. 2014 Jun 25;311(24):2479–2480.

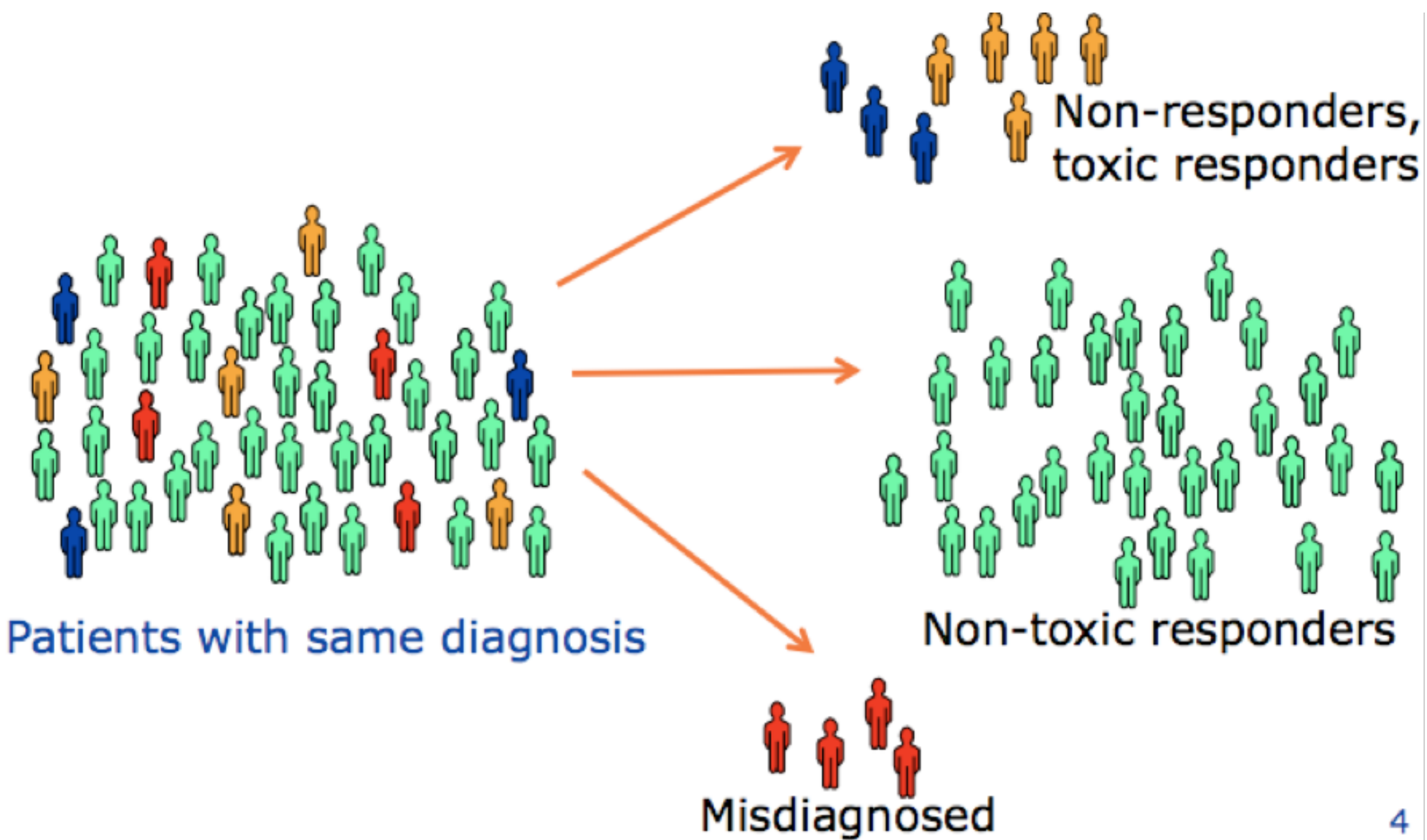
PMID: 24854141



Examples of biomedical data		Ability to link data to an individual	Data quantity
 Pharmacy data	 Health care center (electronic health record) data	 Easier to link to individuals	 More
 Claims data	 Registry or clinical trial data	 Harder to link to individuals	 Less
 Data outside of health care system		 Only aggregate data exists	

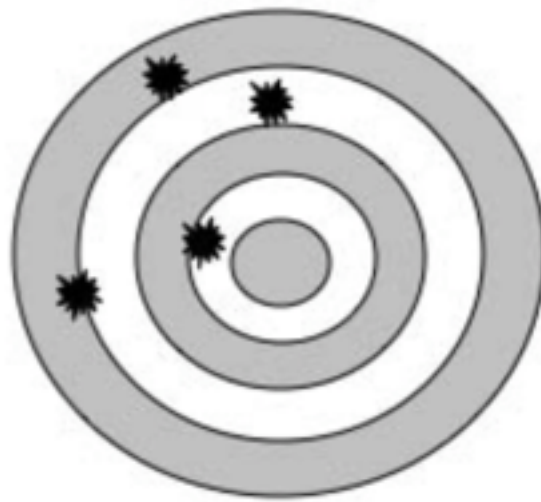
Provider, Patient & Payor Faced With Bewildering Choices: The Current Practice of “Qualitative” Medicine



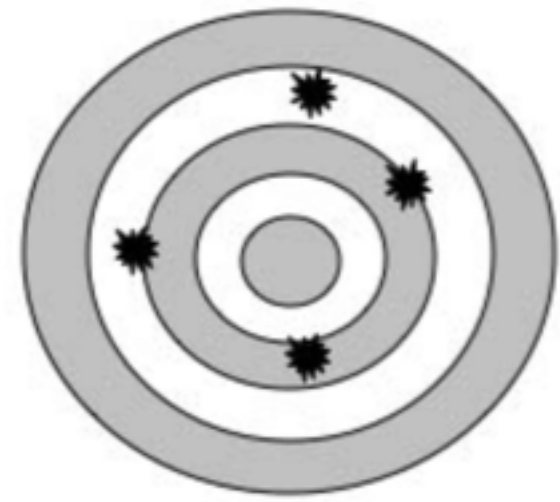


NOAA.gov

https://celebrating200years.noaa.gov/magazine/tct/accuracy_vs_precision.html



**Not Accurate
Not Precise**



**Accurate
Not Precise**



**Not Accurate
Precise**



**Accurate
Precise**

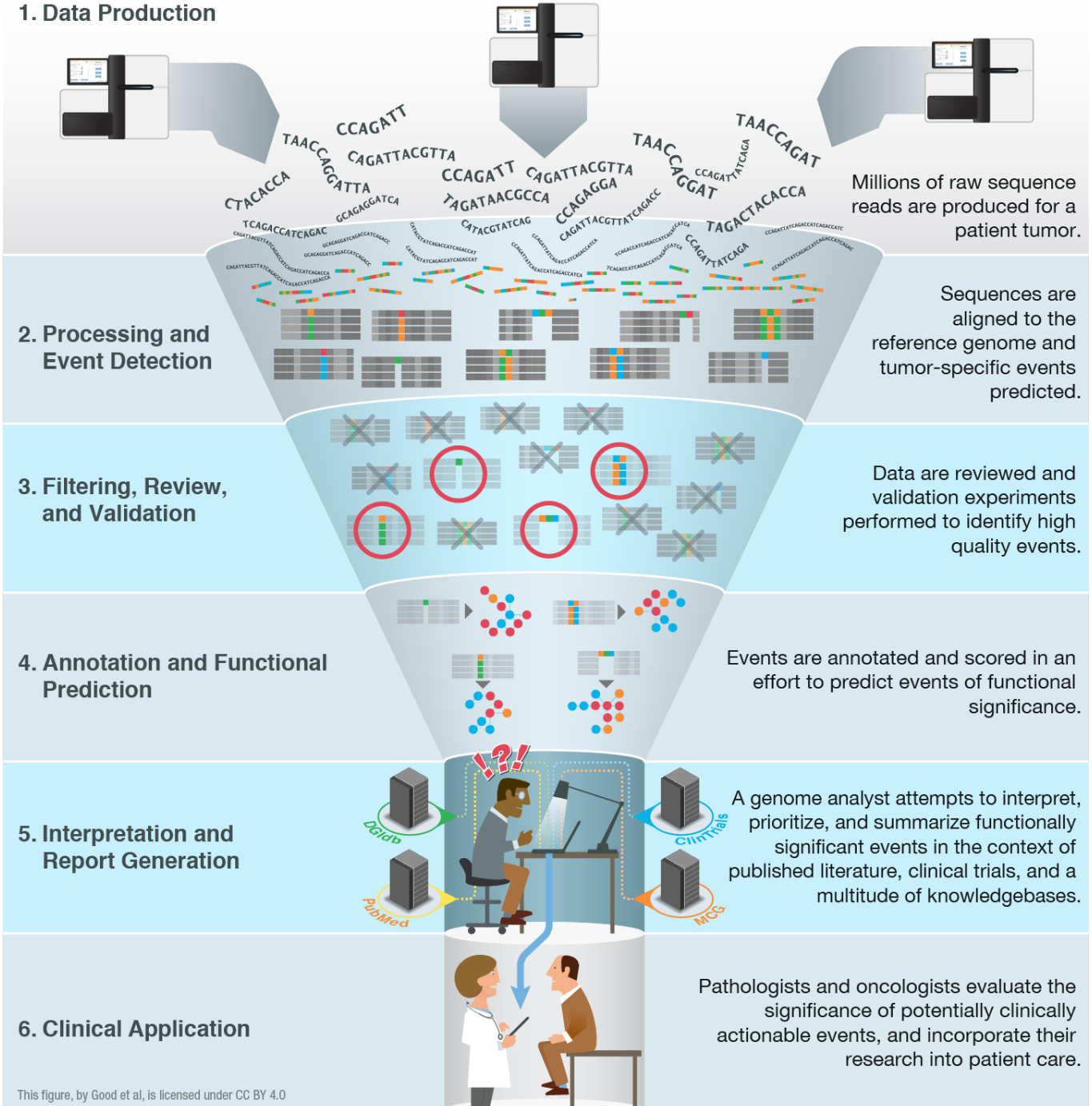
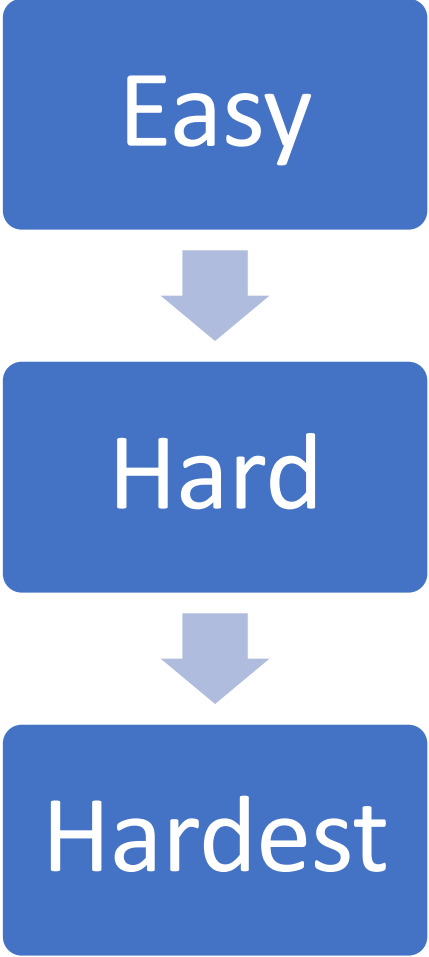
Good BM, Ainscough BJ, McMichael JF, Su AI, Griffith OL.

Organizing knowledge to enable personalization of medicine in cancer.

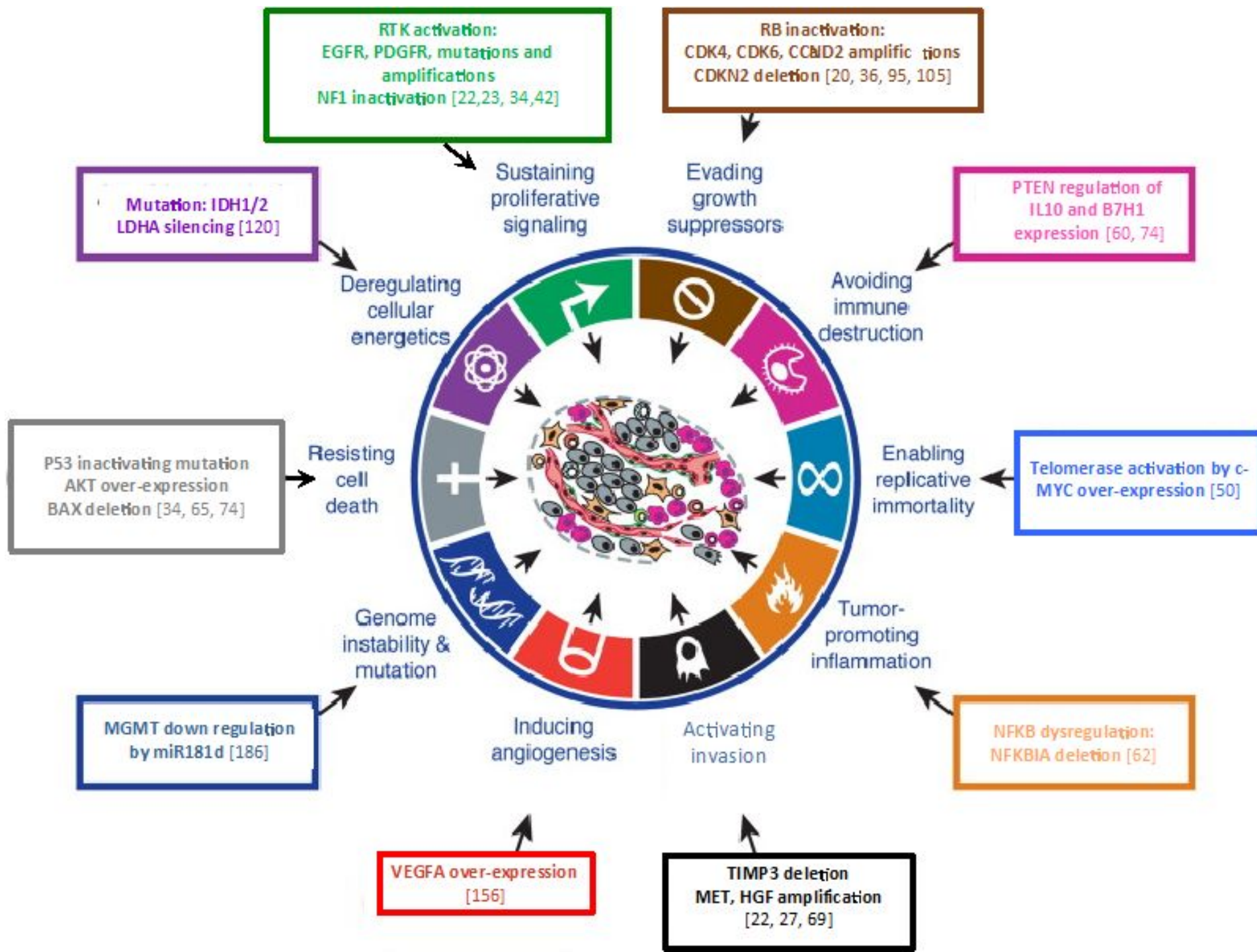
Genome Biol. 2014

Aug 27;15(8):438.

PMCID: PMC4281950



This figure, by Good et al, is licensed under CC BY 4.0



The Promise of Personalized Medicine

- Accelerate drug development, biomarker discovery, and guide diagnosis, treatment, and prevention
 - Detect disease at an earlier stage, when it is easier to treat effectively
 - Shift practice from reaction to prevention
 - Reduce the overall cost of healthcare
-
- (credit Rebecca Crowley Jacobson, VP, UPMC Enterprises)

PRESS RELEASES

NCI Press Release

NCI-MATCH precision medicine clinical trial releases new findings, strengthens path forward for targeted cancer therapies

Posted: June 4, 2018

Contact: NCI Press Office
240-760-6600

The National Cancer Institute's Molecular Analysis for Therapy Choice (NCI-MATCH) trial, the largest precision medicine trial of its kind, has achieved a milestone with the release of results from several treatment arms, or sub-studies, of the trial. The new results offer findings of interest for future cancer research that could ultimately play a role in bringing targeted treatments to patients with certain gene abnormalities, regardless of their cancer type.

Findings from three arms were released at this year's American Society of Clinical Oncology (ASCO) annual meeting in Chicago, adding to [findings from one arm released in November 2017](#). The study was co-developed by NCI, part of the National Institutes of Health, and the ECOG-ACRIN Cancer Research Group, part of the NCI-sponsored National Clinical Trials Network (NCTN). ECOG-ACRIN and NCI are co-leading the trial.

"The outcomes data being released today from this groundbreaking precision medicine trial are an exciting step for NCI-MATCH," said Lyndsay Harris, M.D., of NCI's Cancer Diagnosis Program and NCI study chair. "These findings represent a large collection of data in populations of patients who may not have been studied in conventional clinical trials, and they will have important implications for future precision medicine trials."



Credit: National Cancer Institute

FDA Approves Foundation Medicine's FoundationOne CDx™, the First and Only Comprehensive Genomic Profiling Test for All Solid Tumors Incorporating Multiple Companion Diagnostics

--Landmark approval advances personalized cancer care as an estimated 1 in 3 patients across five common advanced cancers are expected to match with an FDA-approved therapy--

--The Centers for Medicare and Medicaid Services issued a preliminary National Coverage Determination (NCD) for FoundationOne CDx, improving access to molecular information for personalized healthcare--

CAMBRIDGE, Mass.--(BUSINESS WIRE)-- **Foundation Medicine, Inc.** (NASDAQ:FMI) today announced that the U.S. Food and Drug Administration (FDA) approved FoundationOne CDx™, the company's comprehensive companion diagnostic test for solid tumors. FoundationOne CDx is intended for use by health care professionals to help inform cancer treatment management in accordance with professional guidelines for patients with solid tumors. The first and only FDA-approved test of its kind for all solid tumors, FoundationOne CDx is a diagnostic test that acts as:

This press release features multimedia. View the full release here: <http://www.businesswire.com/news/home/20171130006320/en/>



Foundation Medicine is headquartered in Cambridge, MA (Photo: Business Wire)

- a comprehensive companion diagnostic to identify patients who may benefit from treatment with specific FDA-approved targeted therapies;
- a comprehensive genomic profiling (CGP) test that includes genomic biomarkers to help inform the use of other targeted oncology therapies, including immunotherapies;
- a tool for physicians that identifies patient opportunities for clinical trial participation; and,
- an FDA-approved platform for companion diagnostic development for biopharma companies developing precision therapeutics.

FoundationOne CDx assesses all classes of genomic alterations in 324 genes known to drive cancer growth, providing potentially actionable information to help guide treatment decisions. It is also indicated as a companion diagnostic for patients with certain types of non-small cell lung cancer (NSCLC), melanoma, colorectal cancer, ovarian cancer or breast cancer to identify those patients who may benefit from treatment with one of 17 on-label targeted therapies, including 12 therapies currently approved as first-line therapy for their respective indications. FoundationOne CDx also reports genomic biomarkers, such as microsatellite instability (MSI) and tumor mutational burden (TMB), that can help inform the use of immunotherapies; genomic alterations in other genes relevant to patient management; and, relevant clinical trial information.

Based on previous CGP testing conducted by Foundation Medicine, it is estimated that approximately 1 in 3 patients across five common advanced cancers are expected to match with an FDA approved therapy.¹

The number of matched on-label therapies indicated on FoundationOne CDx is expected to increase over time as Foundation Medicine and its biopharma partners pursue FDA approval for additional companion diagnostics on the platform. Today, approximately 50% of new cancer drugs in development are projected to have a companion biomarker.²

News Releases



CancerLinQ Engages Leading Technology Companies
Tempus and Precision HealthAI to Accelerate Data-Driven
Insights to Oncologists and the Cancer Care Community

FOR IMMEDIATE RELEASE

December 21, 2017

"T"EMPUS

Data-driven cancer treatment



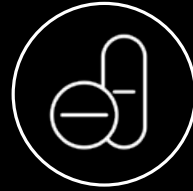
Genomic
Alterations



Targeted
Therapy



Clinical
Trials



Immunotherapy
Options



Patient
Cohort

Patient Cohort aggregates deidentified clinical, genomic, and outcomes data from previously tested patients, allowing you to evaluate treatment regimens for your patients with similar clinical and genomic presentation.

The future includes even more data


- Sequencing of entire exome and entire genome
- Sequencing of individual tumor cells
- Detection of tumor sequence fragments in blood
- Sequencing of multiple areas of a tumor
- Sequencing of metastases and recurrence
- Assembling more integrative analyses across DNA, RNA, protein
- Algorithms to help us untangle the complex molecular changes to find the drugable targets
- (credit Rebecca Crowley Jacobson, VP, UPMC Enterprises)

The Future of Medicine

- Evidence based (data driven)
- Practice based (generation of data)
- Targeted and precise
 - Personalization to individual mutations
 - AI/ML to specific vectors/features
- A Learning Health System
 - “...gets the right care to people when they need it and then captures the results for improvement...” Institute of Medicine/National Academy of Medicine



"We seek the development of **a learning health system** that is designed to generate and apply the best evidence for the collaborative healthcare choices of each patient and provider; **to drive the process of discovery as a natural outgrowth of patient care**; and to ensure innovation, quality, safety, and value in health care."




THE LEARNING HEALTH SYSTEM SERIES

ROUNDTABLE ON VALUE & SCIENCE-DRIVEN HEALTH CARE

DIGITAL INFRASTRUCTURE FOR
THE LEARNING HEALTH SYSTEM

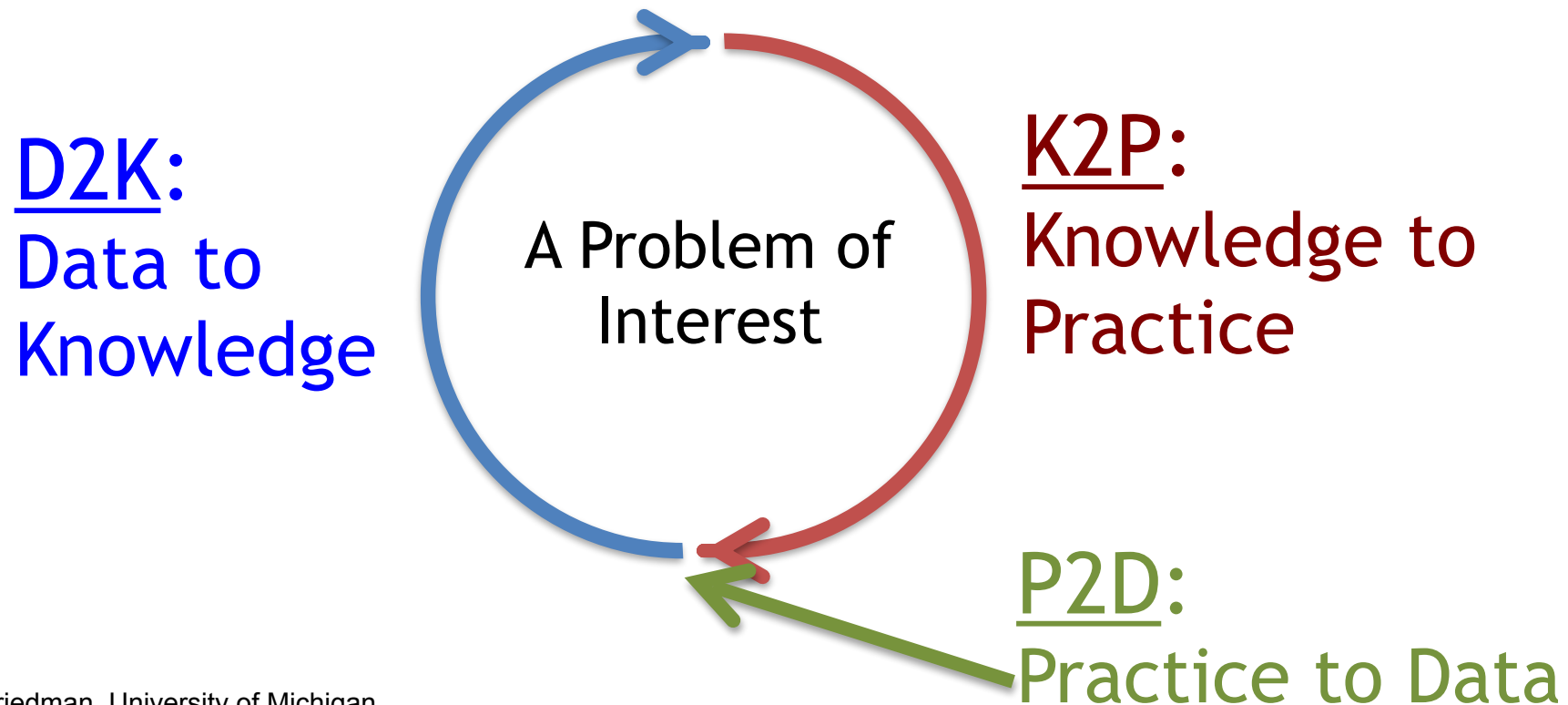
The Foundation for Continuous Improvement
in Health and Health Care

Workshop Series Summary



The LHS Links Discovery to Better Health











Better Health = [D2K] [K2P] [P2D]



Checklist View: Properties of a Health System That Can Learn

- ✓ Every patient's characteristics and experiences are available to **learn** from
- ✓ Best practice knowledge is **immediately available** to support decisions
- ✓ Improvement is **continuous** through ongoing study
- ✓ An **infrastructure** enables this to happen routinely and with economy of scale
- ✓ All of this is part of the **culture**

Core Values of the Learning Health System

 1) Person Focused	 2) Privacy
3) Inclusiveness 	4) Transparency 
 5) Accessibility	 6) Adaptability
7) Governance 	<small>Cooperative & Participatory</small> 8) Leadership 
 9) Scientific Integrity	 10) Value

Learning Health Systems **at Scale:** Research Coupled with Impact

- Scalable expansion of data collection and use
- Collaboration over
 - Millions of patients with
 - New outcomes techniques
- Automated feature modeling
- High-performance analysis
 - clinical, cost, sensor (phone),
 - imaging, and omics
- Tight clinical integration
- From theory and experimentation to observation and simulation, toward a learning health system: “...enables discovery [and innovation] as a natural outgrowth of patient care...” – NAM (formerly IOM)
- Quality Improvement, Decision Support, Population Health, Cost, Safety, Hardening, Personalization and Commercialization

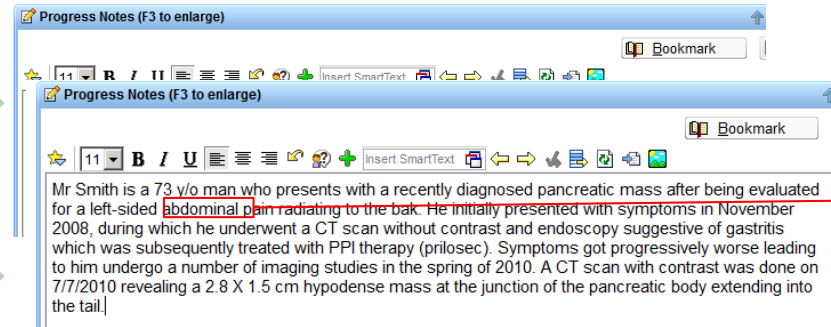
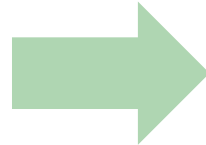
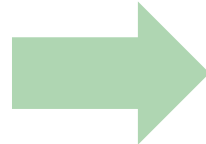
LIFESTYLE

ENVIRONMENT

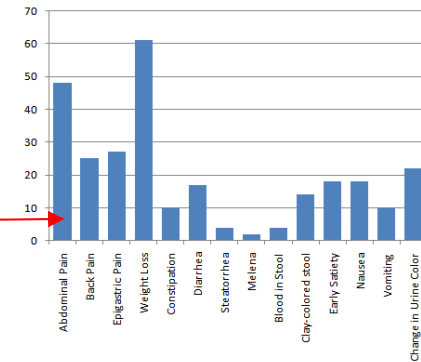
CLINICAL

GENOMIC

Structured Clinical Documentation: Statement of Problem

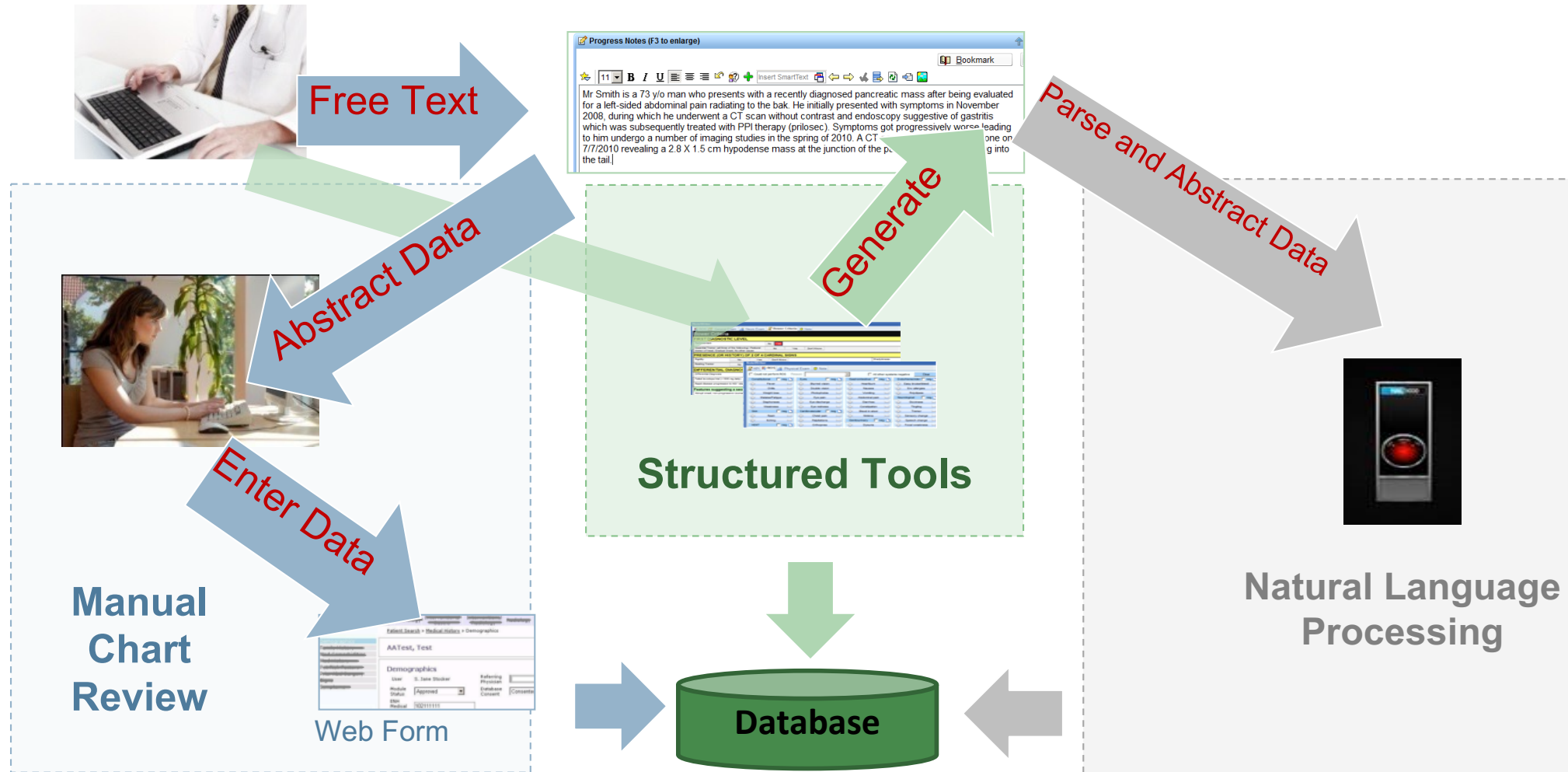


Prevalance of Selected Patient-Reported Symptoms, 309 visits



- Clinical documentation is a rich source of information on interactions between the health system and individual patients.
- Question: How can we capture this information **Consistently and Completely** for analysis—especially the interesting parts of progress notes?
- Answer: Tools Balance Expressivity and Workflow

Three Different Approaches



Research Informatics Office (RIO)

- Mission
 - “to support investigators through innovative collection and use of biomedical data”
- Science-as-a-Service
 - Health Record Research Request (R3)
 - PaTH Network (PCORI CDRN)
 - NMVB, TCRN, Cancer Registry, PGRR
 - UPMC, Enterprises, IPM and PSC relationships
 - Delivery, Help Desk: REDCap, Neptune, ACT, AoU, etc.



Neptune

Research Data
Warehouse

R3

 Honest Broker
Service

PaTH Network
patient empowered research



All of UsSM
THE FUTURE OF HEALTH BEGINS WITH YOU

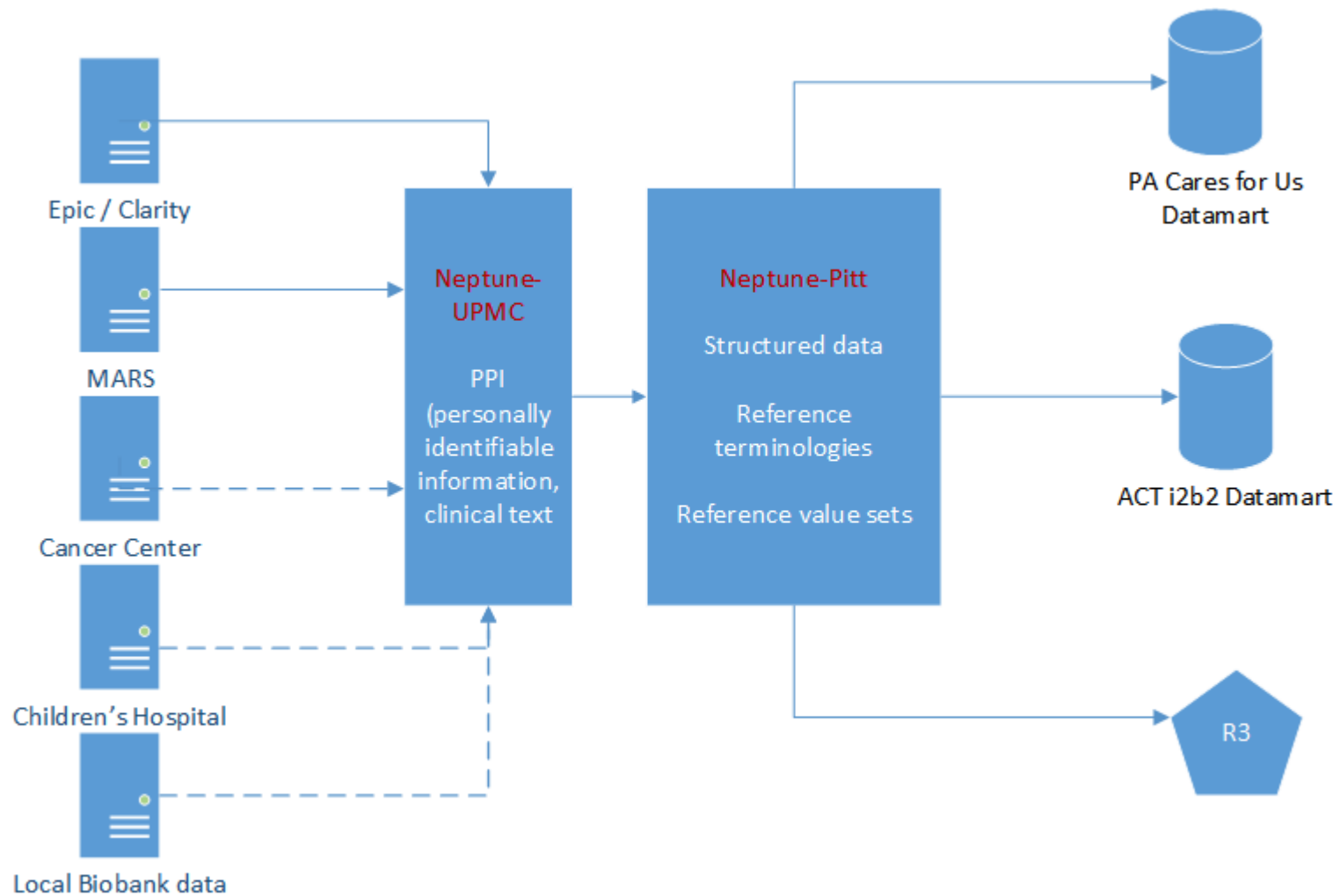
The
Precision
Medicine
Initiative®





Neptune

Research Data
Warehouse





Neptune

Research Data
Warehouse

- Architecture
 - Atomic data warehouse
 - Footprint in both UPMC and Pitt
- Data Domains
 - Personally identifiable data (PPI), demographics
 - Encounters: outpatient, ED, inpatient
 - Diagnoses: billing, encounter based, problem list
 - Procedures: billing
 - Medications: orders/prescriptions, dispensing
 - Laboratory tests: orders, results
 - Social history: tobacco, alcohol
 - Vitals, allergies
 - Clinical text
- Terminologies & Value sets
 - Demographics (race, ethnicity, gender)
 - Encounter types
 - Diagnoses: ICD-9, ICD-10
 - Procedures: ICD-9, ICD-10, CPT-4, HCPCS
 - Medications: RxNorm, NDC
 - Laboratory tests: LOINC



Neptune

Research Data
Warehouse

- Period: January 2004 - November 2017
- Update frequency: monthly
- Patients: 6.35M
- Diagnoses: 190M
- Procedures: 91M
- Laboratory test results: 973M
- Medication orders: 62M

Health Record Research Request (R3)

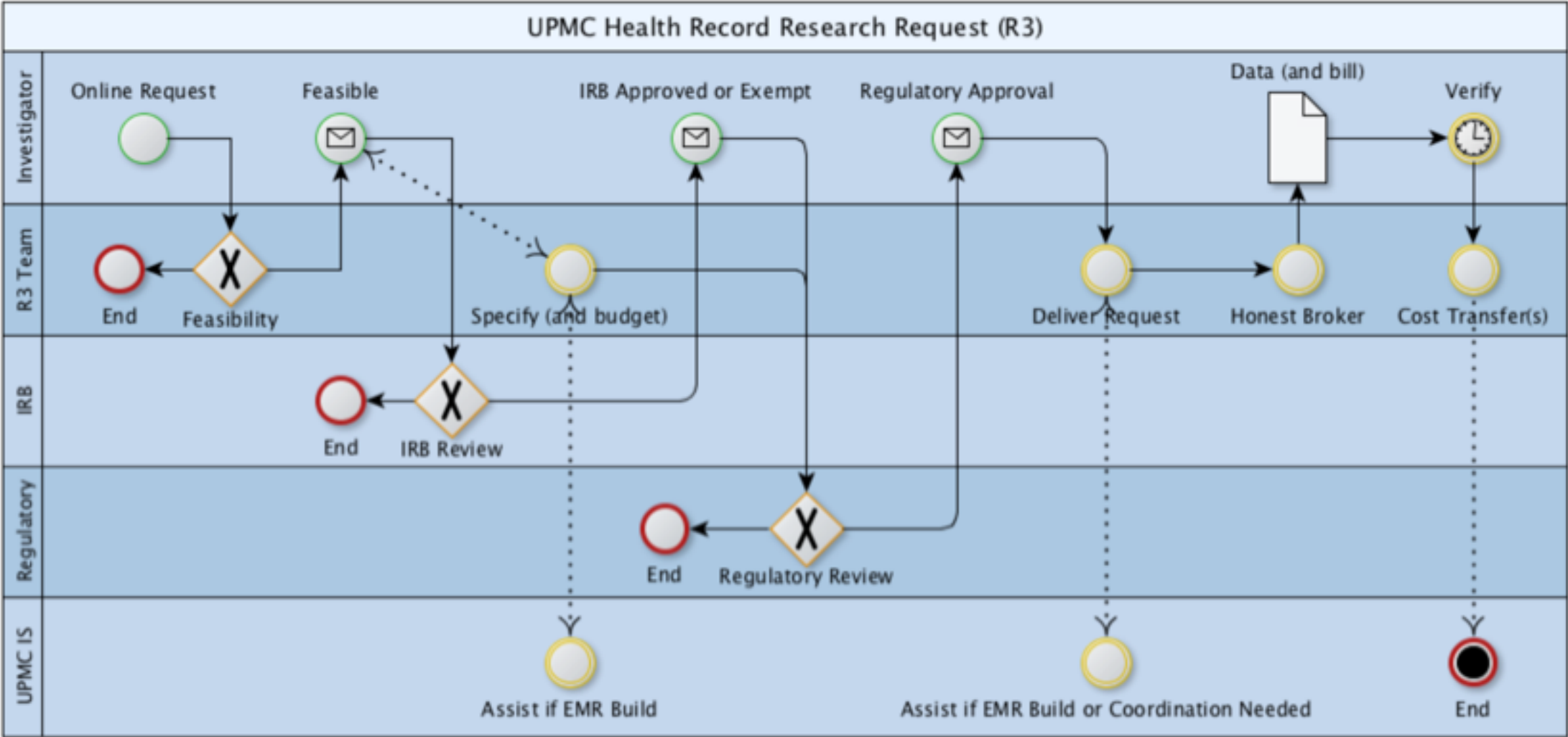
- University and UPMC desire to make certain de-identified clinical data available...for research.
- Under CRIO, on behalf of UPMC, certain DBMI staff operate via HIPAA BAA as Honest Broker
- R3 is this service or process of provisioning data through Neptune and of authorizing additional sources



University of Pittsburgh
Institutional Review Board

Honest Broker Certification

R3 Workflow





At its core, TIES is a natural language processing (NLP) pipeline and clinical document search engine. The software de-identifies, annotates, and indexes your clinical documents, making it easier for your researchers to search for and find the documents and cases. TIES also supports tissue ordering and acquisition and integration with tissue banks and honest brokers. It also works across institutions with separate TIES installations as the TCRN.



What is the TCRN?

The TIES Cancer Research Network (TCRN) is a federated tissue and data sharing network that enables researchers to identify cases, access data and request tissue/materials from one or more network partners.

Authority, responsibility, effectiveness

- National Algorithm Safety Board – Ben Shneiderman
 - Was the “right” data used to train? How does it perform? Can AI’s be responsible (who has the liability? the builder, maintainer, implementer?)
- A “Fundamental Theorem” of Biomedical Informatics (Friedman)

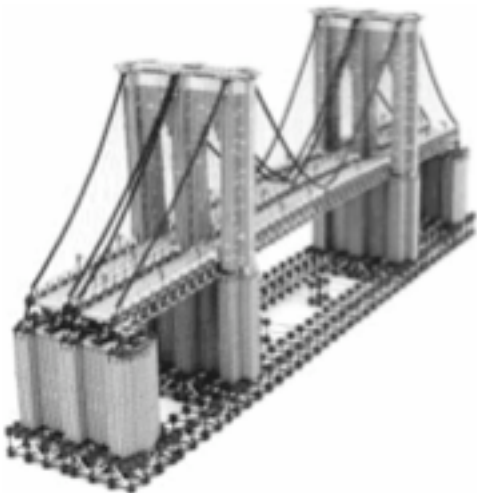


What can we expect?

- “Personalized medicine” or “Precision medicine” will play an increasing role in healthcare, particularly in cancer
- Genome sequencing will become increasingly common; integral to patient care
- Correlation of molecular changes to phenotype will be critical for both research, and also for selecting therapy for patients
- **New technology approaches are required for adaption and scale**

Important characteristics

- We must integrate systems that may not have worked together before
- These are human systems, with differing goals, incentives, capabilities
- All components are dynamic—change is the norm, not the exception
- Processes are evolving rapidly too



We are not building something simple like a bridge or an airline reservation system



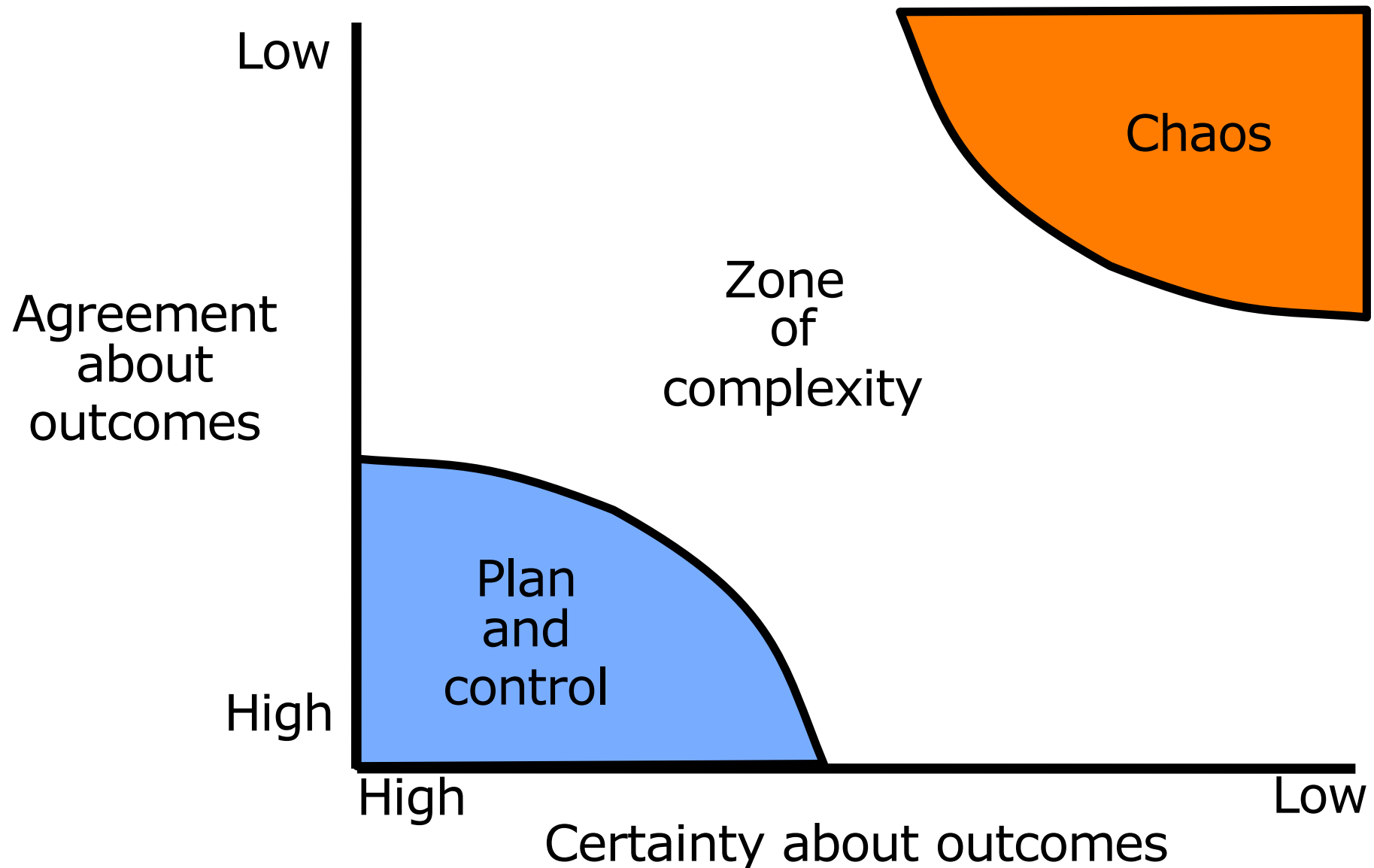
Healthcare is a **complex adaptive system**

A complex adaptive system is a collection of individual agents that have the freedom to act in ways that are not always predictable and whose actions are interconnected such that one agent's actions changes the context for other agents.

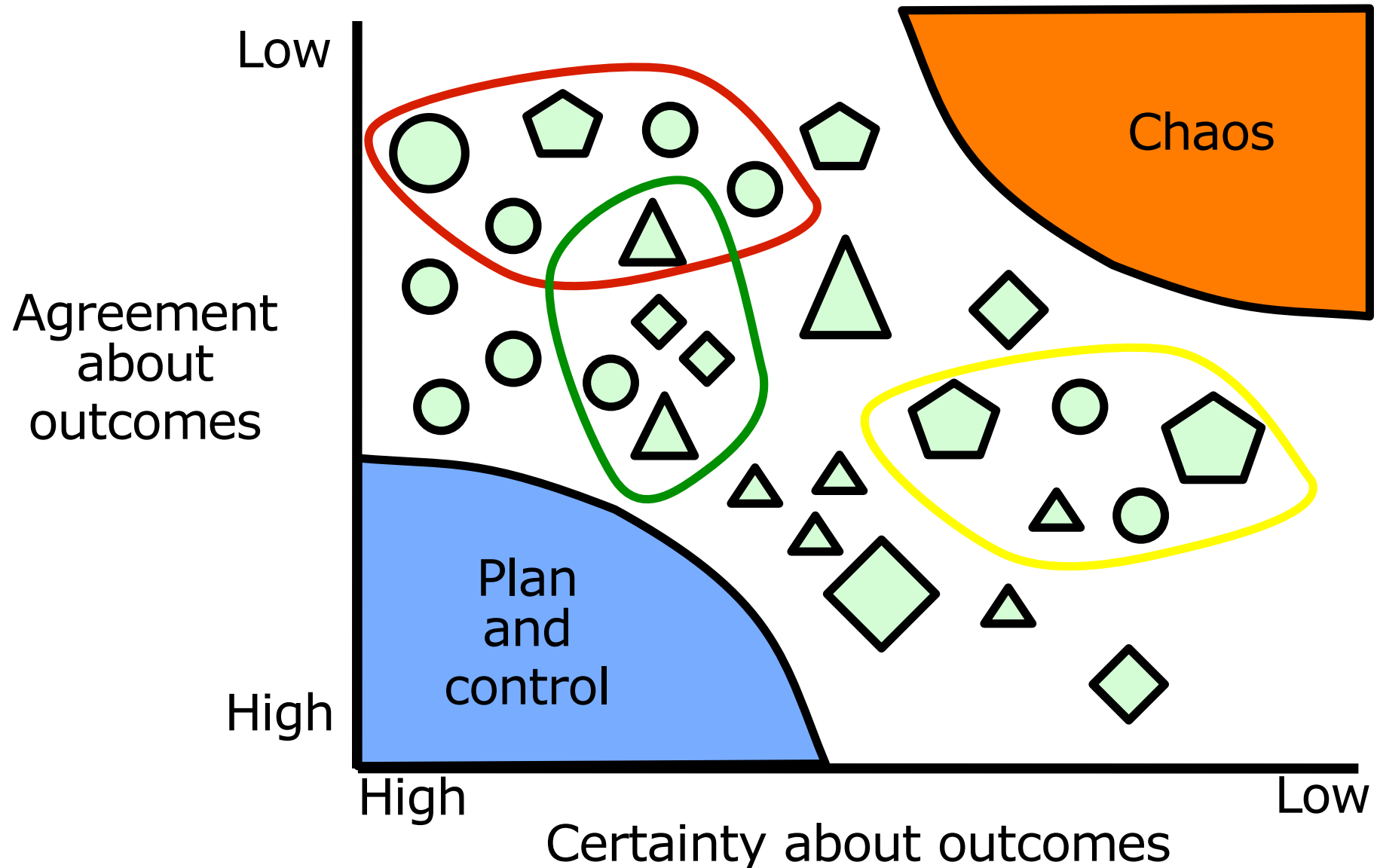
*Crossing the Quality Chasm,
IOM, 2001; pp 312-13*

- Non-linear and dynamic
- Agents are independent and intelligent
- Goals and behaviors often in conflict
- Self-organization through adaptation and learning
- No single point(s) of control
- Hierarchical decomposition has limited value

We need to function in the **zone of complexity**



We need to function in the **zone of complexity**



We call these groupings **virtual organizations (VOs)**

A set of individuals and/or institutions engaged in the controlled sharing of resources in pursuit of a common goal

Healthcare = dynamic,
overlapping VOs, linking

Patient – primary
care

Sub-specialist –
hospital

But U.S. health
system is marked by
fragmented and
inefficient VOs with
insufficient
mechanisms for
controlled sharing

Service Oriented Science

- **New information architectures enable new approaches to publishing and accessing valuable data and programs.** So-called service-oriented architectures define standard interfaces and protocols that allow developers to encapsulate information tools as services that clients can access without knowledge of, or control over, their internal workings. Thus, **tools formerly accessible only to the specialist can be made available to all**; previously manual data-processing and analysis tasks can be automated by having services access services. Such service-oriented approaches to science are already being applied successfully, in some cases at substantial scales, but **much more effort is required before these approaches are applied routinely across many disciplines.** Grid technologies can accelerate the development and adoption of service-oriented science by **enabling a separation of concerns between discipline-specific content and domain-independent software and hardware infrastructure.**

We need hosted federation of services

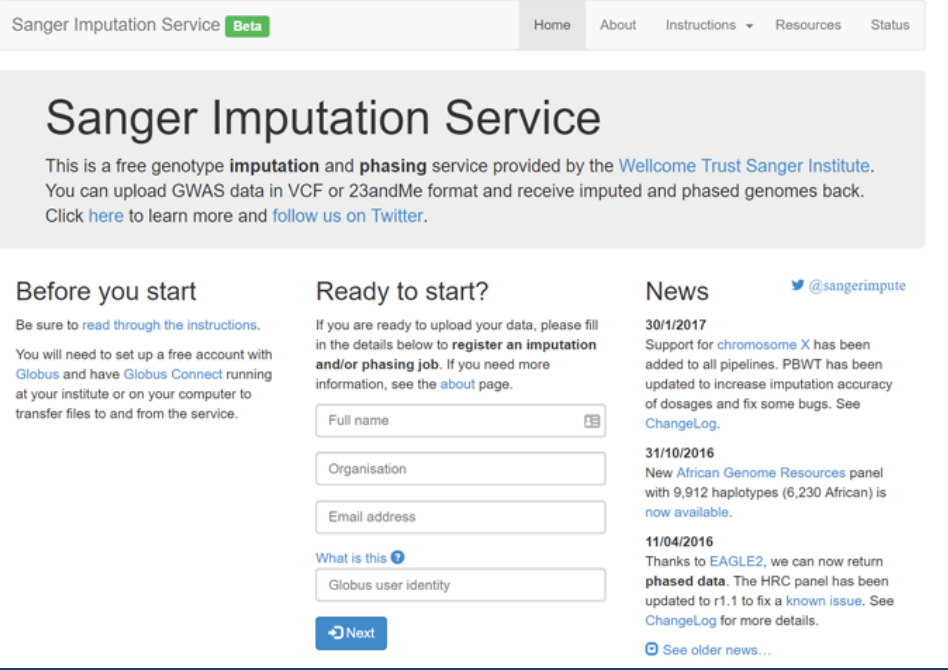
- Attribute-based authorization.
- Distributed identity management.
- End-to-end security.
- Data naming, linking, movement, and integration.
- Flexible, but enforceable policy/sociability.
- Extensibility.
- Redundancy.
- Robust in multiple industries/stability.
- Without central ownership/manageability.

Globus as solution

- Solves issue with third party access to private data
- Complement to other software/systems
- Easy to streamline and scale
- Useful for teams with distributed resources and agents
- Distribution of big data

Many variants possible

- Manage access to data at multiple locations
- Manage access to data on cloud
- Upload data for analysis
- Data download from scientific instruments
- Data publication
- Transfer data to computer for analysis



Sanger Imputation Service Beta Home About Instructions Resources Status

Sanger Imputation Service

This is a free genotype **imputation** and **phasing** service provided by the [Wellcome Trust Sanger Institute](#). You can upload GWAS data in VCF or 23andMe format and receive imputed and phased genomes back. Click [here](#) to learn more and [follow us on Twitter](#).

Before you start

Be sure to [read through the instructions](#). You will need to set up a free account with [Globus](#) and have [Globus Connect](#) running at your institute or on your computer to transfer files to and from the service.

Ready to start?

If you are ready to upload your data, please fill in the details below to **register an imputation and/or phasing job**. If you need more information, see the [about](#) page.

Full name

Organisation

Email address

What is this [?](#)

Globus user identity

[Next](#)

News

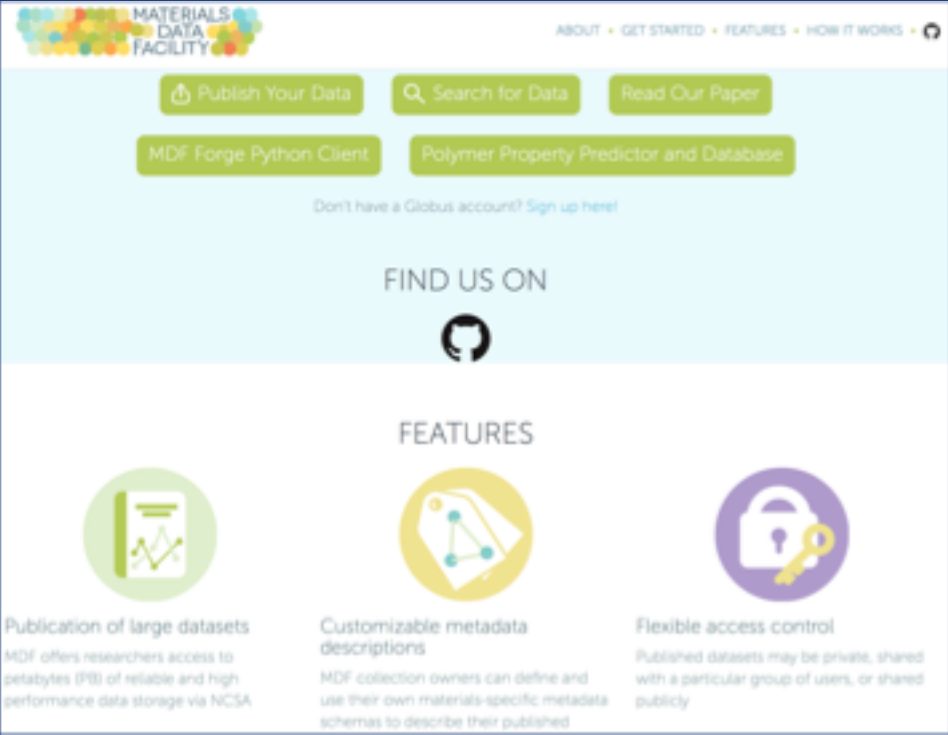
[@sangerimpute](#)

30/1/2017
Support for chromosome X has been added to all pipelines. PBWT has been updated to increase imputation accuracy of dosages and fix some bugs. See [ChangeLog](#).

31/10/2016
New African Genome Resources panel with 9,912 haplotypes (6,230 African) is [now available](#).

11/04/2016
Thanks to [EAGLE2](#), we can now return **phased data**. The HRC panel has been updated to r1.1 to fix a [known issue](#). See [ChangeLog](#) for more details.

[See older news...](#)




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FEATURES



Publication of large datasets
MDF offers researchers access to petabytes (PB) of reliable and high performance data storage via NCSA



Customizable metadata descriptions
MDF collection owners can define and use their own materials-specific metadata schemas to describe their published



Flexible access control
Published datasets may be private, shared with a particular group of users, or shared publicly



A key message: Outsource all that you can

- Outsource responsibility for determining user identities
- Outsource control over who can access different data and services within the portal
- Outsource responsibility for managing data uploads and downloads between various locations and storage systems
- Leverage standard web user interfaces for common user actions



Future: Data, Algorithms, Intelligence, oh my!

- All data is attributed (may be private, but not anonymous).
- Algorithms (machine “partners”) have responsible humans behind them.
- Intelligence is manifest as complex adaptive socio-technical learning collectives including machine “partners”.

- BIG QUESTIONS:
- In the Future: will we want to know we’re interacting with an AI (we seem to want to now, but we also want efficiency...)?
- In the Future: will we value or even tolerate anonymity (e.g. blockchain)?