



Global Alliance
for Genomics & Health
Collaborate. Innovate. Accelerate.

Peter Goodhand, Executive Director GA4GH
President Ontario Institute for Cancer Research



The Challenge



Data from **millions of samples** may be needed to achieve results and progress - showing patterns that would otherwise remain obscure.

That will take new methods and organizational models.

Historically:

- Data is typically in silos: by type, by disease, by country, by institution
- Analysis methods are non-standardized, few at scale
- Approaches to regulation, consent and data sharing limit interoperability

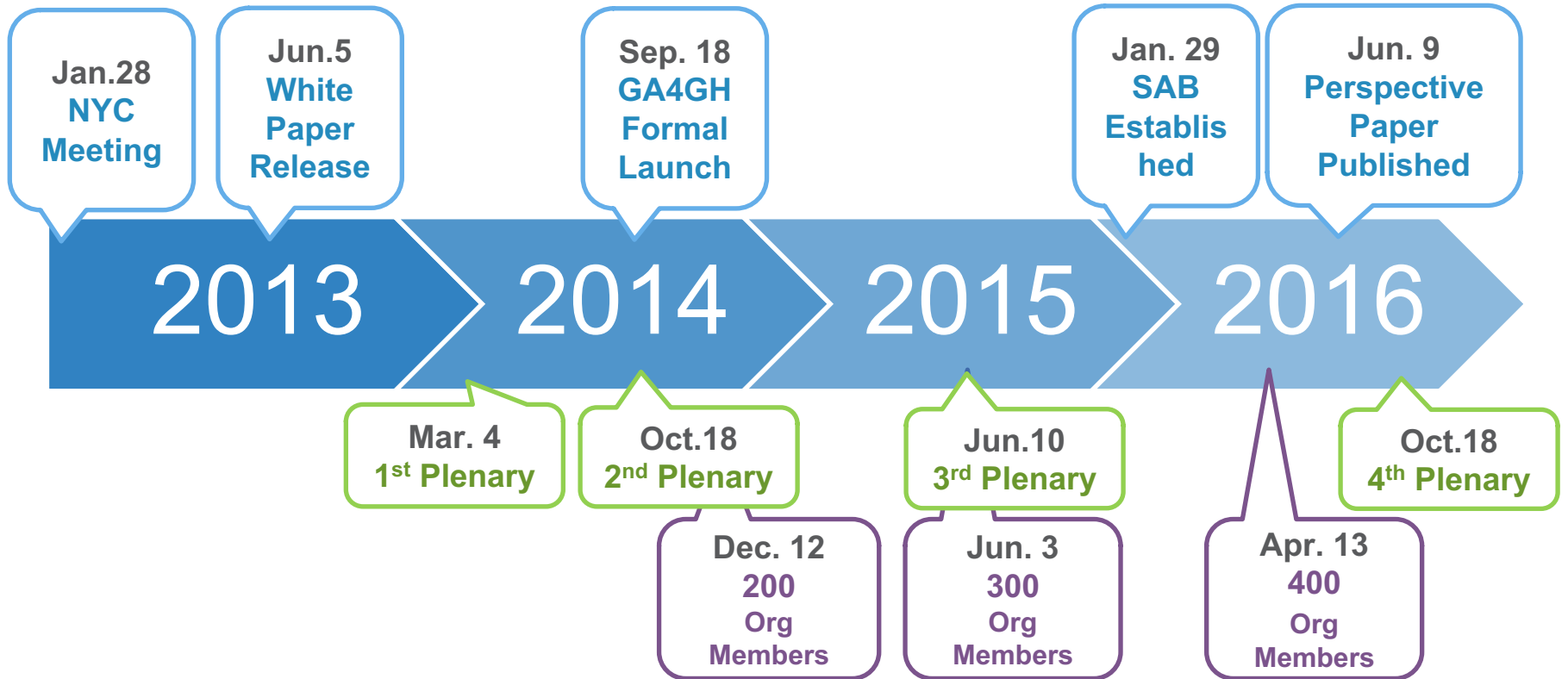
Don't act: an overwhelming mass of fragmented data, such as electronic medical records in many countries

Collective Action: achieve the interoperability of the www or global telecommunications – Smart phones.

LAUNCH A GLOBAL ALLIANCE – 2013

To accelerate progress in human health by helping to establish a common framework of harmonized approaches to enable effective and responsible sharing of genomic and clinical data, and by catalyzing data sharing projects that drive and demonstrate the value of data sharing

Organizational Milestones



470+

Organizational
Members



1000+

Individual
Members



75

Countries



Global Alliance members include

1. Life science and information technology companies (43%)
2. Universities and research institutes (32%)
3. Academic medical centers and health systems (10%)
4. Disease advocacy organizations and patient groups (5%)
5. Consortia and professional societies (5%)
6. Funders and agencies (5%)

Last Update: January 2017

Next Steps



- 5-year Strategic Roadmap
 - GA4GH vision to enable genomic data sharing
 - Cross-cutting 'product lines' to implement goals
 - Development of GA4GH Toolkit
- National Initiatives
 - Engaging national precision medicine and genomics initiatives
 - Preliminary meeting: 8 countries (Oct, 2016)
 - Next meeting (co-hosted by Genomics England and AGHA): 20+ countries (May, 2017)
- 5th Plenary
 - Open meetings to engage community, advance projects
 - Oct 15-17, 2017 (Orlando, Florida)

Coordination with G2MC



- G2MC: *implementation* of genomic medicine into clinical care
- GA4GH: enabling international genomic and health data sharing within and across the research- care continuum
- GA4GH-G2MC collaboration on health informatics
- G2MC secretariat co-localized with GA4GH at



Research & Clinic - Historically Two Solitudes



Genomic Research	Genetic Health Care
<ul style="list-style-type: none">• Subjects, papers, global	<ul style="list-style-type: none">• Patients, providers, local (national or state/system)
<ul style="list-style-type: none">• Rapid change of scale and scope of NGS due to decreased costs and increased compute power	<ul style="list-style-type: none">• Change from single gene to panels
<ul style="list-style-type: none">• Data science primarily academic developed/used	<ul style="list-style-type: none">• Data science developed and delivered by HIT providers
<ul style="list-style-type: none">• BAM/SAM/CRAM, VCF, Aligners, variant callers, pipelines	<ul style="list-style-type: none">• EHR, EMR, HL7, FHIR, SMART

Global Research, Local Healthcare



Human Genome Project (1990)



International HapMap Project (2002)



International Cancer Genome Consortium (2008)



1000 Genomes Project (2008)

Research



Academic Activities

- Journals
- Conferences

Healthcare

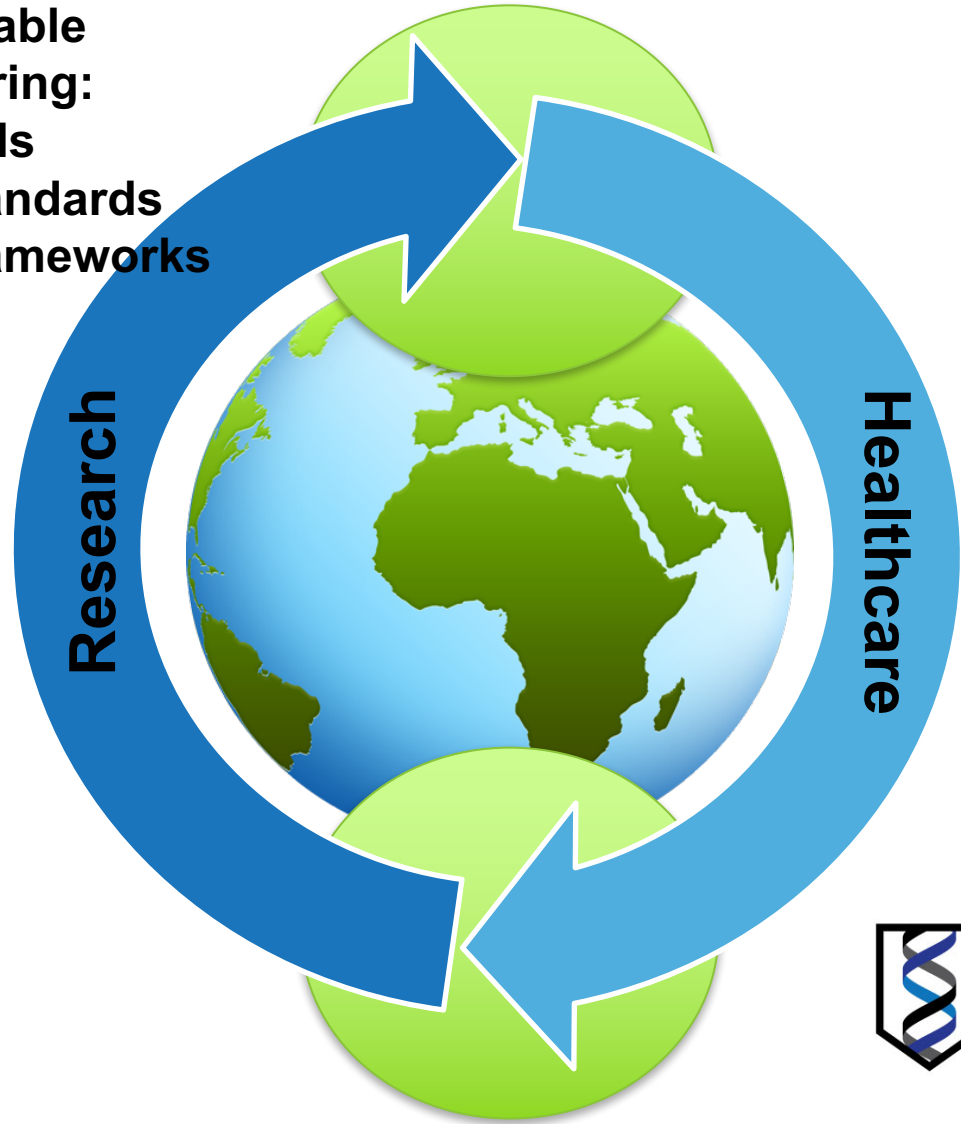
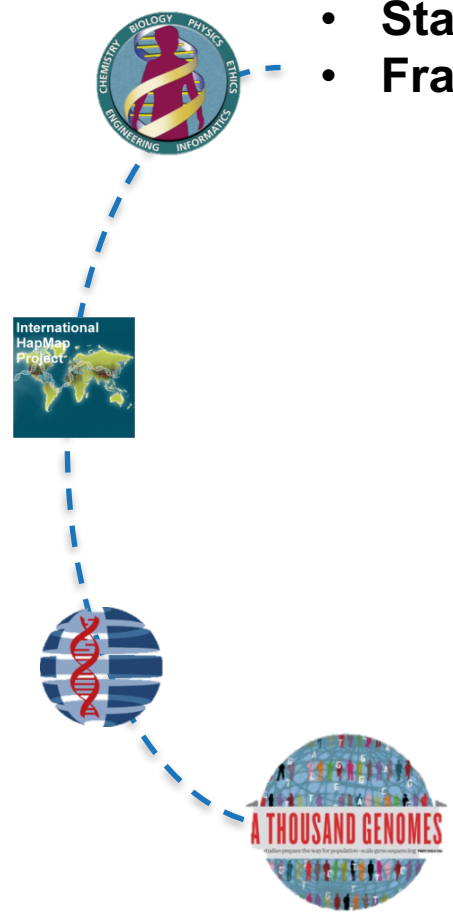
Jurisdictional Challenges:

- Health expenditures
- Legal systems
- Languages
- Infrastructure

“Global Knowledge, Local Care”

Interoperable Data Sharing:

- APIs
- Standards
- Frameworks



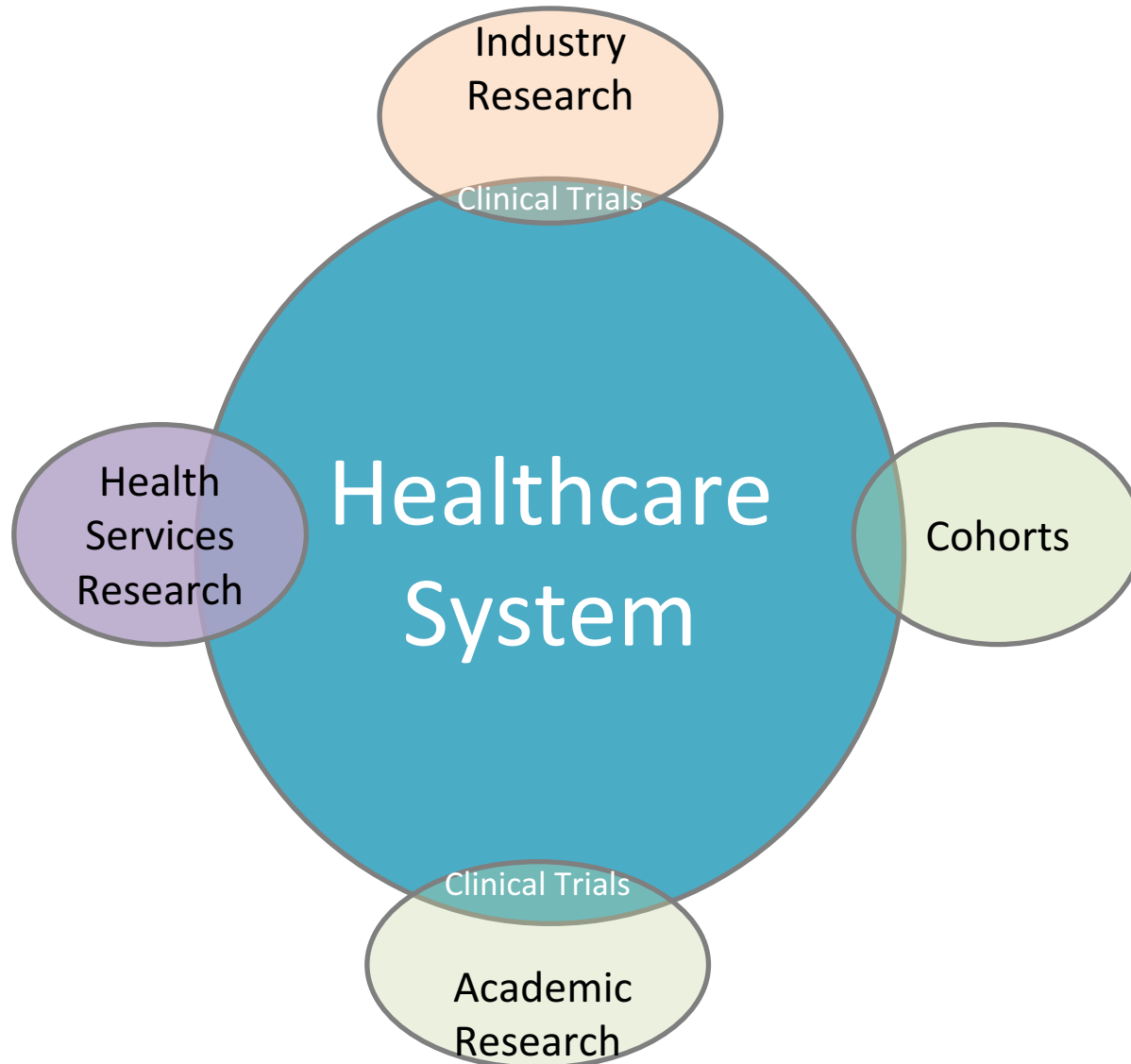
Beacon

BRCA
CHALLENGE

Genomic
Knowledge
Exchanges

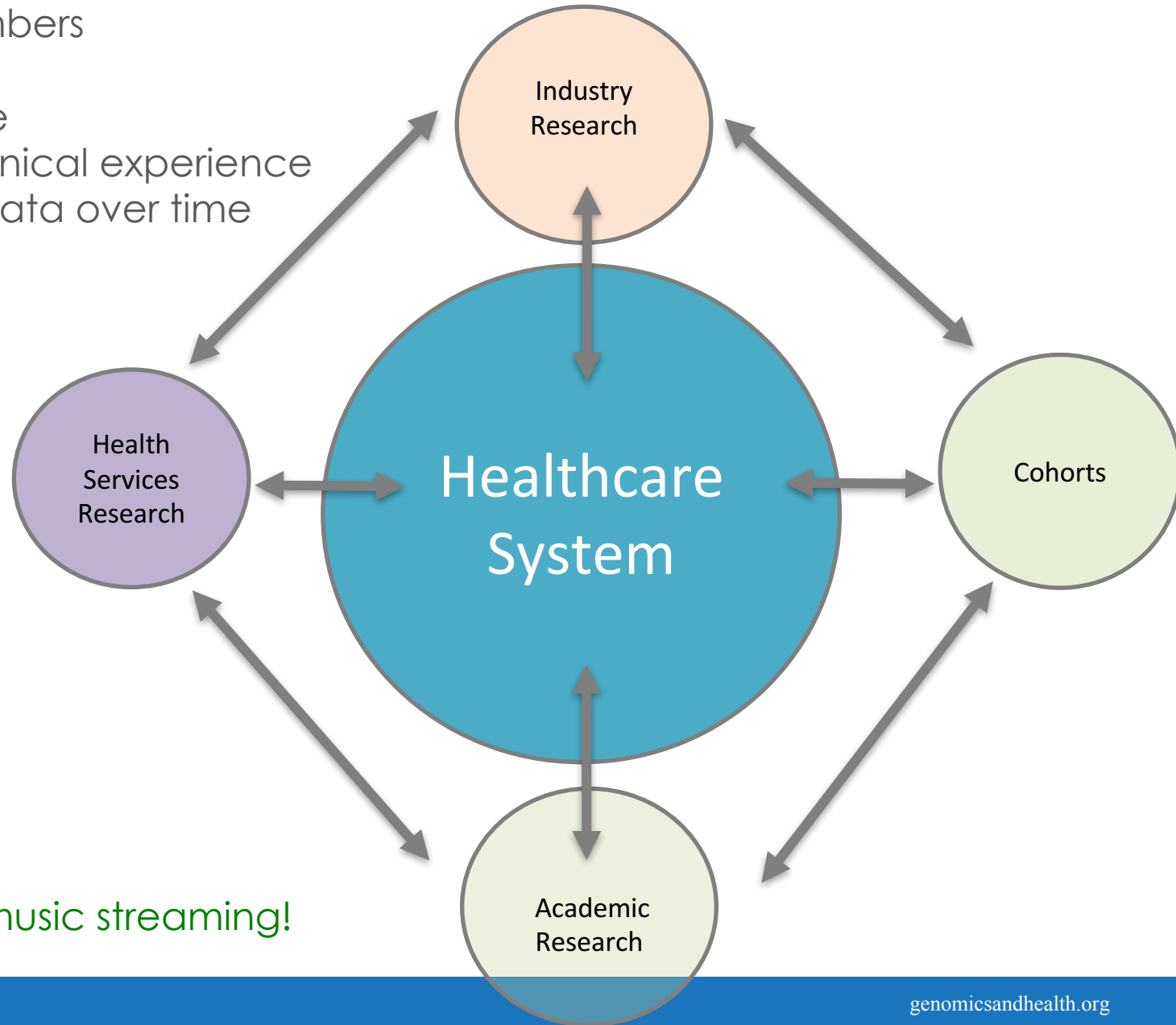
Matchmaker
Exchange

Cancer
Gene
Trust



Tremendous opportunity in digitized data

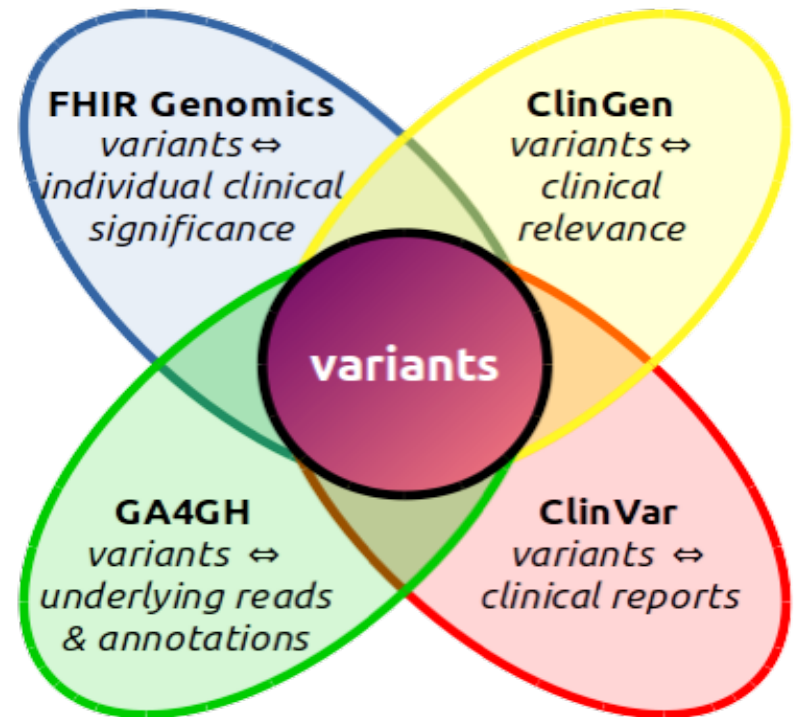
- Power in numbers
- Dynamic
- Interoperable
- Linkage of clinical experience & research data over time



CD's vs digital music streaming!

Variant Modelling Collaboration

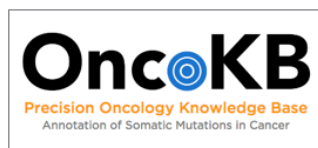
- Gil Alterovitz, Harvard Medical School, FHIR Genomics
- Larry Babb, Sunquest, ClinGen
- Karen Eilbeck, University of Utah
- Gaston Fiore, Boston Children's Hospital, FHIR Genomics
- Bob Freimuth, Mayo Clinic, ClinGen, HL7/FHIR
- Reece Hart, Invitae, GA4GH
- David Kreda, Harvard Medical School, FHIR Genomics
- Jennifer Lee, NCBI, ClinVar
- Peter Robinson, Jackson Labs



Courtesy of Reece Hart

Variant Interpretation for Cancer Consortium (VICC)

Building on the work of [Genotype-to-Phenotype \(G2P\)](#) Task Team led by Adam Margolin



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Variant Interpretation for Cancer

- Gene
- Variant
- Cancer subtype
- Clinical implication: drug sensitivity, drug resistance, adverse response, diagnostic, or prognostic
- Source (e.g., PubMed identifier)
- Curation group

<http://ga4gh.org/#/vic>

ga4gh-dwg-vic@genomicsandhealth.org

VICC Co-chairs:

- Obi Griffith
- Nuria Lopez-Bigas
- David Tamborero
- Malachi Griffith

Goals/Principles:

- Clinical cancer variant interpretation
- Standards and guidelines
- Open content
- Interoperability



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genomicsandhealth.org

- eHealth task team (Chair: John Mattison)
- Objectives:
 - Support the discovery, access, and sharing of genomic and clinical information within and among EHR systems
 - Survey current activities in eHealth and promote best practices
 - Harmonize efforts where possible
- Deliverables to date:
 - Catalogue of 85 global resources for sharing clinical and genomic eHealth data
 - Catalogue of 23 family health history tools
 - Family history collection tools: statement of best practice
- Possible focus areas for 2017-2018:
 - Resources to support users in assessing/evaluating products for clinical care
 - Collaborations HL7-FHIR, SMART on FHIR, Sync for Genes

Coordinating with external e- health initiatives (Lead: Grant Wood)

- HL7 Clinical Genomics workgroup (FHIR)
- SNOMED
- DIGITizE, IGNITE, EMERGE
- ClinGen, ClinVar
- Farr Institute, Swiss Institute of Bioinformatics
- Healthcare Services Platform Consortium
- Genetic Alliance (S) and other patient groups
- National Precision Medicine Initiatives
- HUGO, HVP/GV, HGVS

Aligning Data in Research and Health

- Variant Modelling
- Variant Interpretation, Clinical Actionability
- Phenotype Ontologies
- Genotype, Phenotype, “Envirotype”
- Pathogenicity Classifications – 5 vs 2
- Rare/Undiagnosed disease: discovery and matching underway (MME)– P&F facebook, facial recognition
- Cancer: pre-disposition in germline; heterogeneity, clonal evolution, primary, mets, recurrence in somatic

Genomic data in Healthcare Systems

- In the point of care EMR
- In the integrated/federated EHR
- Data Repositories for Genomic Data - GACS
- Direct and Immediate CDS
- Warnings, Alerts, Updates – New Knowledge
- System responsibilities for security and privacy of genomic data
- Patient/family role in use of their data for research

Many Questions

Research Perspective

How can we gain access to well annotated clinically generated genomes?

How can we interrogate and learn from longitudinal data in the health record - with genomic data?

How can we connect with patients (families) to engage them directly in the research process?

How can we leverage the investment of industry in Clinical Trials?

Clinical Perspective

How can we share clinically generated data in a way that will benefit the care we deliver?

How will complex and at times evolving genomic knowledge be made accessible to my practice?

What should I advise my patients regarding participation in research, incidental findings, return of results?

How should we regard traditional RCTs vs RWD?

What are the issues in your country (state/province/system/region) to accomplish the goals of a longitudinal genomic patient records?