

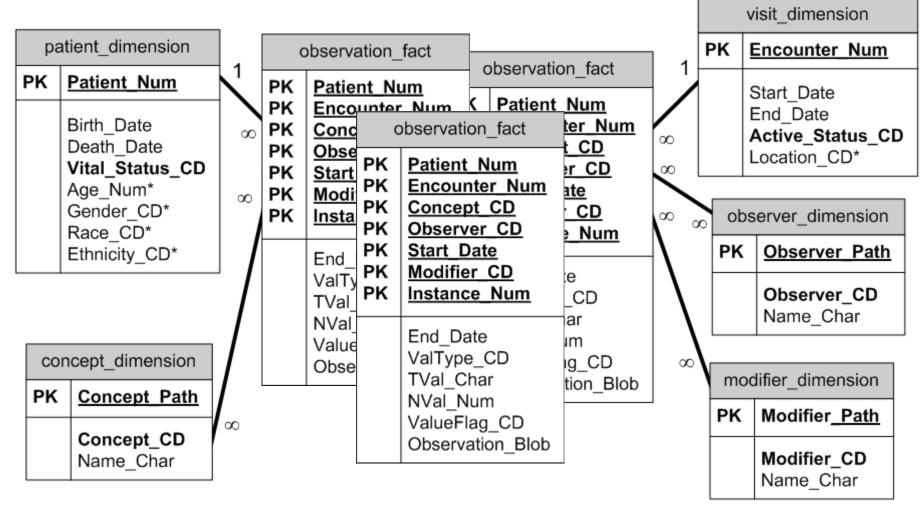
Advanced i2b2 Topics

Shawn Murphy MD, Ph.D.

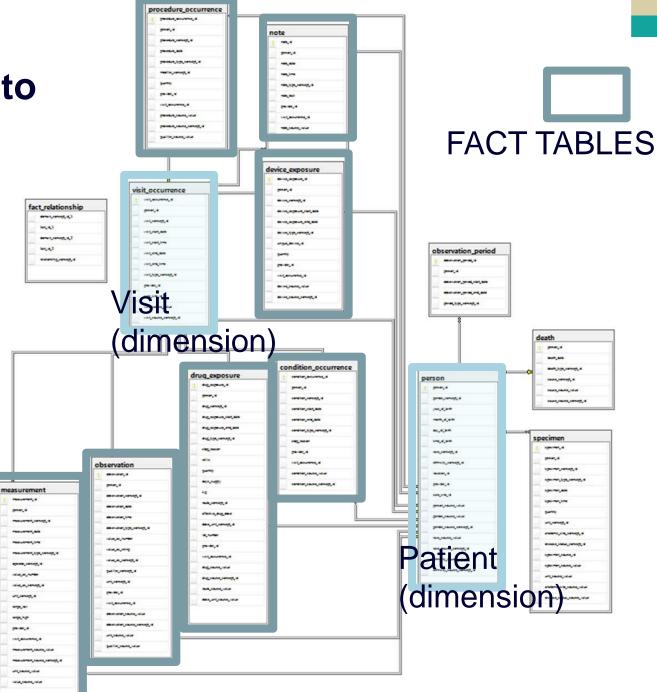
Mass General Brigham and Harvard Medical School

November 14th, 2020

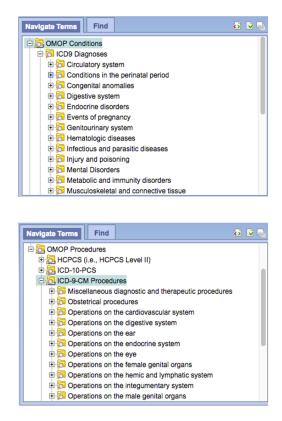
i2b2 Star Schema can adapt to Multiple Fact Tables

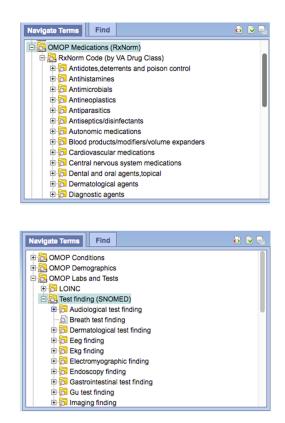


Adapts to OMOP



Connect to OMOP by building ontology of OMOP standard concepts

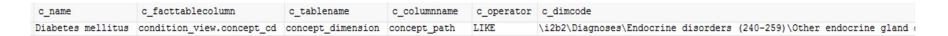


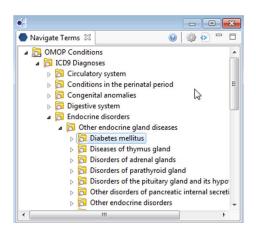


Ontologies covering the condition, procedures, drug, measurement and observation domains. All terms are mapped to standard concepts using OMOP's mapping tables

Use Ontology Tables to direct Queries to proper Fact Table view

Prepend c_facttablecolumn with OMOP domain view and modify CRC to parse into 'domain view' and 'c facttablecolumn'

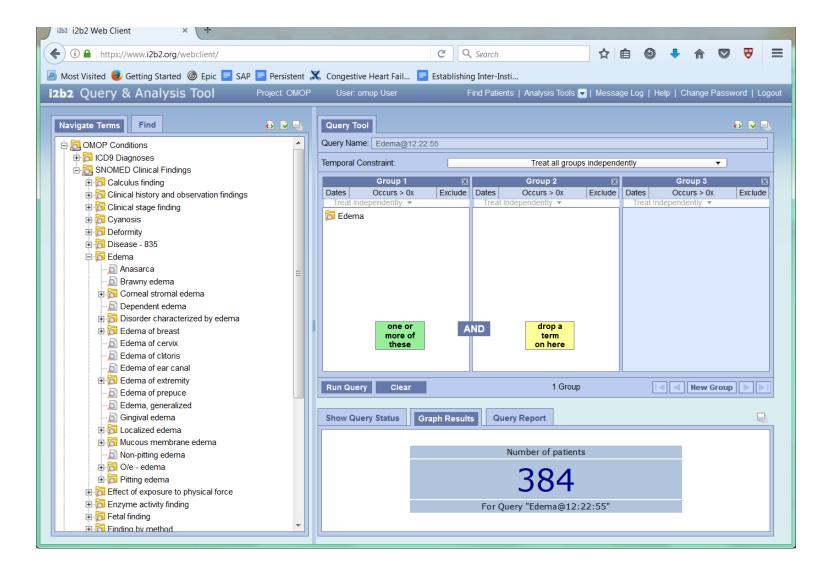




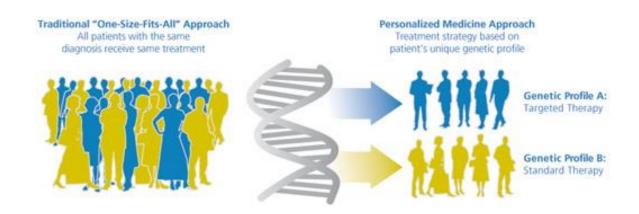
select patient_num from condition_view where concept_cd IN (select concept_cd from concept_dimension where concept_path like '\i2b2\Diagnoses\Endocrine disorders (240-259)\Other endocrine gland diseases (250-259)\(250) Diabetes mellitus\%')

https://community.i2b2.org/wiki/display/OMOP

OMOP Linked into i2b2 software

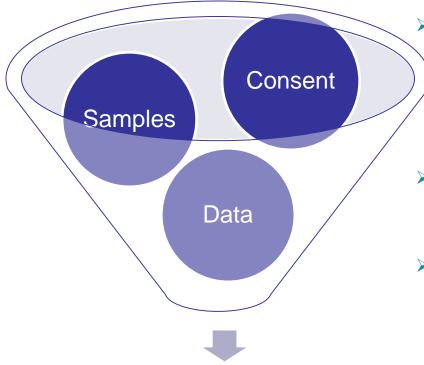


Personalized Medicine and Genomic technology are critical to managing populations



- Managing a population involves improving health outcomes of the group as a whole by identifying, monitoring and addressing health needs of individuals through:
 - Subpopulation stratification
 - Targeted, evidence-based treatment protocols
 - Predictive analytics

The Partners Biobank



Research Discoveries

The Partners Biobank provides samples (plasma, serum, and DNA) collected from consented patients.

 84,000 patients have consented to date

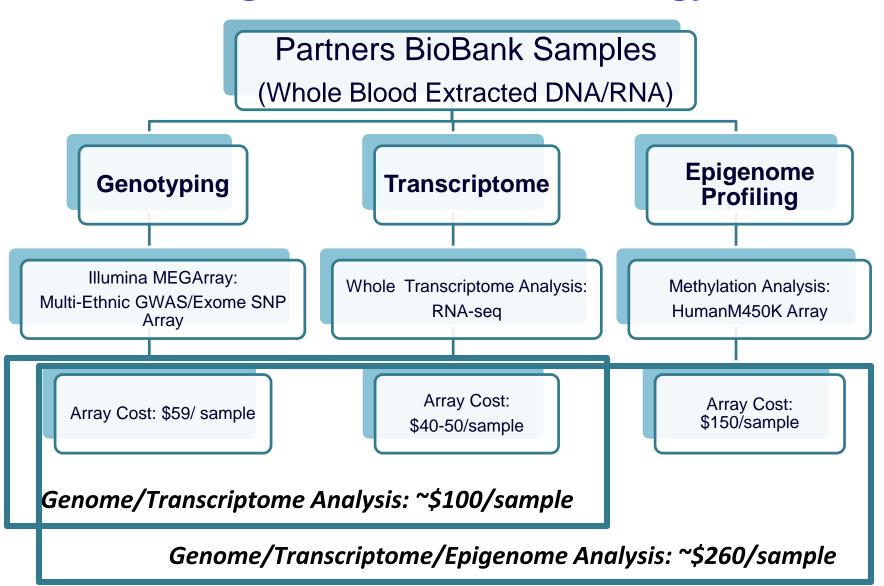
Samples are available for distribution to Partners investigators* to help identify novel Personalized Medicine opportunities that reduce cost and provide better care



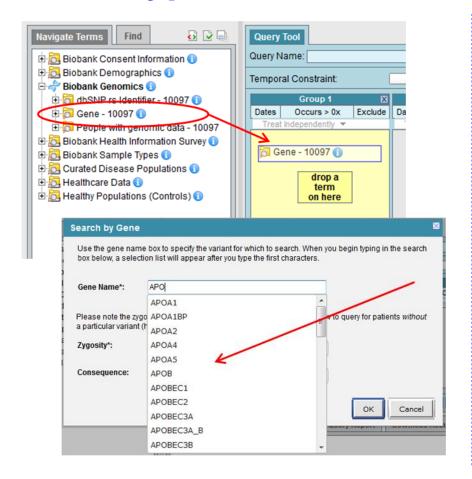
*with required approval from the Partners Institutional Review Board (IRB).

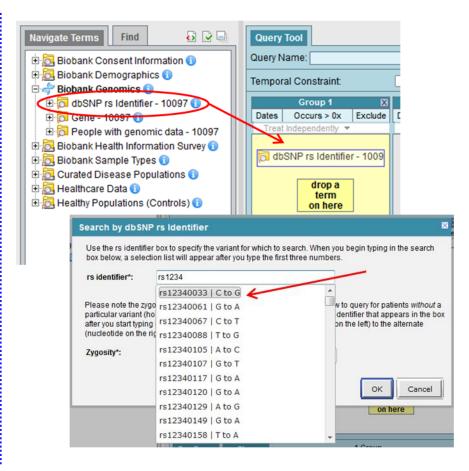
Improved Clinical Care for All Patients

Biobank Integrative Genomics Strategy

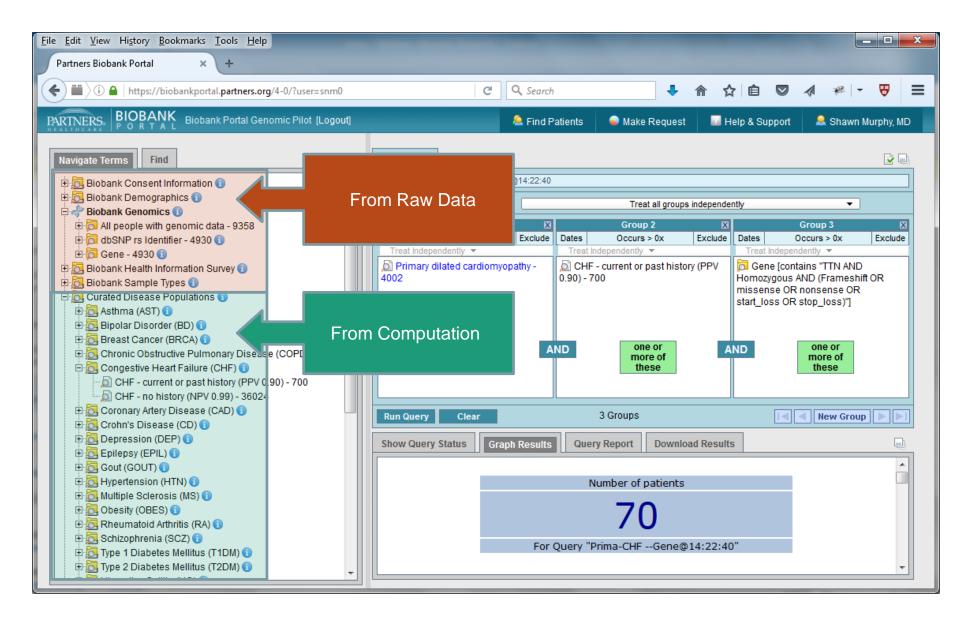


Genotype Data

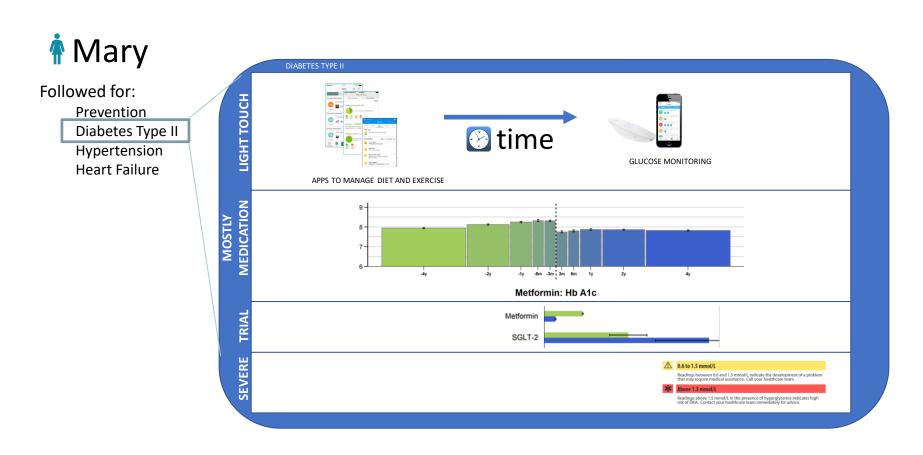




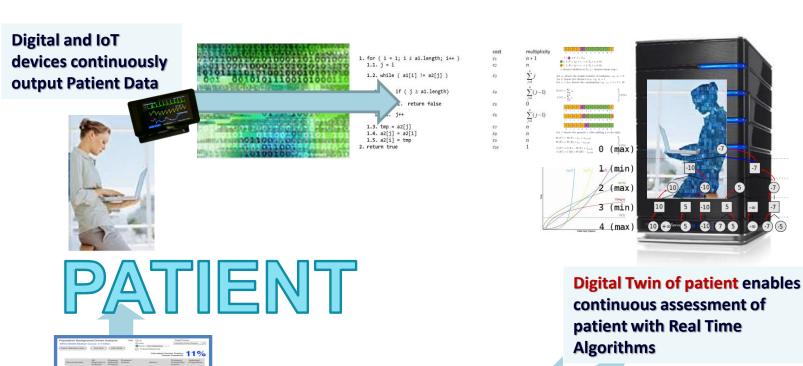
https://community.i2b2.org/wiki/display/IGD/Loading+Genomic+VCF+Files+into+i2b2



In Digitally Driven Healthcare, Disease Labels determine Algorithms for Managing Patient

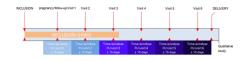


Innovations Enabling Digital Care



Navigator Model dramatically increases Frequency and Convenience for Patient Communication

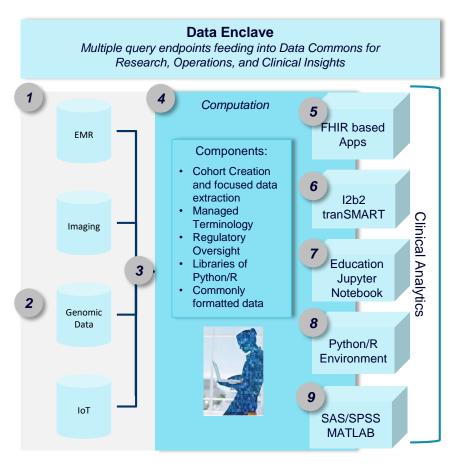






System drives Pragmatic Clinical Trials Leading to Continuous Process Improvement

Digital Twin for Continuously Assessing Patient



Data Enclave	
1	Enable Data Extracts, perhaps some are Federated
2	Combine and Link data, put in common OMOP/i2b2 format
3	Conduit to Data Enclave
Technical Solution Development	
4	Healthcare Ready Bundle
Research and Clinical Application Projects	
- 5 -	FHIR based SMART Apps
6	I2b2 tranSMART with Fractalis plugin (next version of SmartR plugin)
7	Jupyter Notebook with AI Visualizations – code can advance to production
- 0 -	Python/R Environment full interactive development in Data Lake

Many, many contributions ...

- Management
 - ■Isaac Kohane
 - Susanne Churchill
 - Griffin Weber
 - Lee Nadler
 - DouglasMacFadden
 - ■Christopher Herrick
 - Victor Castro
 - ■Roy Perlis
 - ■Beth Karlson
 - Paul Avillach
 - ■Jeff Klann
 - Michael Mendis

- Biobank/Phenotyping
 - Scott Weiss
 - Beth Karlson
 - Jordon Smoller
 - Natalie Bouton
 - Vivian Gainer
 - Nich Wattanasin
- Healthcare Innovation Team
 - Kavi Wagholikar
 - Randy Gollub
 - Calum Macrea
 - Sandy Aronson

Special thanks to...

Core i2b2 Developers over past 14 years:

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- Nich Wattanasin
- William Simons
- Christopher Herrick
- Bill Wang

- Kavi Wagholikar
- Wayne Chan
- Bhaswati Ghosh
- Reeta Metta
- Barbara Benoit
- David Wang
- Jeff Klann

