A National Center for Biomedical Computing





Developing Specialized Hives to Contribute to the i2b2 Ecosystem

Shawn Murphy Michael Mendis

Distributed Query System links together Specialized i2b2 Hives





Patient Databases of Coded Data

Patient Databases of Highly Specialized Data

Patient Databases of Unstructured Text Data Specialized Patient Databases Created by Extracting Features from Big Data

Analytic Results of Patient-specific Calculations

Patient Databases with fluid data types from spreadsheets

Greater Specialization





I2b2/RPDR Text-based Queries

- 130 Million Notes Available alongside 2.4 Billion Facts to query in i2b2/RPDR
- Researchers search for phrases of interest in a variety of clinical notes
 - Cardiology, Discharge Summaries, Endoscopy, LMR, Operation, Pathology, Pulmonary, and Radiology
- Transparent to users add text based searching to query tool
- Notes are updated nightly in special notes repository
- Several steps taken to conceal PHI through query
 - Researchers will not be able to query on numbers of any kind
 - Names are excluded from text index making them "unsearchable"
 - Search terms are also matched against a PHI database
 - □ Separate database is securely maintained with latest patient PHI, including names, addresses, numbers, nicknames, etc.
 - Search terms are queried in Notes Repository and PHI database
 - Patients that are returned from both databases are excluded from results
 - Final aggregate counts are obfuscated to hide true totals and limit ability to identify one individual

Policy can be enforced at Query Tool when item is used in query

Search for Diabetes, A1C, and SGLT2 Inhibitors



Enhance w/ LMR Note Search

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RPDR – Additional Patients Returned





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Query 1.68 billion rows of Genomic Data for Specific Variants





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Put together a new Hive very quickly with BRISSKIT

BRISSKIT is an open source project that was developed in Leicester, UK

- Allows Spreadsheet to be uploaded into i2b2 from a web client
- Creates an instant i2b2 repository of spreadsheet data



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Imaging – DICOM Index allows images of high quality to be distinguished in a PACS database

- Investigators will be able to define sets of patients who are relevant to their research by defining the specific type of image required for their analysis (e.g. high resolution).
- Through the Big Data Commons, Investigators will be able to link this patient cohort to other available data (genomic data, biobank samples, other research data, EHR data, etc)





Unpredictable Quality Using Raw ICD9/10 Codes

Phenotype	Count with ICD-9/ICD-	Count (90% positive	Count with Genotype
	10 Code	predictive value)	Data
Asthma	7618	3322	805
Bipolar Disorder	1754	219	84
Breast Cancer	2101	1711	378
Congestive Heart Failure	10160	4597	1859
Coronary Artery Disease	1435	803	236
Crohn's Disease	5177	700	350
Depression	11154	4273	1074
Epilepsy	2351	1211	381
Gout	2464	1828	566
Hypertension	20788	16995	4553
Multiple Sclerosis	602	320	58
Obesity	10245	12179	3191
Rheumatoid Arthritis	3475	878	261
Schizophrenia	509	83	14
Type 1 Diabetes	2196	232	61
Type 2 Diabetes	7123	4385	1268
Ulcerative Colitis	1359	624	157

May 4, 2016, n ~ 40,000

Predictably Quality - Computed Phenotypes

Phenotype	Count with ICD-9/ICD-	Count (90% positive	Count with Genotype
	10 Code	predictive value)	Data
Asthma	7618	3322	805
Bipolar Disorder	1754	219	84
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May 4, 2016, n ~ 40,000

Definitive Disease States Associated with Genomic Data

Phenotype	Count with ICD-9/ICD-	Count (90% positive	Count with Genotype
	10 Code	predictive value)	Data
Asthma	7618	3322	805
Bipolar Disorder	1754	219	84
Breast Cancer	2101	1711	378
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May 4, 2016, n ~ 40,000

High Quality Data Available for Genomics Queries





Patient Databases of Coded Data

Patient Databases of Highly Specialized Data

Patient Databases of Unstructured Text Data Specialized Patient Databases Created by Extracting Features from Big Data

Analytic Results of Patient-specific Calculations

Patient Databases with fluid data types from spreadsheets

Greater Specialization

The Distributed Patient Information Commons is Managed by the Ontologies

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Getting Data from the Patient Information Commons

Using spreadsheets to extract matrixes of data

Using data transfer from one node to another to run the analysis on one node

Mega-analysis by Julia with direct access to data

Using SMART to get insights into clinical care

Partners Biobank Portal – Specify Data for Download

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Tribute to...

- RPDR/I2b2 Core Team
 - Christopher Herrick
 - Michael Mendis
 - Lori Phillips
 - Janice Donahoe
 - Nich Wattanasin
 - Wayne Chan
 - Vivian Gainer
 - Alyssa Goodson
 - Mariah Mitchell
 - Martin Rees
 - Charles Wang
 - Laurie Bogosian
 - Stacey Duey
 - Andrew Cagan
 - David Wang

- Biobank Team
 - Natalie Boutin
 - Victor Castro
 - Scott Weiss
 - Beth Karlson
- SMART Team
 - Ken Mandl
 - Josh Mandel
 - Kavi Wagholikar
- Genomics Innovation Team
 - Sandy Aronson
 - Heidi Rehm
 - Calum MacRea

... and many more.



Centre for Health Technologies University of Pavia

4th i2b2 EU AUG Meeting Sept 15th 14:30 - Sept 16th 12:30 Pavia – ITALY Invited Talks by Paul Avillach Shawn Murphy

FILMEFI

EX

Also featuring Ulli Prokosch Ulrich Sax Christel Daniel, ... further info http://cht.unipv.it

Sponsored by:



FOUNDED BY BRIGHAM AND WOMEN'S HOSPITAL AND MASSACHUSETTS GENERAL HOSPITAL

I2b2 and SMART Information and Software on the Web

i2b2 Homepage (<u>https://www.i2b2.org</u>) i2b2 Software (<u>https://www.i2b2.org/software</u>) i2b2 Community Site (<u>https://community.i2b2.org</u>) SMART Platforms Homepage (<u>http://smarthealtit.org</u>)

