

Capturing Well-structured Patient Data in 5 minutes (not including questions)

Michael Brudno

Scientific Director, Centre for
Computational Medicine
Hospital for Sick Children

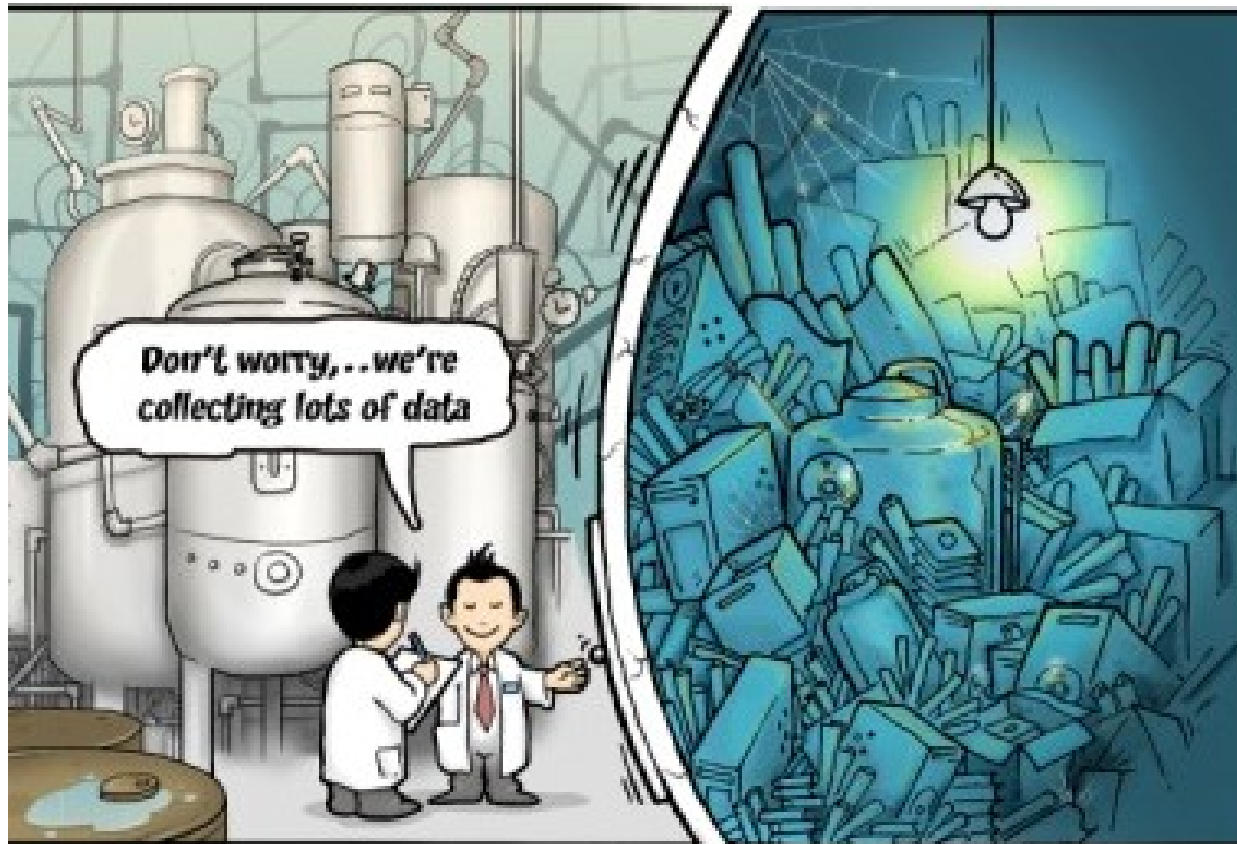
Associate Professor



SickKids[®]

Patient Data

- Has to come from somewhere



Previous State of Clinical Phenotyping

- Important to collect phenotype data to cross-reference against cases observed across sites
- Currently Two Alternatives: free text or checkboxes

Dysmorphic features

- df
- dysmorphic
- dysmorphic faces
- dysmorphic features

Congenital malformation/anomaly:

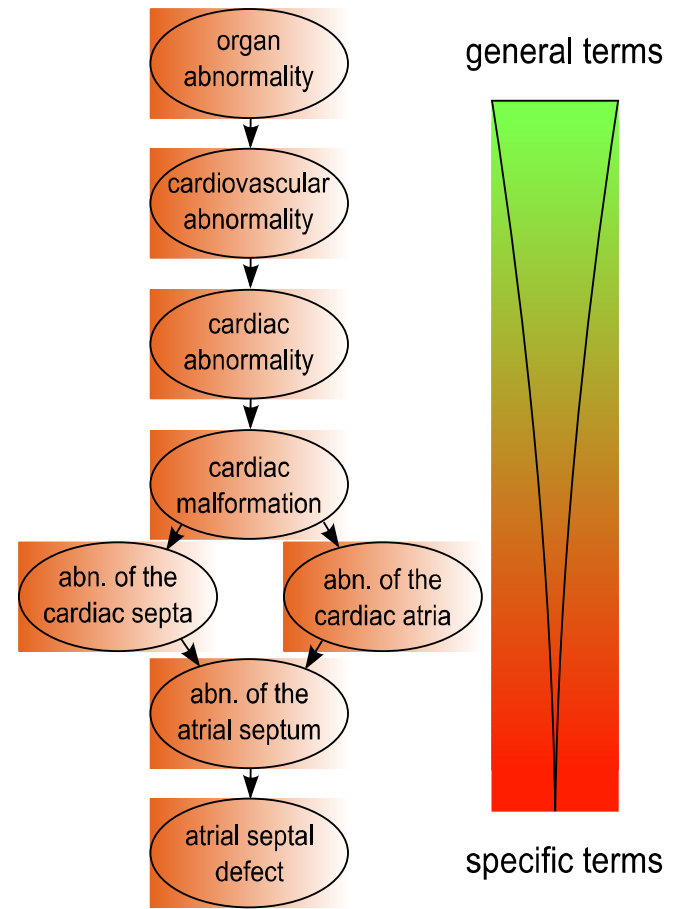
- congenital anomaly
- congenital malformation
- congenital anomoly
- congenital anomly
- congenital anomaly
- congenital anomaly
- cong. m.
- cong. Mal
- cong. malfor
- congenital malform
- congenital m.
- multiple congenital anomalies
- multiple congenital abnormalities
- multiple congenital abnormalities

Phenotypic description (Clinical symptoms)

<p>Behavior, Cognition and Development</p> <p><input type="checkbox"/> Global development delay</p> <p><input type="checkbox"/> Fine motor delay <input type="checkbox"/> Gross motor delay</p> <p><input type="checkbox"/> Language delay</p> <p><input type="checkbox"/> Learning disability</p> <p><input type="checkbox"/> Mental retardation</p> <p>Examples of lists:</p> <p><input type="checkbox"/> Mild</p> <p><input type="checkbox"/> Moderate</p> <p>* <input type="checkbox"/> Cong. malfor. behav. pro.</p> <p>* <input type="checkbox"/> dd, mental retardation</p> <p>* <input type="checkbox"/> Attention deficit hyperactivity disorder</p> <p>* <input type="checkbox"/> Autism</p> <p>* <input type="checkbox"/> df < delayed puberty</p> <p>* <input type="checkbox"/> Pervasive developmental delay</p> <p>* <input type="checkbox"/> Psychiatric disorders (Specify below)</p> <p>* <input type="checkbox"/> dd df mr</p> <p>* <input type="checkbox"/> Other: mental retard. short stature</p> <p>Neurological</p> <p><input type="checkbox"/> Hypotonia</p> <p><input type="checkbox"/> Seizures</p> <p><input type="checkbox"/> Ataxia</p> <p><input type="checkbox"/> Dystonia</p> <p><input type="checkbox"/> Chorea</p>	<p>Cardiac</p> <p><input type="checkbox"/> ASD</p> <p><input type="checkbox"/> VSD</p> <p><input type="checkbox"/> AV canal defect</p> <p><input type="checkbox"/> Coarctation of aorta</p> <p><input type="checkbox"/> Tetralogy of fallot</p> <p><input type="checkbox"/> Other: _____</p> <p>Craniofacial</p> <p><input type="checkbox"/> Craniosynostosis</p> <p><input type="checkbox"/> Cleft lip <input type="checkbox"/> Cleft palate</p> <p><input type="checkbox"/> Microretrognathia <input type="checkbox"/> Retrognathia</p> <p><input type="checkbox"/> Facial dysmorphism (Specify below)</p> <p><input type="checkbox"/> Other: _____</p> <p>Eye Defects</p> <p><input type="checkbox"/> Blindness</p> <p><input type="checkbox"/> Coloboma</p> <p><input type="checkbox"/> Epicanthus</p> <p><input type="checkbox"/> Eyelid abnormality (Specify bellow)</p> <p><input type="checkbox"/> Other: _____</p>
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Human Phenotype Ontology

- 10,200+ terms
- 57,000+ links to 5,000+ OMIM Disorders



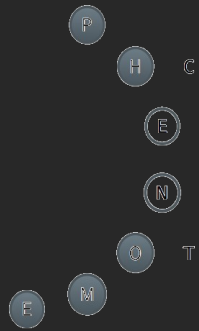
PhenoTips: Key Features

- Ontologies are large (HPO has > 10,000 terms) and difficult to use
- Predictive Search (with synonyms): allowing rapid phenotype entry with HPO terms
- Extensive measurements (e.g. height, weight, eye distance, arm length...), with charting and automated selection of relevant phenotypic terms
- Built-in Diagnosis Assistance: suggest a list of differentials to consider
- Built-in Exam Assistance: suggest related but unselected phenotypes, so none are missed

What PhenoTips Enables

- Fast search of previously seen patients to identify cohorts for retroactive studies
- High-throughput identification of genetic causes of disorders via mining of HPO/nosology connections
- Effective sharing of patient phenotype data between many databases (Global Alliance/IRDIRC Matchmaker project)
- Identification of second families with undiagnosed genetic disorders based on phenotype matching

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