

Data sharing in Canada through the Canadian Open Genetics Repository



COGR
Canadian Open
Genetics Repository

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the Canadian Open Genetics Repository Working Group*

opengenetics.ca

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Ontario **Genomics** Institute

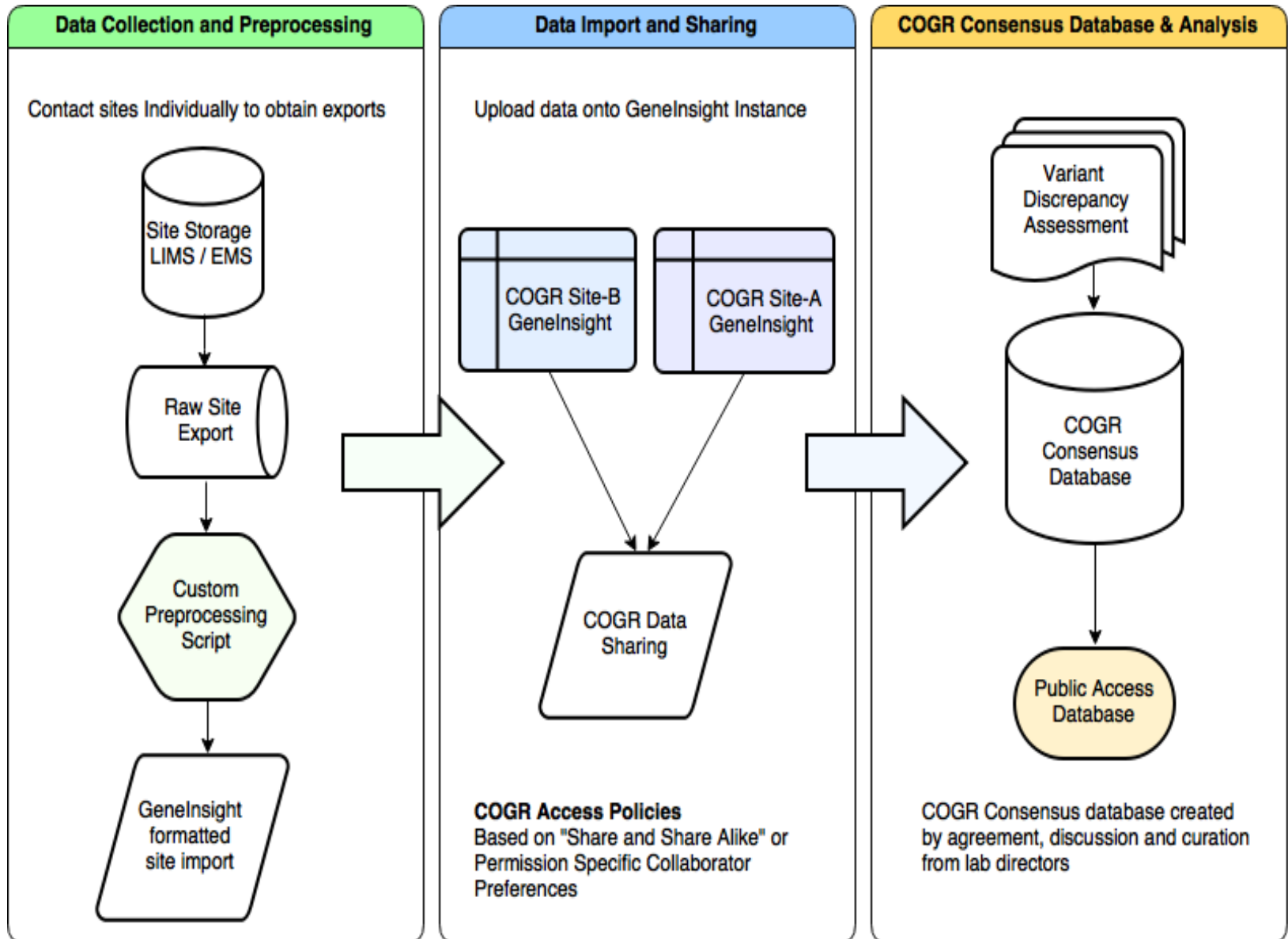


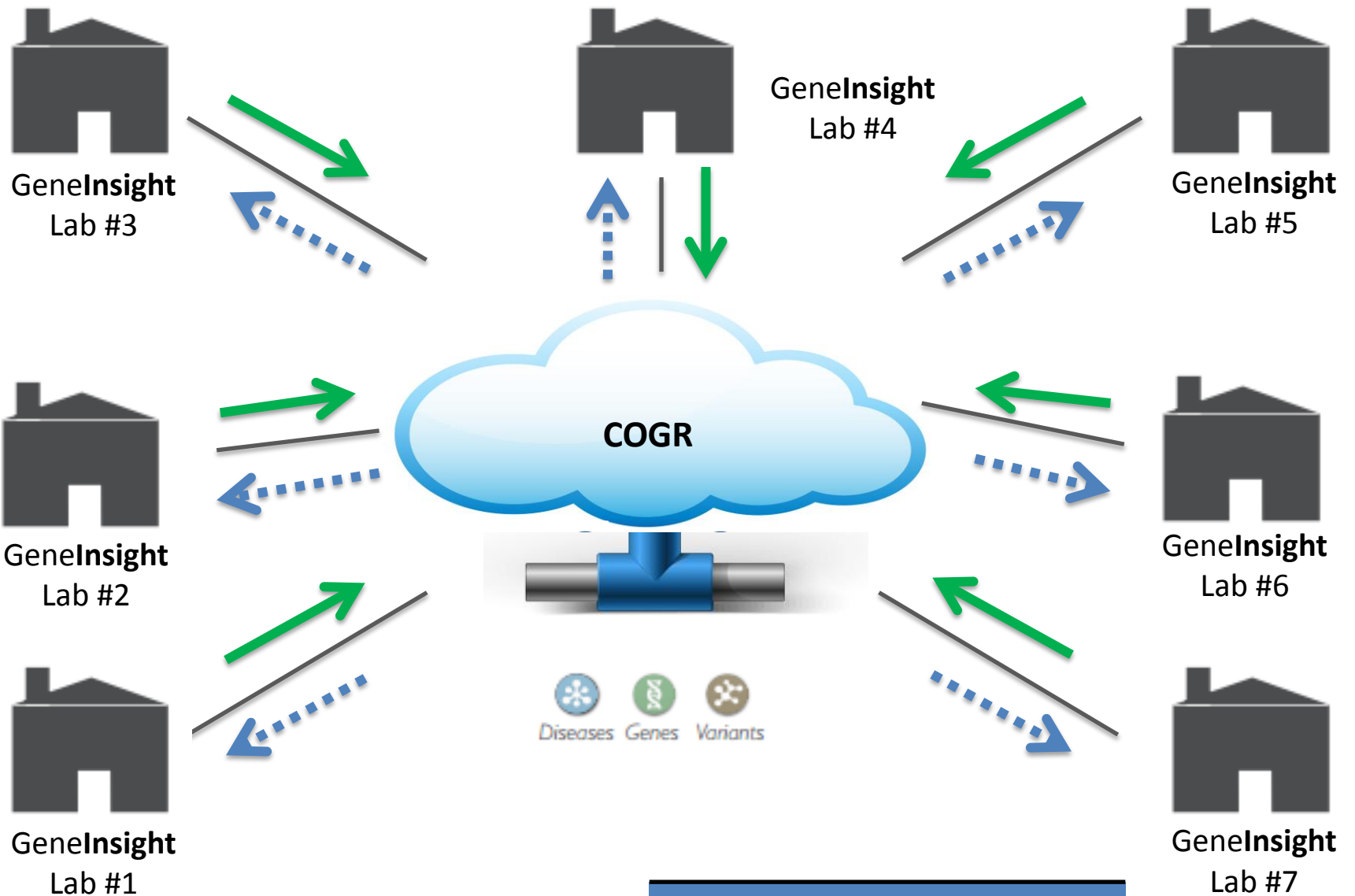
GenomeCanada



Objectives

- To create Canada's unified, open-access, clinical-grade genetic database using a commonly shared platform
- **Standardize** variant assessment procedures
- **Data extraction** and variant classification **consensus** building
- **Data Access** and **Disseminate** results to a large public repository

Workflow Overview

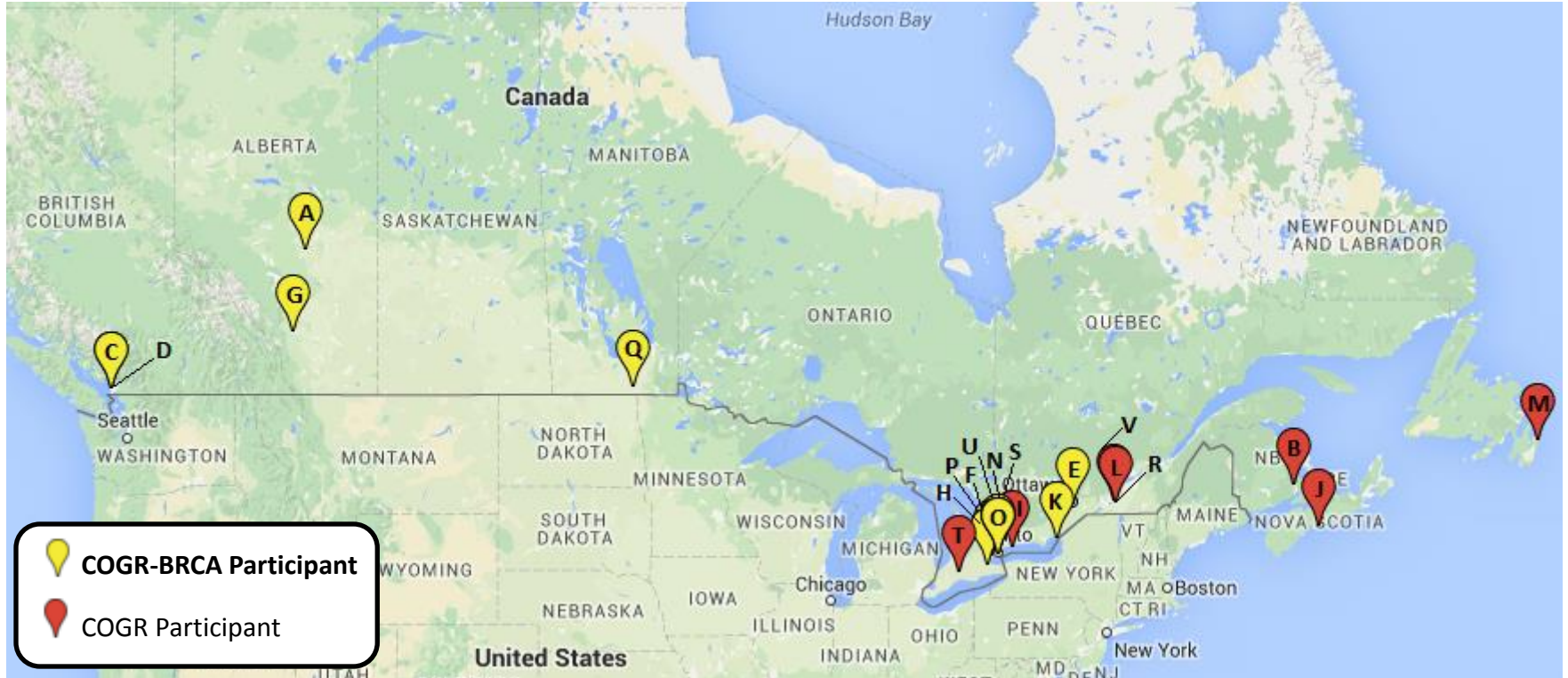




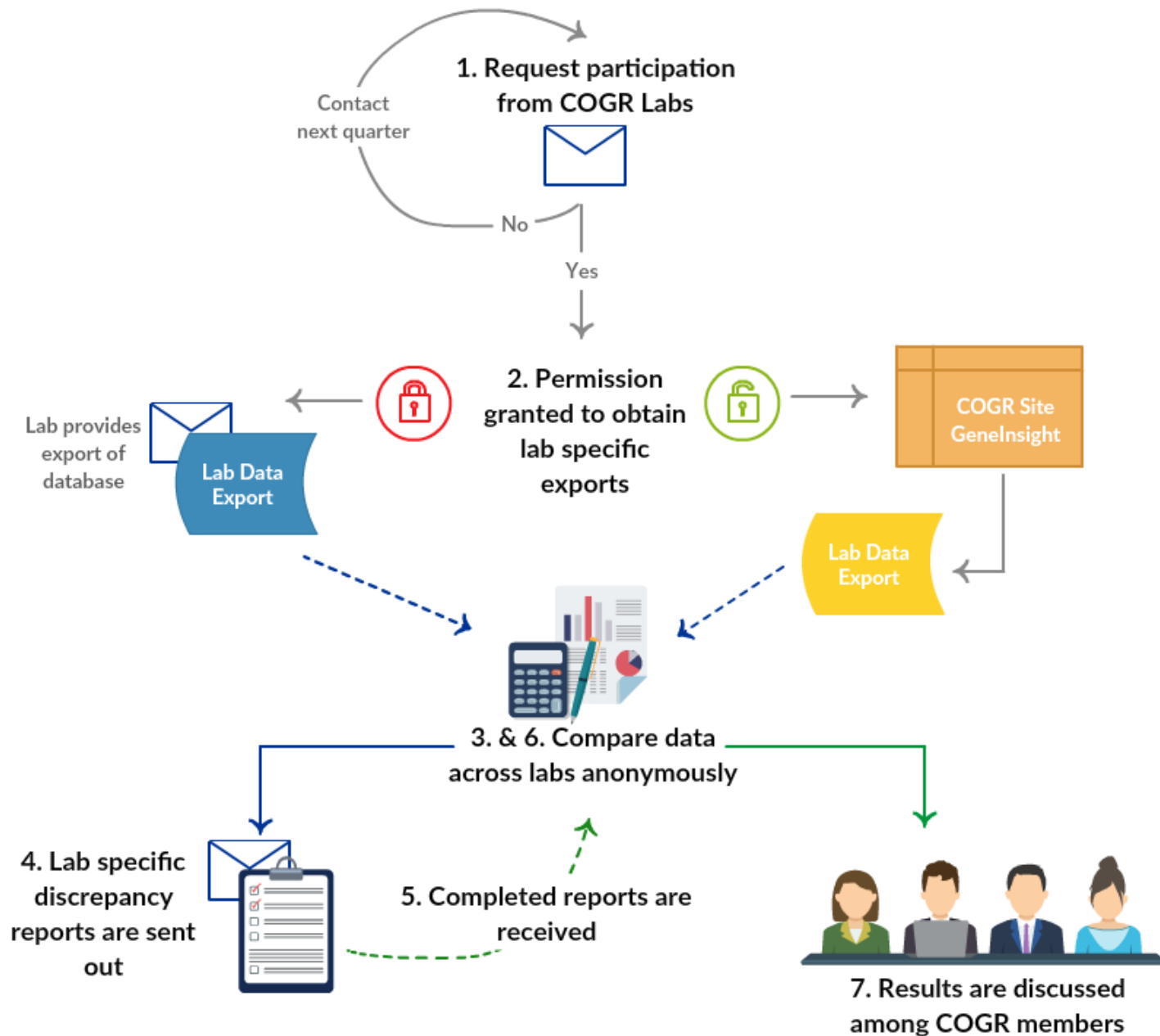
 Clinically validated, de-identified, and approved data
 Read-only viewing with ability to validate and import

Shared Values	#
Interpreted Variants	~20,000
Genes	1,266
Diseases	66

COGR - BRCA Participating Institutions



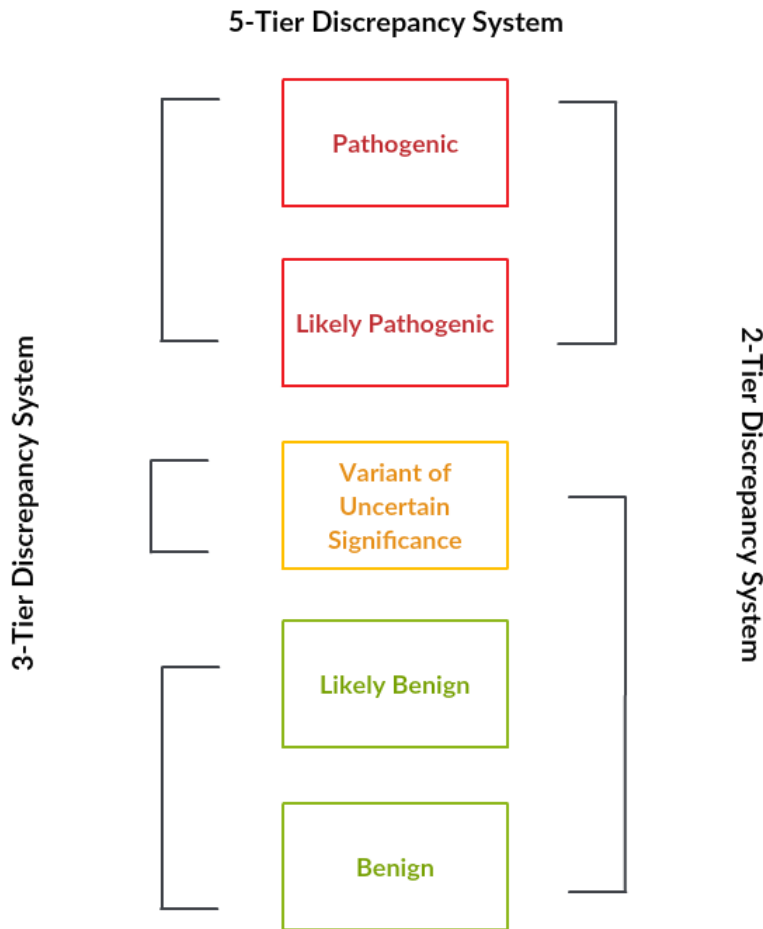
- A. Alberta Children's Hospital (Calgary, AB)**
- B. Atlantic Cancer Research Institute (Moncton, NB)**
- C. British Columbia Cancer Agency (Vancouver BC)**
- D. Children's & Women's Health Centre of BC (Vancouver BC)**
- E. Children's Hospital of Eastern Ontario (Ottawa ON)**
- F. Credit Valley Hospital, Trillium Health Centre (Mississauga ON)**
- G. Dept of Medical Genetics, University of Alberta (Edmonton, AB)**
- H. Hamilton Health Sciences, McMaster University (Hamilton, ON)**
- I. Impact Genetics Inc. (Bowmanville, ON)**
- J. Izaak Walton Killam Health Centre (Halifax, NS)**
- K. Kingston General Hospital, Queen's University (Kingston, ON)**
- L. McGill University Health Complex (Montréal, QC)**
- M. Memorial Health University Medical Center (St. John's, NL)**
- N. Mount Sinai Hospital, University of Toronto (Toronto, ON)**
- O. North York General Hospital (Toronto ON)**
- P. Ontario Institute of Cancer Research (OICR) (Toronto, ON)**
- Q. Regional Health Authority, University of Manitoba (Winnipeg, MB)**
- R. Sainte-Justine Hospital, University of Montreal (Montréal, QC)**
- S. SickKids Hospital and McLaughlin Centre (Toronto, ON)**
- T. University Hospital, Western University (London, ON)**
- U. Women's College Hospital, University of Toronto (Toronto, ON)**
- V. Jewish General Hospital, Montreal (Montréal, QC)**



Discrepancy Report Results Overview

- **11** participating labs across 4 provinces
 - ON, BC, AB, MB
- Received total of **5,554** *BRCA1/2* variants
 - 3014 unique variants
 - 1,148 seen in >2 labs
 - 110 to 1072 variants per lab (505 on average)

Tiered Discrepancy Models



5-tiered

- Initially implemented

3-tiered

- Recommended by participants

2-tiered

- Clinical management

Data Received

5-tier

- 900 variants had 2 or more classifications
- 350 (38.9%) were discordant
 - 1410 observations of these variants

3-tier

- 240 (26.7%) discordant

2-tier

- 45 (5%) discordant

After analysis

330 (23.8%) discordant variant observations changed classification after analysis

5-tier

- Number of discordant variants decreased from 38.9% to 30.7%

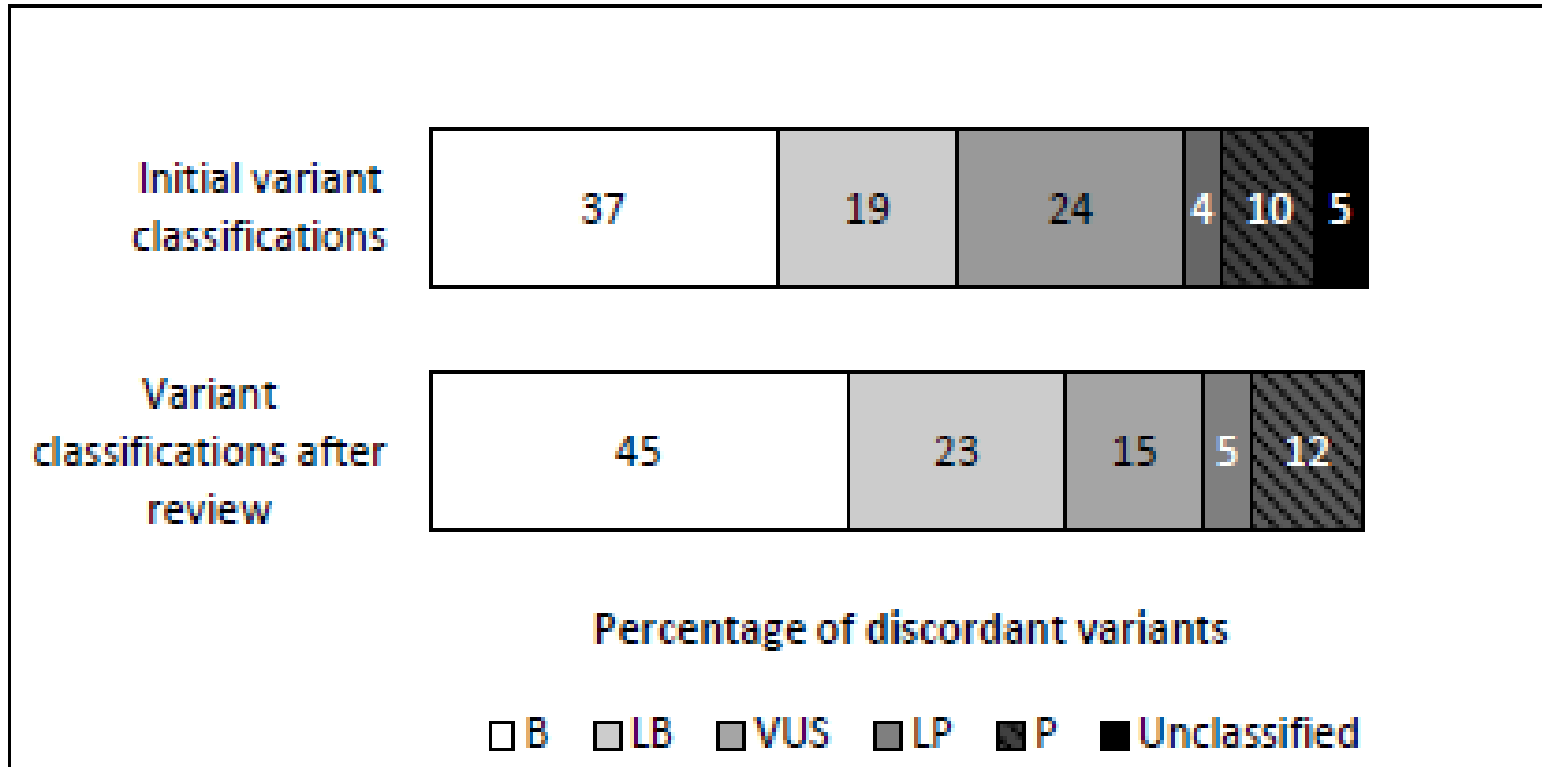
3-tier

- Decrease from 26.7% to 12.45%

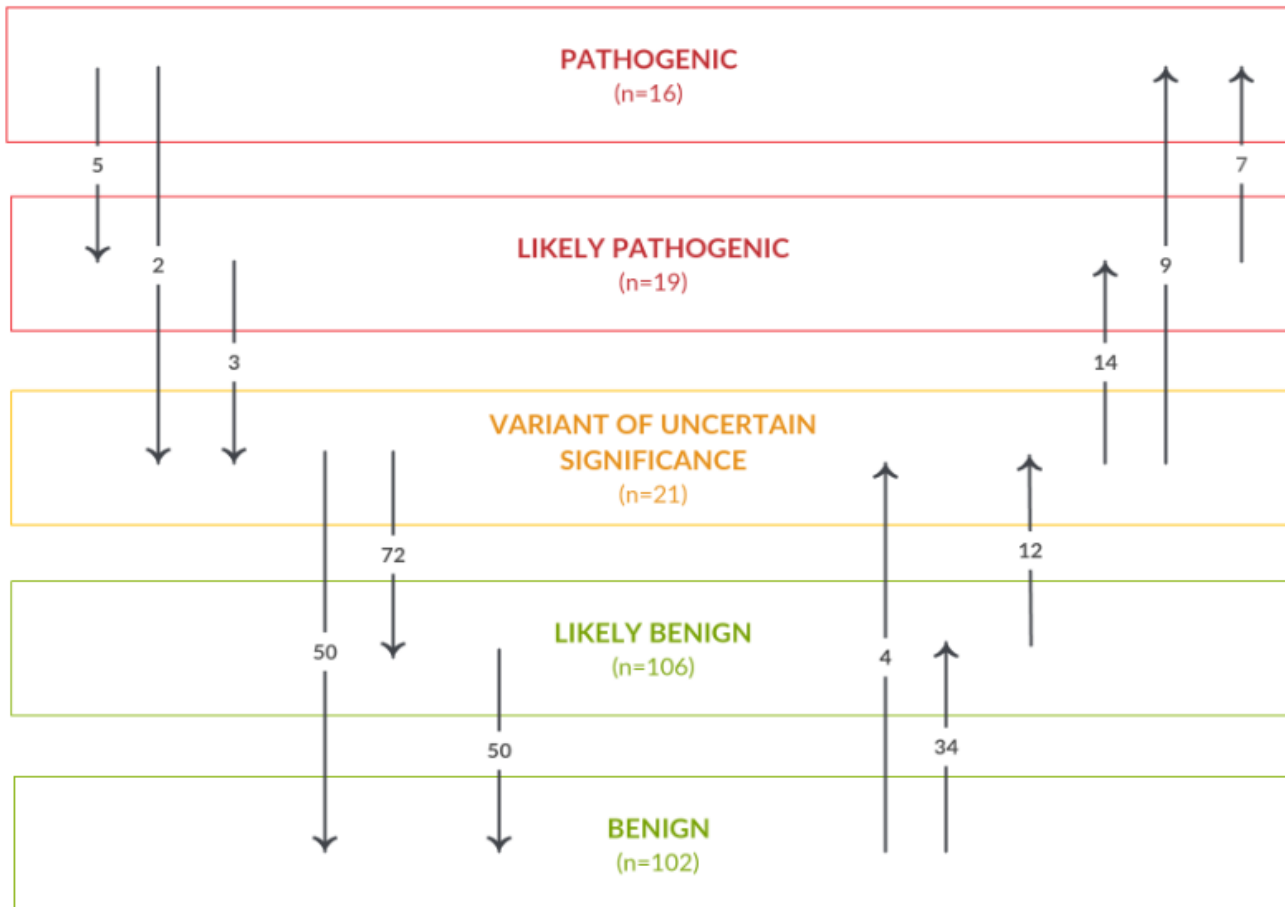
2-tier

- Decrease of from 5% to 1%

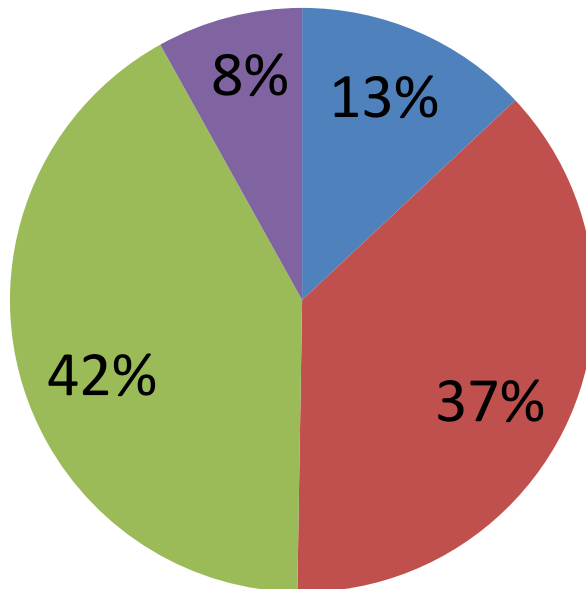
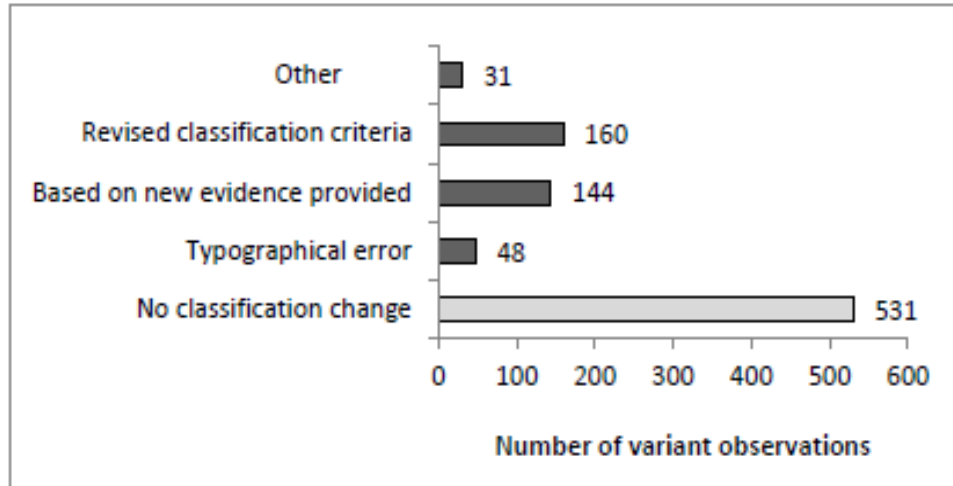
Classification Changes



Direction of Classification Changes

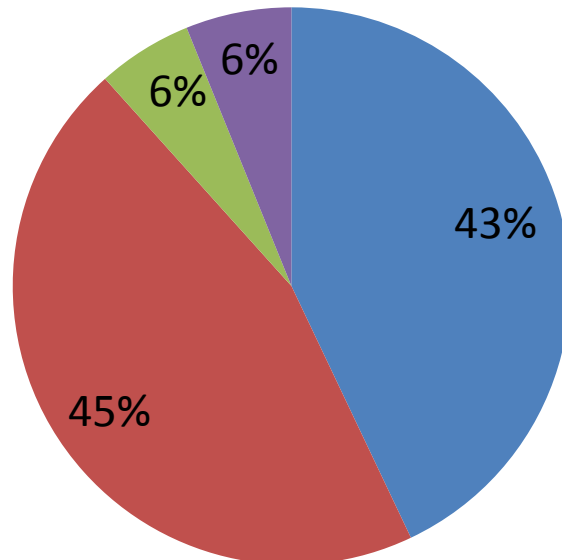
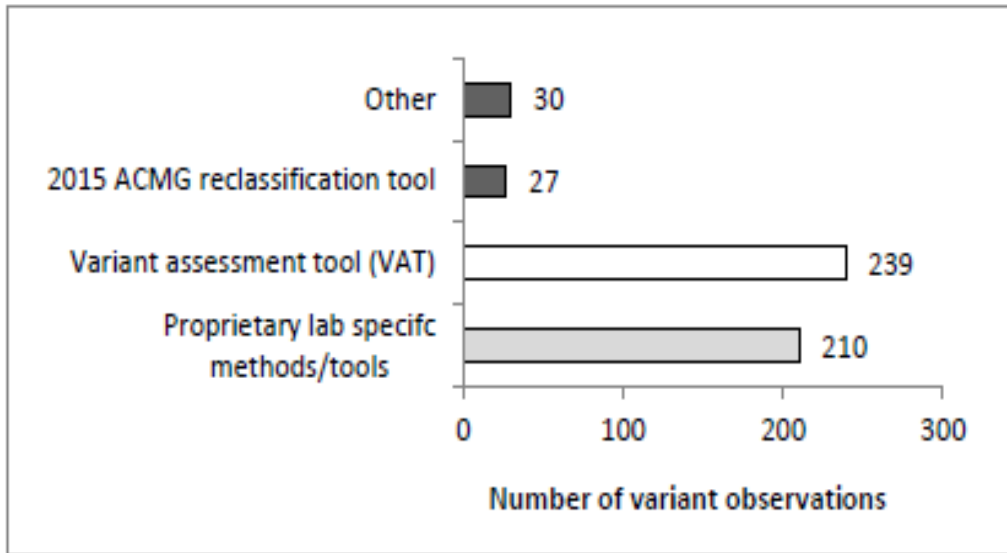


Reason for Reclassification



- Typographical error
- Based on new evidence provided
- Revised classification criteria
- Other (Please specify)

Methods for Reassessment



- We used our own methodology or tools to reassess
- We used the COGR variant assessment tool (VAT)
- We used the ACMG reclassification tool provided

Take Home

- A Canadian inter-institutional quality improvement program
- Importance of periodic review, tracking and maintaining versioned variant information including for variant classification
- mandatory data submission - proficiency testing, laboratory accreditation
- Real-time reporting of variant data - quality assessment program
- We encourage individual institutions to share their data holdings – generate, maintain and preserve knowledge

Collaborating Organizations



GenomeCanada



OntarioGenomicsInstitute



American College of Medical
Genetics and Genomics
Translating Genes Into Health®

ClinVar

Clinically relevant variation

```
CTGATGGTATGGGGCCAAGAGAT  
AGGTACGGCTGTCATCACTTAGAC  
AGGGCTGGGATAAAAGTCAGGGC  
CATGGTGCATCTGACTCCTGAGG  
CAGGTTGGTATCAAGTTACAAGA  
GCCTGACTCTCTCGCCTATTGG
```



Beacon Network



ClinGen

Clinical Genome Resource



DNASTACK



BRCA
EXCHANGE



THE
HUMAN VARIOME
PROJECT

VARIANTWIRE



Global Alliance
for Genomics & Health