

# Canadian Prenatal Genomic Microarray and Sequencing Symposium

*Working Towards a Consensus*

October 2, 2014



**8:30 am - Auditorium**

*Exploring the Key Issues*

**12:00 pm - Gallery**

*Lunch*

**12:45 pm - Auditorium**

*Reaching Consensus in the UK and Belgium*

**2:00 pm – Gallery**

*Breakout Session to Reach Consensus*

**4:00 pm - Auditorium**

*Group Presentations*

**5:10 pm – 13th Floor**

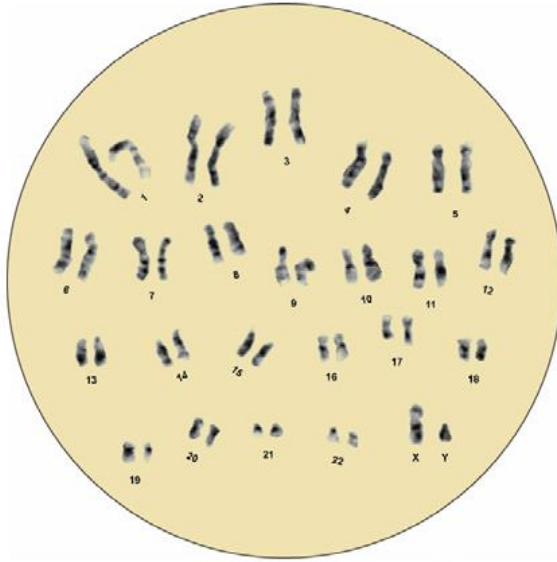
*Wine and Cheese*



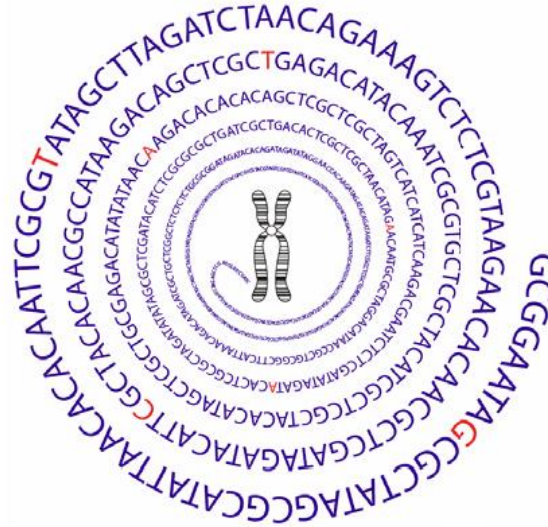
McLaughlin Centre  
UNIVERSITY OF TORONTO

# Why are you here?

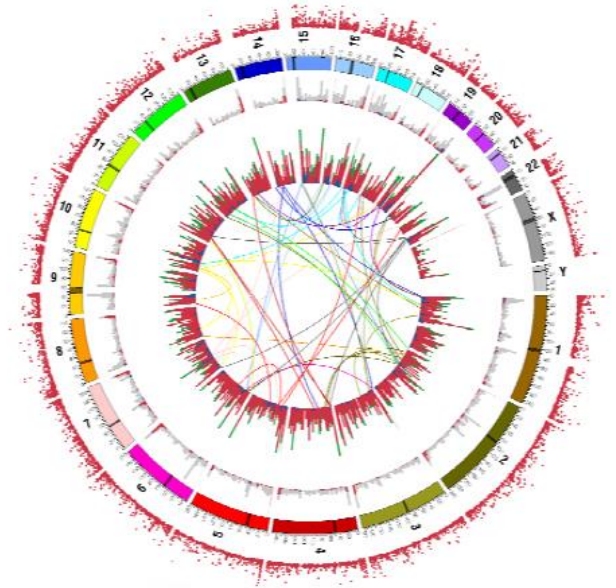
**Relax, contemplate, deliberate on prenatal genomics**



**Chromosomes and alterations**



**SNPs and gene mutations**



**Unified genetic variation maps of genome  
i.e. genomics  
*Nature 2006;, Nature 2010***

## Deliverables?

**Good Canadian common sense**

**(and maybe a paper and some funding ideas)**



# McLaughlin Centre

[Home](#) [Director's Message](#) [Governance](#) [News](#) [Operations](#) [Announcements](#) [Partnerships](#) [Contact](#)

## *Advancing Genomic Medicine through Research and Education*



## News

MCLAUGHLIN CENTRE DIRECTOR  
NAMED POTENTIAL NOBEL PRIZE  
RECIPIENT

[...more](#)

THE ROYAL SOCIETY OF CANADA  
ANNOUNCES 2014 MCLAUGHLIN  
MEDALLIST

[...more](#)

[More News](#)

## Announcements

MCLAUGHLIN CENTRE BIENNIAL  
REPORT 2013-15

[...more](#)

PRENATAL GENOMIC  
DIAGNOSTICS SEMINARS 2014

[...more](#)

[More Announcements](#)



# Key McLaughlin Centre Meetings



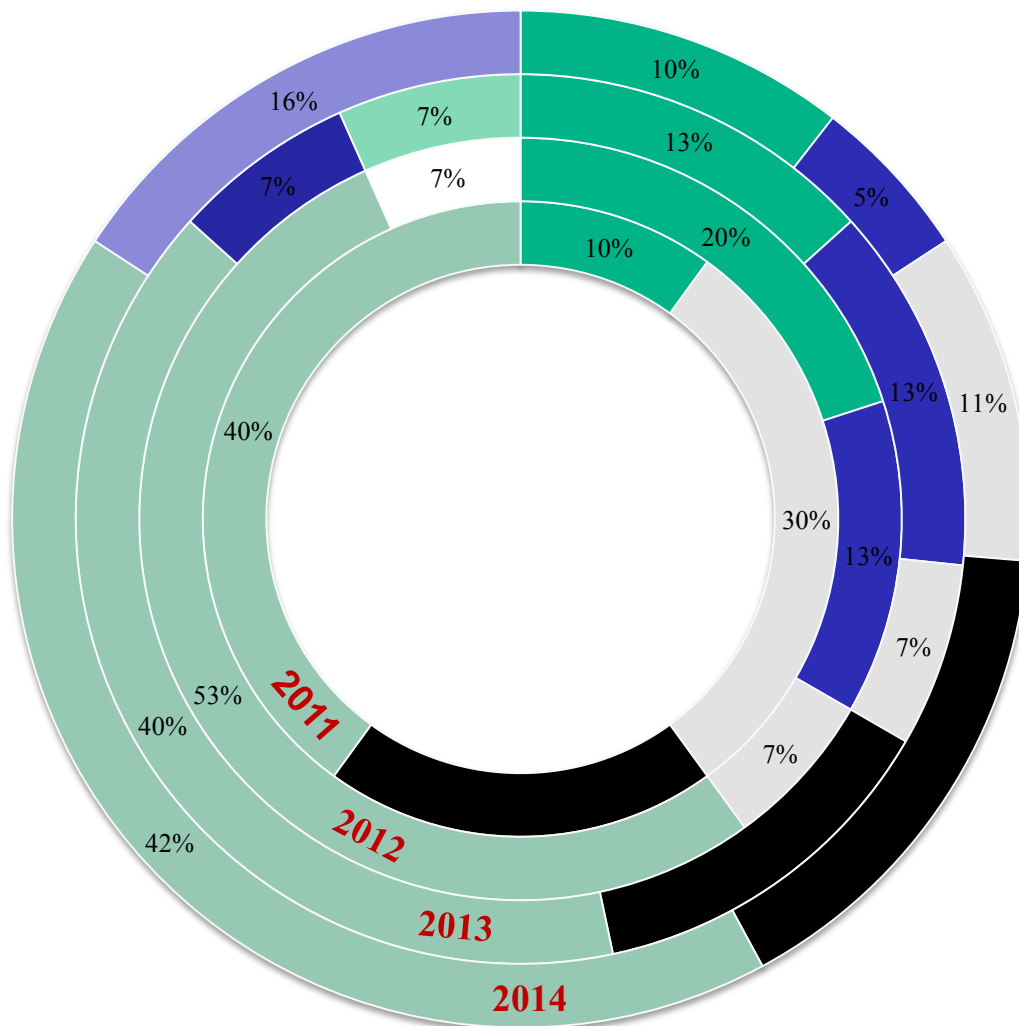
- Genome Structural Variation Symposium, July 2005
- Microarray Use in Canadian Clinical Cytogenetics Laboratories, July 2010
- Genetic risk Factors for Autism: Translating Discoveries into Diagnostics, September 2010
- **Canadian Prenatal Genomics Microarray and Sequencing Symposium, October 2014**



## McLaughlin Center Genomic Medicine Peer-Reviewed Accelerator Grants by Lead Institutions 2011-14

<u>Year</u>	<u>Total Funded</u>
<b>2011</b>	<b>10</b>
<b>2012</b>	<b>15</b>
<b>2013</b>	<b>15</b>
<b>2014</b>	<b>19</b>
<b>Total:</b>	<b>59</b>

### **Institutions receive funding**



# Launch of the Personal Genome Project

# Globe and Mail Saturday December 8<sup>th</sup>, 2012

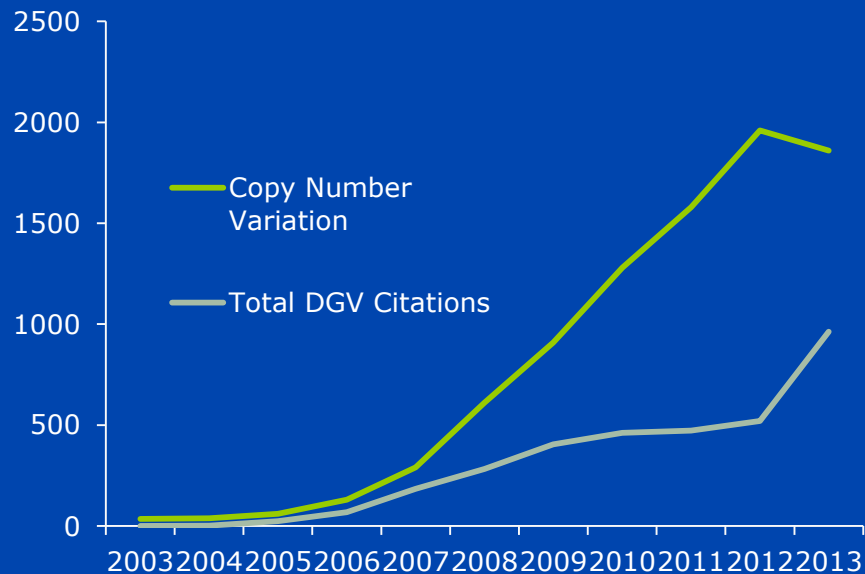


*“Canada is the only G8 country with no law against genetic discrimination”*

- PGP-C is a joint venture between the University of Toronto McLaughlin Centre, The Centre for Applied Genomics at the Hospital for Sick Children, Medcan Clinic and many other important partners, most notably the USA-PGP.

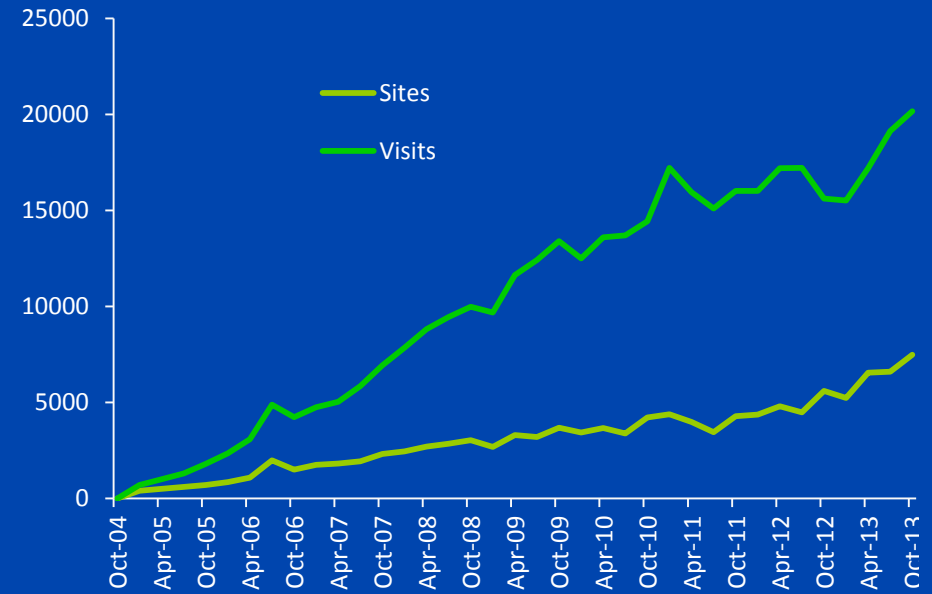
# Database of Genomic Variants

Post 2004 Publication/Citations



DGV is acknowledged >400 times per year in the peer-reviewed literature.

Visits to The Database of Genomic Variants



*Nucleic Acids Research*, 2013, 1–7  
doi:10.1093/nar/gkt958

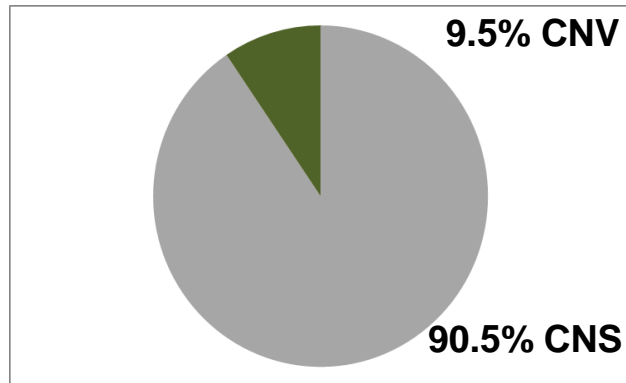
## The database of genomic variants: a curated collection of structural variation in the human genome

Jeffrey R. MacDonald<sup>1</sup>, Robert Ziman<sup>1</sup>, Ryan K. C. Yuen<sup>1</sup>, Lars Feuk<sup>2,\*</sup> and Stephen W. Scherer<sup>1,3,\*</sup>

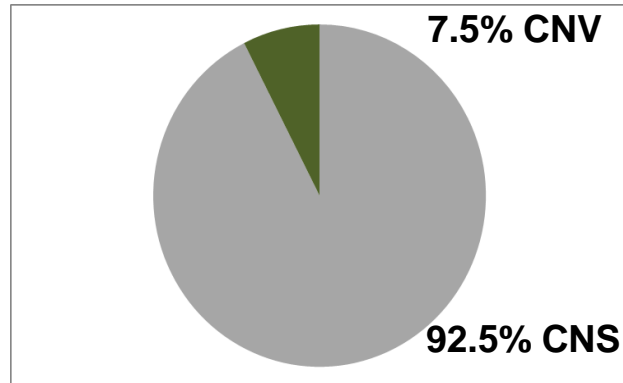
<sup>1</sup>The Centre for Applied Genomics, Peter Gilgan Centre for Research and Learning, The Hospital for Sick Children, 686 Bay Street, Toronto, Ontario M5G 0A4, Canada, <sup>2</sup>Department of Immunology, Genetics and Pathology, Science for Life Laboratory, Uppsala University, Uppsala SE-751 08, Sweden and <sup>3</sup>Department of Molecular Genetics, University of Toronto, Toronto, Ontario M5S 1A8, Canada

# A Chromosome Imbalance Map of the Human Genome

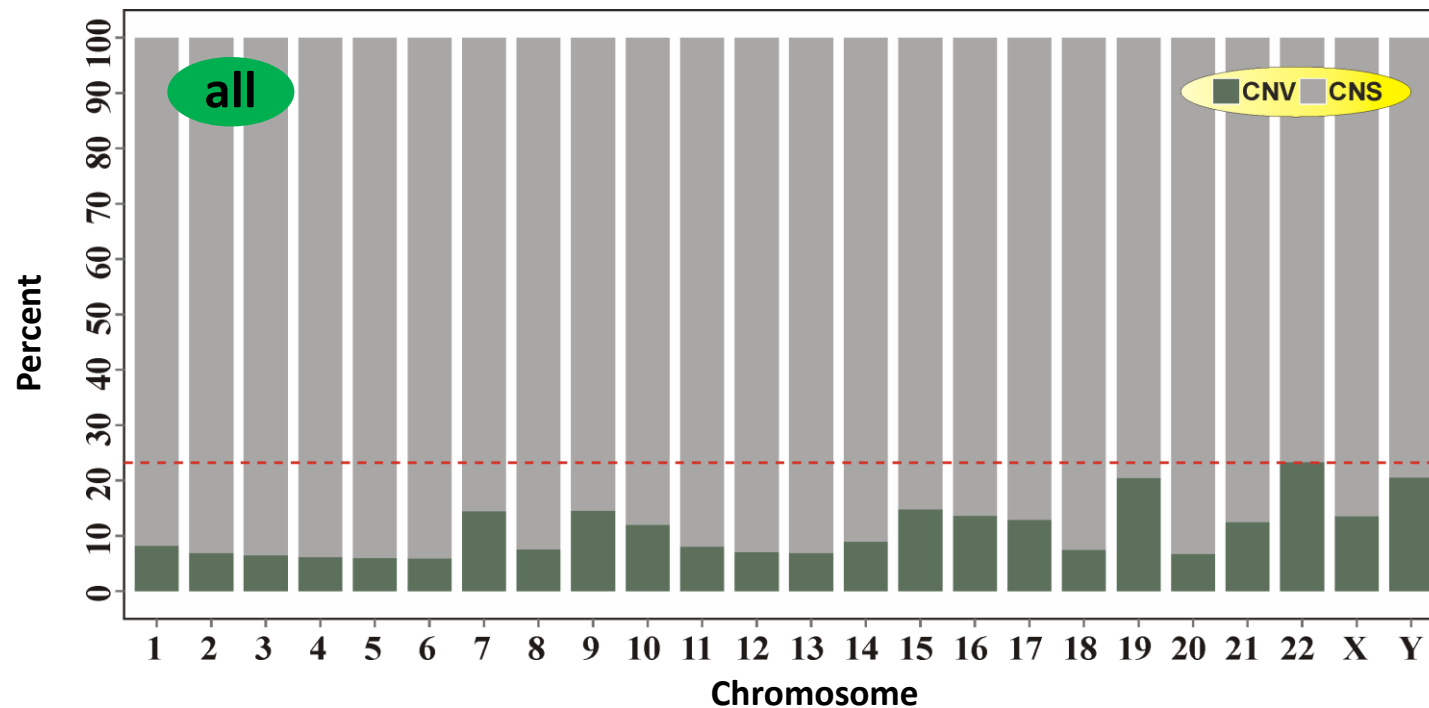
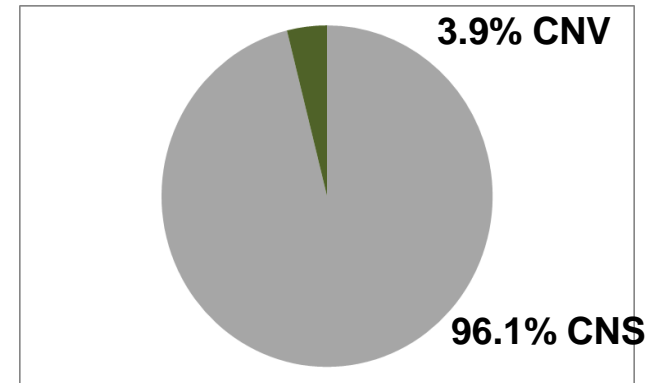
All variants



Loss



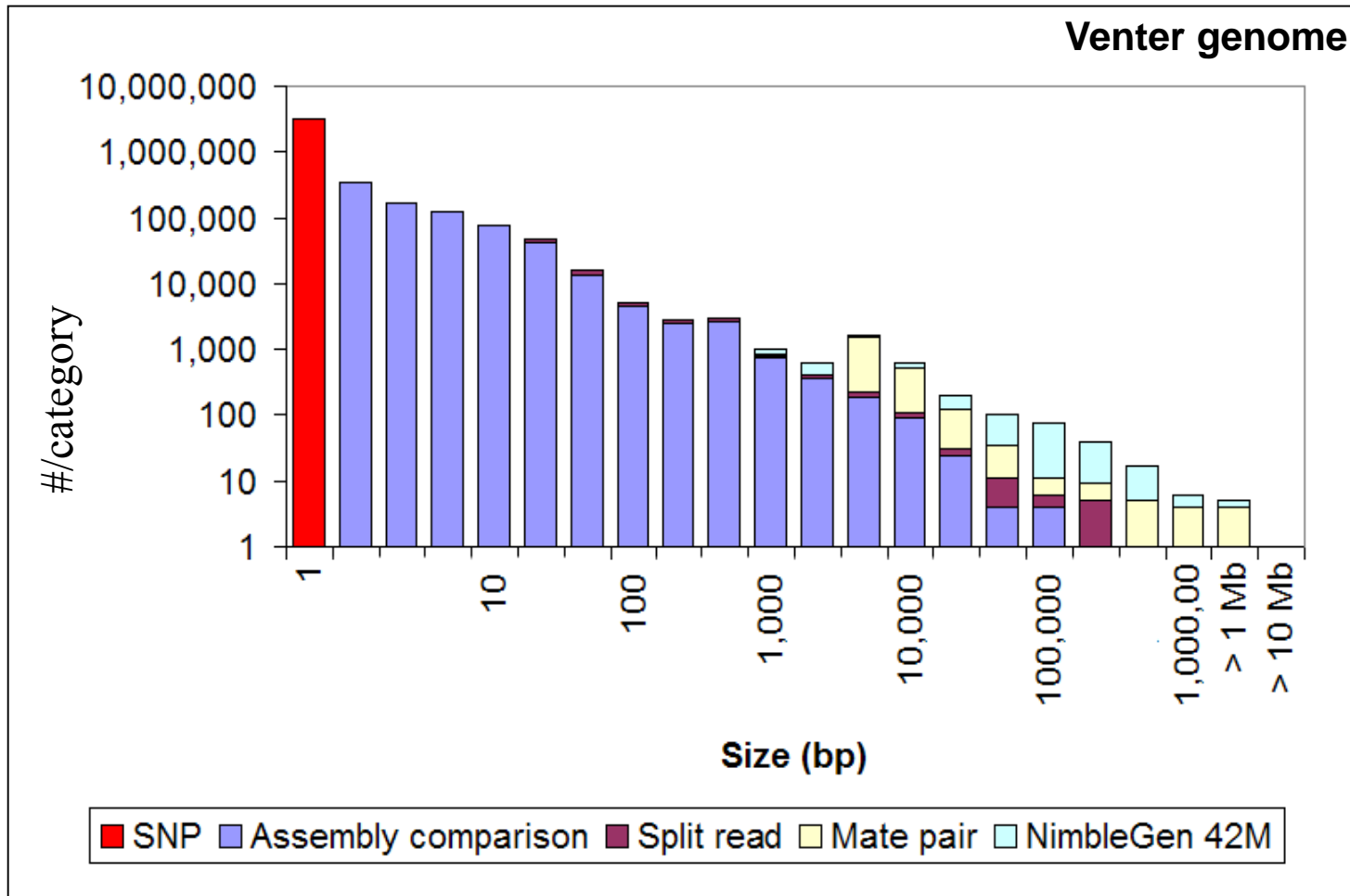
Gain



*Unpublished data*



# How much genetic variation per genome?



-3,213,401 bp (0.11%) variable due to SNPs

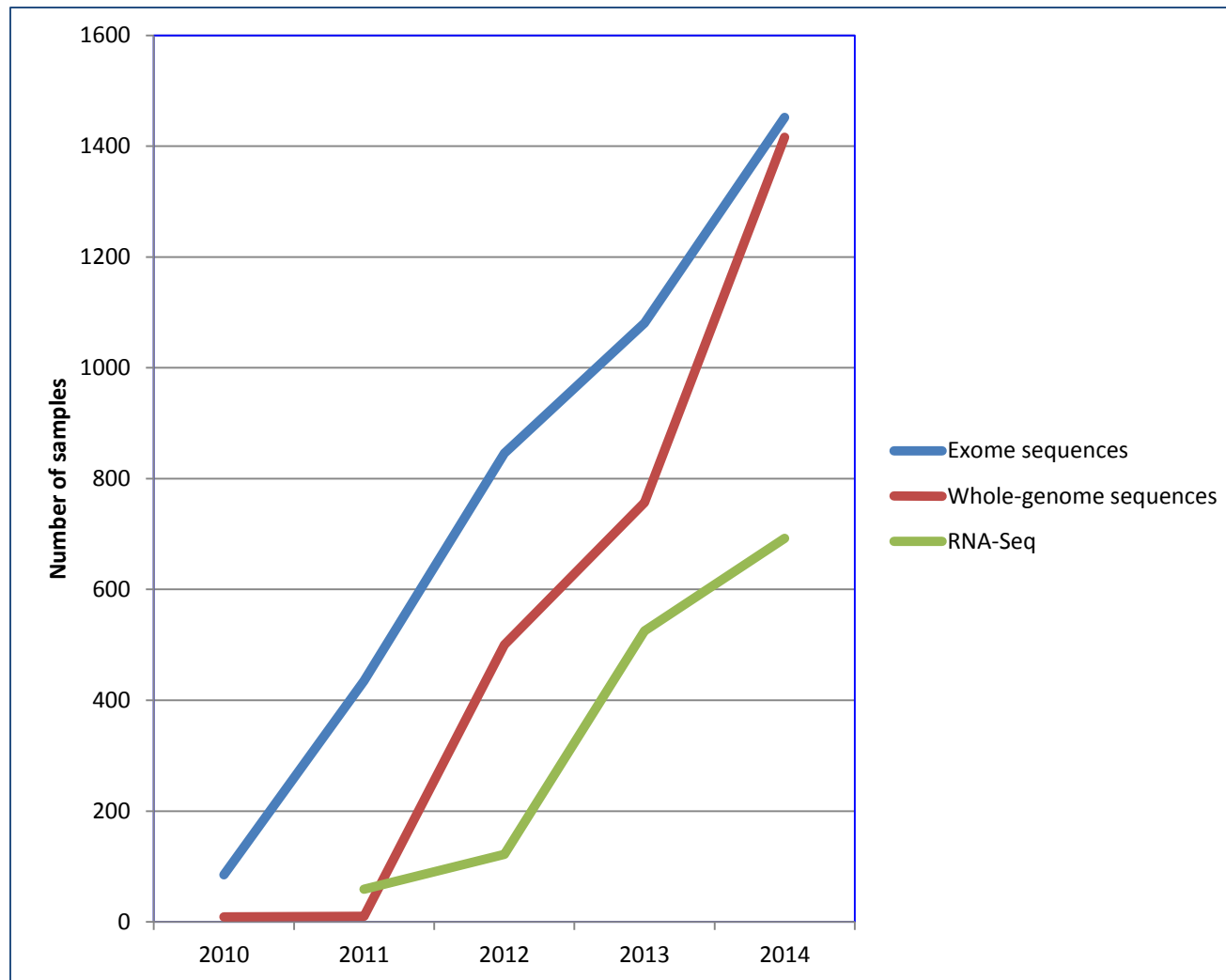
-40,568,593 bp (807,999 events; 1.35%) CNV/indel

-0.3% as inversion

-4,867 genes (~20%) impacted by CNV/indel

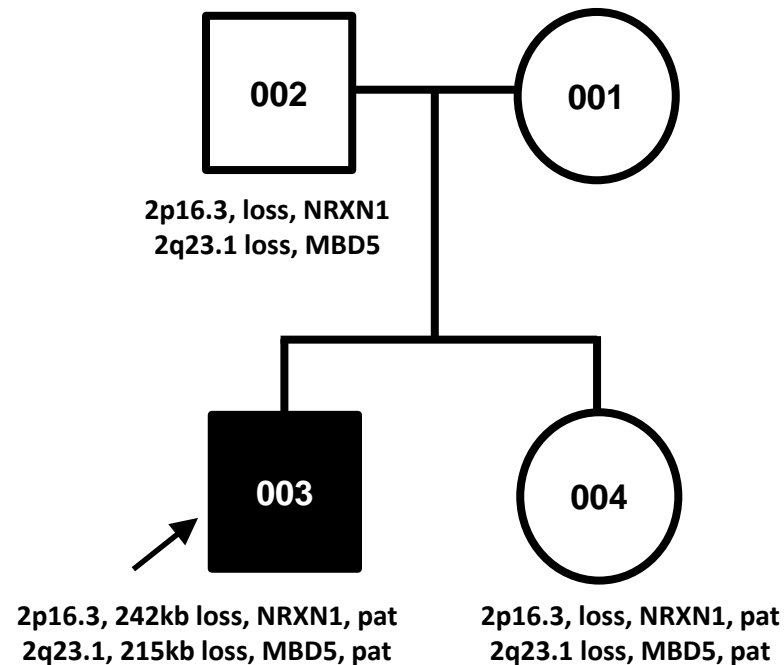
*With Venter team  
Science, 2003  
Nature Genetics, 2006  
PLOS Biology, 2007  
Genome Biology, 2010  
Human Mutation, 2013*

# Clinical and Research: Shift to Genome-wide Analysis



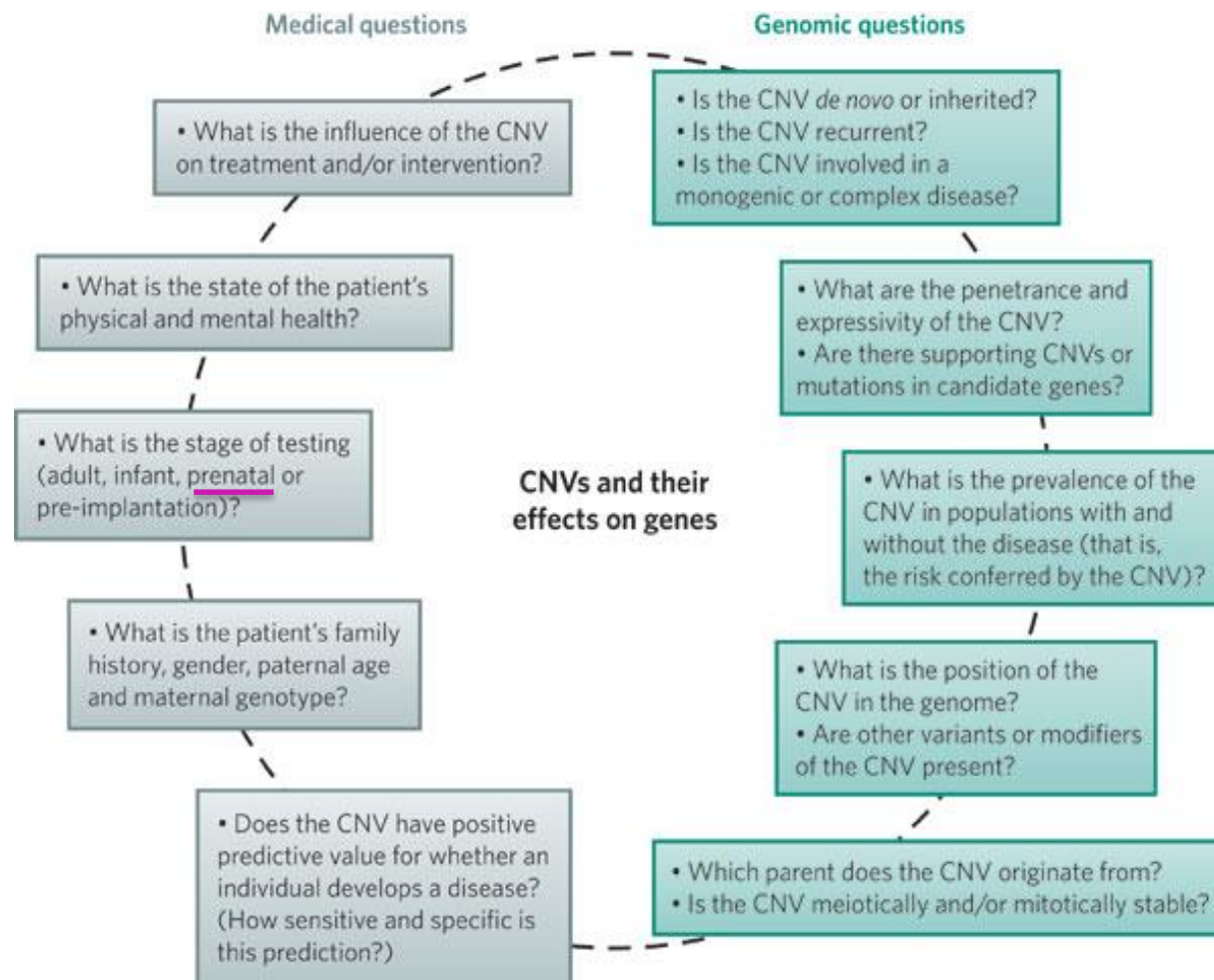
Data from SickKids Hospital (DPLM + TCAG)

# A Clinical Genomic Enigma Wrapped in a Question Mark? *(especially in prenatal setting)*



**...and a terrific research opportunity.**

# Clinical Context of CNVs (and other genetic variants)



*Nature, 2008*