Agenda

9:30 am	Opening Remarks: **Steve Scherer**, Director, The Centre for Applied Genomics (TCAG), The Hospital for Sick Children and Director, McLaughlin Centre for Molecular Medicine, University of Toronto
9:45 am	**John McPherson**, Director Cancer Genomics, Ontario Institute for Cancer Research (OICR) *The Changing Landscape of High Throughput Sequencing*
10:30 am	**Benjamin J. Blencowe,** Professor, Banting and Best Department of Medical Research, Department of Molecular Genetics, CCBR, University of Toronto *Using high throughput sequencing to study alternative splicing complexity and regulation*
11:00 am	Morning Break (please visit sponsors' tables)
11:15 am	**Andrew Smith**, Banting and Best Department of Medical Research, University of Toronto *Highly Multiplexed Barcode Sequencing*
11:45 am	**Mehrdad Hajibabaei**, Director of Technology Development, Biodiversity Institute of Ontario (BIO) University of Guelph *Next-generation Biodiversity Analysis*
12:15 am	Lunch
1:15pm	**Mike Brudno**, Assistant Professor and Canada Research Chair in Computational Biology, University of Toronto *Finding Insertion and Deletion Variants with NGS Data*
1:45 pm	**David Guttman**, Director, Centre for the Analysis of Genome Evolution and Function and Canada Research Chair in Comparative Genomics, University of Toronto *Tying Down Arabidopsis Mutants with Chastity Belts*
2:15 pm	Break (please visit sponsors' tables)
2:45 pm	**Ben Evans**, Assistant Professor, Department of Biology, McMaster University *Next Generation Insights into Evolution of Polyploid Frogs (Xenopus)*
3:15 pm	**Nils Homer**, Department of Computer Science, UCLA *U87MG Decoded: The Genomic Sequence of a Cytogenetically Aberrant Human Cancer Cell Line*
3:45 pm	Closing Remarks: **Sergio Pereira**, Facility Manager Next Generation Sequencing, TCAG
4:00 pm	End

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