



AGENDA

October 3, 2008	
8:00 am	Registration, breakfast with Exhibitors & Sponsors
9:00 am	Joan Kean-Howie, Director General, NRC-IMB, Welcome
Session 1 – Technologies on the Leading Edge Session Chair: Robert K. Boyd, Researcher Emeritus, NRC	
9:05 am	Keynote Address – Christie Hunter , Applied Biosystems <i>Targeted Protein Expression Profiling using MRM: Biological Variation of Plasma Protein Levels Assessed using Twin Samples</i>
9:50	Jack Henion – Advion BioServices Inc <i>A Novel Chromatographic Method Allows Online Reanalysis In Proteomic Investigations</i>
10:15	Coffee Break & Poster Session
10:45	Mark Wall – Dalhousie University <i>Organic/aqueous Solvent System for Incomplete Tryptic Digestion and Improved Protein Identification</i>
11:10	David Barnett – Atlantic Cancer Research Institute <i>An alternative peptide precursor ion selection strategy for protein identification by mass spectrometry</i>
11:35	John Marshall , Ryerson University <i>Human Serum Proteins Fractionated by Preparative Partition Chromatography prior to LC-ESI-MS/MS</i>
12:00	Lunch with Exhibitors & Sponsors - Posters
Session 2 – Biological Insights Through Proteomics Session Chair: Alan Doucette	
1:30	Keynote Address – Guy Poirier , Université de Laval <i>Systems Biology of Poly(ADP-ribose) binding proteins</i>
2:20	Patrick Murphy – Dalhousie University/NRC <i>Quantitative Proteomics of the IGF-1R Signalling Network.</i>
2:45	Coffee Break & Poster Session
3:15	Paola Marignani , Dalhousie University <i>Identification of novel protein partners for the tumour suppressor kinase Ikb1 by mass spectrometry and reverse chemical genetics</i>
3:40	Jeurgen Kast – University of British Columbia <i>Improving Platelet Storage by Forward and Reverse Proteomic Approaches and Drug Intervention</i>
4:05	Jim Fawcett – Dalhousie University <i>Polarity proteins and their role in Schizophrenia - a mass spectrometry approach</i>
4:30	Dev Pinto, Closing Remarks