The 2014 Workshop on Clinical and Applied Proteomics

Friday Oct. 24th, 2014  8:00 am – 5:00 pm

M3.01 - JSL Browne Amphitheater, H4 – Employee/Nurse lounges for booth & social events
The Royal Victoria Hospital, 687 Pine Avenue, Montreal

Clinical Proteomics
The RI-MUHC & McGill University
Table of Contents

Welcome from the Workshop Organizers – P3
Workshop Program – P4
The Bio-sketches of the Speakers – P5-7
Thank You to Our Sponsors/Partners – P8
Booth Layout and Lunch/Drink Stations – P9
Map and parking information – P10
Major Sponsor/Partner Profile & Advertisement – P11
  • The Clinical Proteomics – P12
  • The Systems Biology Training Program – P13
  • Dr. Peter Metrakos Laboratory – P14
  • ThermoFisher Scientific – P15
  • The Research Institute of McGill University Health Centre – P16
  • McGill University-Génome Québec Innovation Centre – P17
  • The Chemical Biology Program at McGill – P18
  • The Faculty of Medicine – McGill University – P19
List of Attendees – P20-25
Welcome From The Workshop Organizers

Dear Workshop Attendees,

It is with a great pleasure that we welcome you to the 2014 Workshop on Clinical and Applied Proteomics. This is our fourth annual Workshop on various aspects of clinical proteomics and affiliated disciplines, all of which have implications in better understanding the mechanisms of disease and potentially better diagnosis and treatment.

The objective of our workshop is to gather together clinicians, researchers and trainees who share an interest in applied proteomics/interactomics from a disease-related perspective, as well as those who are interested in learning new technologies, sharing research ideas with others, interacting and collaborating within the RI-MUHC/McGill and externally in Montreal and beyond.

We are grateful for the participation of our guest speakers (Drs. Mathias Dreger, Thomas Kislinger, Liang Li, Gavin Reid and Igor Stagljar), who will present their latest discoveries and new trends in Clinical and Applied Proteomics.

We would like to acknowledge our sponsors/partners. Without their generous support it would not be possible to hold this Workshop free of charge for the participants.

Finally, we would like to thank Pamela Cameron, Bobbie Carin, Shi Bo Feng, Fariba Kalantari, Stavroula Kalantzis, EunJoo Lee, Lily Li, Alexander Mazur, Daniel Moses, Filomena Sargo for their help and assistance, ensuring the success of this Workshop.

Enjoy the Workshop!

Jing Liu
Kurt Dejgaard
Amani Batarseh
Pierre Chaurand
Peter Metrakos
David Thomas
Tommy Nilsson
The 2014 Workshop on Clinical and Applied Proteomics

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<table>
<thead>
<tr>
<th>Time</th>
<th>Speaker</th>
<th>Affiliation</th>
<th>Presentation or Activity</th>
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<tbody>
<tr>
<td>08:00</td>
<td>All</td>
<td>All</td>
<td>Light Breakfast and Registration</td>
</tr>
<tr>
<td>08:55</td>
<td>Tommy Nilsson</td>
<td>RI-MUHC and McGill University</td>
<td>Welcome and Introductory Remarks</td>
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<tr>
<td>09:00</td>
<td>Thomas Kislinger</td>
<td>University of Toronto</td>
<td>Systematic Development of SRM Assays for the Detection of Aggressive Prostate Cancers</td>
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<tr>
<td>09:40</td>
<td>Liang Li</td>
<td>University of Alberta</td>
<td>Challenges and Opportunities of Metabolomics Research for Clinical Applications</td>
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<tr>
<td>10:20</td>
<td>All</td>
<td>All</td>
<td>Coffee Break and Booth Visit</td>
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<tr>
<td>10:50</td>
<td>Pierre Chaurand</td>
<td>Université de Montréal</td>
<td>Imaging Mass Spectrometry Stratification of Colorectal Cancer Liver Metastasis</td>
</tr>
<tr>
<td>11:30</td>
<td>Mathias Dreger</td>
<td>Caprotec, Berlin, Germany</td>
<td>Profiling of Small Molecule-Protein Interactions using Tri-functional Capture Compounds</td>
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<tr>
<td>12:10</td>
<td>All</td>
<td>All</td>
<td>Lunch and Booth Visit</td>
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<tr>
<td>13:45</td>
<td>Amani Batarseh</td>
<td>RI-MUHC and McGill University</td>
<td>A Shotgun Lipidomics Approach to Study NAFLD</td>
</tr>
<tr>
<td>14:00</td>
<td>Gavin Reid</td>
<td>Michigan State University/University of Melbourne</td>
<td>Deciphering the Functional Role of Lipidome Profile Alterations in Colorectal Cancer Malignancy and Metastasis</td>
</tr>
<tr>
<td>14:40</td>
<td>Igor Stagljar</td>
<td>University of Toronto</td>
<td>Membrane Protein Interaction Networks in Health &amp; Disease: the EGFR Story</td>
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<tr>
<td>15:20</td>
<td>David Thomas</td>
<td>McGill University</td>
<td>Closing Remarks</td>
</tr>
<tr>
<td>15:25</td>
<td>All</td>
<td>All</td>
<td>Door Prize Announcement, Wine and Cheese</td>
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Bio-sketches of the Speakers

Dr. Pierre Chaurand (Ph.D. 1994, Université Paris Sud, Orsay, France) is Associate Professor of Chemistry at Université de Montréal (2009 - present). His expertise’s are in fundamental and analytical MS. He is one of the pioneers of the IMS technology. His research interests are focused on the development of new strategies and methods to improve the specificity and sensitivity of IMS with applications in clinical biology. Dr. Chaurand has over 65 peer reviewed publications in the field of MS, with over 35 in the specific field of IMS including several with cancer relevance.

Dr. Mathias Dreger heads the CCMS Biology department at the biotechnology company caprotec bioanalytics GmbH (Berlin, Germany). Prior to his affiliation with caprotec, he performed research in the fields of neurochemistry, subcellular biochemistry, and proteomics, with a focus on subcellular proteomics. He holds a PhD in Biochemistry, obtained at the Free University Berlin Germany. In 2003, he moved on to the University of Oxford, UK, to set up and run a proteomics core facility lab at the Department of Physiology, Anatomy, and Genetics. His focus was novel analytical strategies in proteomics. In 2008 he moved back to Berlin to join caprotec, focusing on chemical proteomics. At caprotec, he is responsible for the design and conduction of capture experiments using tri-functional Capture Compounds to identify the protein targets of small molecules in biological samples.

Dr. Thomas Kislinger received his MSc in Analytical Chemistry from the University of Munich, Germany (1998). He completed his PhD in 2001, investigating the role of Advanced Glycation Endproducts in diabetic vascular complications at the University of Erlangen, Germany and Columbia University, New York. Between 2002 and 2006 he completed a post-doctoral fellowship at the Banting and Best Department of Medical Research at the University of Toronto. In 2006 he joined the Princess Margaret Cancer Center as an independent investigator. Dr. Kislinger holds positions as Senior Scientist at the Princess Margaret Cancer Center and as Associate Professor at the University of Toronto in the Department of Medical Biophysics. He is a Tier 2 Canadian Research Chair in Proteomics in Cancer Research. The research interests in the Kislinger lab are focused on the application of proteomics and allied computational tools. His research interests include the development of novel proteomics technology, the application of proteomics to cancer biology and biomarker discovery and the use of mass spectrometry for the identification of dynamic changes in plasma membrane proteins.
Dr. Liang Li obtained his PhD from the University of Michigan, Ann Arbor, Michigan and joined the University of Alberta in July 1989, where he is now a Professor of Chemistry, Adjunct professor of Biochemistry and Tier I Canada Research Chair in Analytical Chemistry. Dr. Li served as Director, Alberta Cancer Board Proteomics Resource Laboratory, from 2000 to 2005 and is Chair of the Analytical Chemistry Division at the U of A since 2007. Dr. Li’s research interest is in the area of developing analytical mass spectrometry for proteomics and metabolomics applications. He has worked on the development of MALDI-TOFMS for protein and polymer characterization. He has also involved in developing enabling MS techniques and methods for proteome analysis, including a shotgun protein sequence mapping method based on microwave-assisted acid hydrolysis of proteins combined with MS. He is active in developing new methods for quantitative and comprehensive metabolome profiling of bio-systems for biological studies and discovery of disease biomarkers. He has published 201 papers and given over 228 invited lectures. He holds 4 US patents. Dr. Li has served on editorial boards of several journals and is an editor of Analytica Chimica Acta since 2005. Dr. Li has received several awards, including the Rutherford Memorial Medal in Chemistry from the Royal Society of Canada (2003), the F.P. Lossing Award from the Canadian Society for Mass Spectrometry (2006), the Maxxam Award from the Canadian Society of Chemistry (2009) and the Gerhard Herzberg Award from the Canadian Society for Analytical Sciences and Spectroscopy (2010).

Dr. Gavin E. Reid is the Professor of Bioanalytical Chemistry in the School of Chemistry and the Department of Biochemistry and Molecular Biology, and member of the multidisciplinary Bio21 Molecular Science and Biotechnology Institute, at The University of Melbourne, Australia. Over the past twenty-seven years, he has held a number of technical research positions and academic appointments in Australia and the USA, including 10 years (2004-2014) as an Assistant and Associate Professor in the Department of Chemistry and the Department of Biochemistry and Molecular Biology at Michigan State University. The primary research interests of the Reid group are the development of novel analytical biochemistry, mass spectrometry and associated chemical strategies for proteome and lipidome analysis, and their application toward understanding the role of proteins and lipids in disease, including cancer, diabetes and diabetic complications. To date, this inter-disciplinary and highly collaborative research has resulted in 140 peer-reviewed publications, presentations at 77 national and international conferences, 52 invited seminars and 3 patents. Gavin received an American Society for Mass Spectrometry Research Award in 2007, and the Bowie Medal from the Australian and New Zealand Society for Mass Spectrometry in 2011. He is currently an Associate Editor for the Journal of the American Society for Mass Spectrometry, and a member of the editorial advisory board for the European Journal of Mass Spectrometry.
Dr. Igor Stagljar is a Croatian/Canadian molecular biologist, and a Professor in the Departments of Biochemistry and Molecular Genetics at the Donnelly Centre in the University of Toronto, Canada. He is one of the world leaders in the field of interactive proteomics and development of methods to investigate protein-protein interactions. In particular, he is internationally known for the development of the split-ubiquitin Membrane Yeast Two-Hybrid (MYTH) and Mammalian Membrane Two-Hybrid (MaMTH) technologies, powerful tools for the identification of the interactors of membrane proteins and one of the key interactive proteomics technologies. This has lead to many groundbreaking discoveries and the elucidation of functions of various membrane proteins involved in human health and disease. Dr. Stagljar is currently involved in major proteomics projects to map how integral membrane proteins interact to produce either healthy or diseased cells. To that end, his lab is using high-throughput interactive proteomics, genetic, and biochemical tools to understand how cell signaling and membrane transport pathways control cell behavior in normal and disease cells.

Dr. Stagljar is the author of more than 80 PubMed cited scientific papers and is the recipient of several national and international science awards. In addition, Dr. Stagljar is a member of the Editorial board of BioTechniques, Molecular Genetics and Genomics, BMC Biotechnology, Journal of Molecular Biology and Biochemical and Biophysical Research Communications. Lastly, Dr. Stagljar is a co-founder of Dualsystems Biotech Inc, and is currently planning to co-found another company based on the Mammalian Membrane Two-Hybrid (MaMTH) technology.

Dr. Amani Batarseh received her Ph.D. from Georgetown University in 2010 under the supervision of Prof. Vassilios Papadopoulos, working on the Transcriptional Regulation of Translocator Protein 18kDa (TSPO), realizing 6 peer-reviewed publications. Dr Batarseh has had postdoctoral appointments at McGill University 2010-2011 studying the role of TSPO in Benign Prostate Hyperplasia (BPH), followed by a position at Harvard Medical School studying type II diabetes from 2011-2012 before returning to McGill University as a postdoctoral fellow in the lab of Prof. Tommy Nilsson since January 2013. Dr Batarseh’s current role is to establish a shotgun Lipidomics approach within the Clinical Proteomics Platform focusing on studying Non-Alcoholic Fatty Liver Disease (NAFLD), being a hidden disease characterized by steatosis and its progression to Non-Alcoholic Steatohepatitis (NASH). Since NAFLD is a silent disease characterized by fat accumulation in the liver, with liver biopsy being the most definite test to identify its presence, Amani aims to find diagnostic and prognostic markers that could predict the disease earlier by investigating changes in the lipidome of liver lipid droplets from different stages of the disease using a QExactive mass spectrometer, and to find correlations with lipid changes in other bodily fluids such as plasma or urine, since they are more accessible and less invasive than a liver biopsy. This work is being done as a close collaboration with Thermo Fisher Scientific as part of a consortium of Scientists and Doctors at McGill University and University of Montreal, as well as training and exchange with Dr Andrej Shevchenko at the Max Planck Institute in Germany.
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Booth Layout
Nurse/Employee Lounges, H4 - Royal Victoria Hospital
Friday October 24, 2014 8:00 am – 5:00 pm

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5. Olympus
6. Mettler Toledo
7. EMD Millipore
8. ThermoFisher
9. AB Sciex
10. Fisher/Corning
11. GE Healthcare
12. Promega
13. Leica
14. Bio Basic
15. Pall/VWR
16. Life Technologies
17. Bruker
18. Zeiss
20. Eppendorf
21. MediMabs/Bio-Rad
22. Agilent

Stage

Food & drink

Food & drink

Parking P3 entrance (from Pine Avenue)
Parking: $21/day
Thank You to Our Major Sponsors/Partners
Mission Statement: founded in 2011, Clinical Proteomics (www.clinprot.org) is a joint venture between the Research Institute of the McGill University Health Centre (RI-MUHC) and McGill University. Clinical Proteomics is distributed over two sites, one at the Royal Victoria Hospital (housed in M3) and one at the McGill University and Genome Quebec Innovation Centre (MUGQIC, on the 5th floor). We specialize in discovery-based proteomics using state of the art instrumentation and streamlined bioinformatics coupled with extensive on hand expertise. Internal projects include the elucidation of the molecular pathophysiology of non-alcoholic fatty liver disease using patient and donor-derived liver material as well as proteomics-related R&D projects. Clinical proteomics is also open to both academia and industry on a user-fee basis with highly competitive fees.

What we offer with respect to open access: Advanced mass spectrometry and proteomics technologies for protein and lipid identification and quantitation, as well as affinity enrichment and MS identification of phosphopeptides.
  • Advice and strategy for proteomics research
  • Support in grant applications
  • Experimental design and troubleshooting
  • Advanced proteomic analysis and data interpretation
  • Training

Who we are:
  • Director: Dr. Tommy Nilsson
  • Core Operator: Amy Wong, M.Sc. (maternity leave)
  • Core Operator: Daniel Moses, M.Sc.
  • Bioinformatician: Dr. Alexander Mazur
  • Post doctoral fellow: Dr. Amani Batarseh
  • Operation/Business/Administration and Contact Person: Dr. Jing Liu: jing.liu2@mcgill.ca
What is Systems Biology?
Systems biology describes the scope of inquiry and range of techniques employed to address large-scale research projects. It cuts across disciplines by integrating traditional approaches with new technologies and quantitative analysis. Systems biology allows coordinated research teams to perform cutting edge health research by integrating data from gene to patient.

How is Systems Biology applied to biomedical research?
Systems biology approaches to medical research have proven very powerful. In place of traditional "reductionist" paradigms, systems biology employs a whole-organism approach, employing sophisticated tools and methods to provide more detailed and sophisticated understanding of how genes and proteins interact dynamically in both health and disease. These data are often correlated or verified with clinical data, resulting in a gene-to-patient approach to health. Systems biology approaches are increasingly producing clinical successes for a range of diseases, including cancer, diabetes, cystic fibrosis, and many orphan/neglected diseases.

What is the Systems Biology Training Program?
The Systems Biology Training Program is designed to provide McGill PhD students with a broad understanding of a systems biology approach to health research. The program will provide training and experience in the three components of systems biology, as depicted in the program's logo: life science, quantitative science, and technology development and application. These three components combine to understand and inform clinical observations in health research. Students will identify a supervisor that participates within an interdisciplinary team of researchers at McGill. For Systems Biology, such "teams" or "groups" usually involve one or more basic life science researchers, quantitative scientists (statisticians, computer scientists, mathematicians) and technology developers (individuals who develop or use high-throughput technologies). Ideally the identified team also has some "translational" expertise related to clinical and medical applications. Students will participate in lab rotations "within their team" and collaborate with these researchers from differing specializations, in order to become fluent in the methods and philosophy of systems biology. As such, the Systems Biology training program is light-weight since it allows students to remain in close to contact with their research throughout the various training exercises. By the end of their studies, trainees will be well-positioned to initiate and drive interdisciplinary systems biology research programs, and will be prepared to fully engage in the future of health research.

Who should apply to the Systems Biology Training Program?
The Systems Biology Training Program is designed for McGill PhD students working in biomedical research-related fields, whether in traditional wet-bench disciplines like Biochemistry and Biology, quantitative disciplines like Computer Science and Biostatistics, or technological disciplines like Chemistry and Engineering. Students from various departments will have access to the facilities and technologies at the new McGill Life Sciences Complex and Genome Centre, and at various other locations on campus.

How do I apply to the Systems Biology Training Program?
Students who have recently begun, or are planning to begin, their PhD studies are encouraged to apply. More detailed information is available in our website including information about the program requirements and description, course outlines, and application information.
Dr. Peter Metrakos, MD CM, FACS, FRCSC is a Professor of Surgery, Pathology and Anatomy and Cell Biology at McGill University. At the McGill University Health Centre he is the Director of the Multi-Organ Transplant Program and Director of Hepatopancreatobiliary Surgery. As well, he is the Program Leader of the Cancer Axis for the Research Institute of the MUHC.

Dr. Metrakos graduated in General Surgery from McGill University Faculty of Medicine where he also completed a 3 year Basic Science Fellowship in the Department of Experimental Surgery focusing on the physiology of the pancreas and cellular transplantation. Additional Fellowships were carried out in London, Ontario for liver, kidney and small bowel transplantation and in Minnesota for whole organ pancreas and islet cell transplantation.

Dr. Metrakos runs a high volume hepatopancreatobiliary and transplant surgery service. His research interests include the role and interaction of chemotherapy and liver resection in the management of colorectal cancer liver metastases: The Effect of Neoadjuvant Chemotherapy on Liver Metastases (Journal of Gastrointestinal Surgery 2006), The pathology of liver nodules after neoadjuvant chemotherapy is associated with outcomes for colorectal cancer metastases (HPB 2007), A "New era" In the treatment of colorectal cancer liver metastasis: The gloves are off! (Annals of Surgery. 2009), Perioperative bevacizumab containing chemotherapy and liver resection for colorectal cancer liver metastasis (HPB 2010), Portal Vein Embolization stimulates tumour growth in patients with colorectal cancer liver metastases (HPB Oxford 2012), and Staged Hepatectomy for bilobar colorectal hepatic metastases (HPB Oxford 2012). Other research interests are in HCC, Neuroendocrine tumours and targeted therapies, and include: SHARP Investigators Study Group. Sorafenib in advanced hepatocellular carcinoma (New England Journal of Medicine 2008), Sunitinib Malate for the Treatment of Pancreatic Neuroendocrine Tumors (New England Journal of Medicine 2011) and eIF4E/4E-BP ratio predicts the efficacy of mTOR targeted therapies (Cancer Res. 2012).

Dr. Metrakos also studies the role of ischemia/reperfusion of non-alcoholic fatty liver disease in liver transplantation and liver resections. This is funded by a strategic FRQS systems medicine grant.

Biology is complex and understanding it is a big challenge. Identify and quantify more proteins and complexities such as PTMs faster and more accurately with our new portfolio of LC-MS instruments, sample prep solutions and software. HRAM solutions using Thermo Scientific™ Orbitrap™ MS quantifies all detectable proteins and peptides with high specificity and fewer false positives, while triple quadrupole MS delivers SRM sensitivity and speed to detect targeted proteins more quickly. Join us in meeting today’s challenges. Together we’ll transform proteomics.

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In partnership with McGill University, Génome Québec has built a world renowned research centre offering genomics, proteomics and bioinformatics services to the scientific, academic and industrial communities, both at national and international levels. Inaugurated in 2002, the Innovation Centre is the only one of its kind in Canada, regrouping so many services under one roof. Its science and technology platforms as well as its expertise have enabled the Innovation Centre to position itself as a world leader. Composed of competent professionals, the Centre offers to its users, the needed expertise, guidance and follow up to cover all the critical aspects to ensure successful experiments.

**Sequencing Services**
The Innovation Centre has implemented a wide range of the most recent Next-Generation Sequencing technologies (3 Roche GS-FLX, 15 Illumina HiSeq2000/2500, 1 MiSeq, 1 PacBio and 1 Ion Torrent). The Centre has standardized several key protocols and applications for Whole Genome Sequencing, Exome-Seq, RNA-Seq, ChIP-Seq and Methyl-Seq. The Centre has recently joined the ranks as an Illumina Genome Network Partner (IGN). The Centre continues to offer Sanger DNA sequencing services as well as a DNA extraction service. With its fleet of four ABI DNA Analyzers and a capacity exceeding 1 million high quality reads per year, the Centre can rapidly sequence PCR products, plasmids, BACs and can assist you in the completion of your SNP discovery projects. The Centre also offers a 24-hour sequencing service: BaseXpress.

**Gene Expression Analysis Services**
The Array Technologies Unit of the Innovation Centre offers various functional genomics platforms (Affymetrix, Illumina and Agilent) allowing global transcriptional regulation studies. Personalized services adapted to your project needs are proposed.

**Genotyping Services**
The Innovation Centre operates several platform technologies for the genotyping of small numbers of markers such as SNPs (Single Nucleotide Polymorphisms), Indels (insertion/deletion polymorphisms) and microsatellites, all the way to genome-wide analyses of up to 5 million polymorphisms in parallel. Copy number variation (CNV) and quantitative methylation analyses of genomic DNA are also available.

**Proteomics Services**
The Innovation Centre provides timely and affordable tools for protein fractionation, identification and quantitation. The Unit offers guidance to users, from the experimental design phase all through to the data compilation and interpretation stage.

**Bioinformatics Services**
Over the years, the bioinformatics team developed the web-based application Nanuq and the stand-alone application FlexArray for microarray data analysis and visualization. Many tools and applications pipelines have more recently been developed to facilitate and accelerate the analysis of next-generation sequencing data.
The Chemical Biology Program at McGill

http://www.mcgill.ca/biochemistry/chemicalbiology

The Chemical Biology Graduate Option is centered on the pursuit of an original research project under the direction of one or more Mentors. Students may currently register for the Option through any of the Departments of Biochemistry, Chemistry, Physiology or Pharmacology and Therapeutics. At present roughly 40 graduate students and a smaller number of postdoctoral fellows are enrolled in the program. The program is supported by McGill University and by the Canadian Institutes of Health Research (CIHR) through its Strategic Training Initiatives program.

The program of training incorporates several important features, including a diverse curriculum and programs of seminars, workshops and discussion groups designed to provide students with a well-rounded exposure to both the chemical and biological aspects of the discipline.
AT THE HEART OF 21ST CENTURY LIFE SCIENCES
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<th>Affiliation</th>
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<tbody>
<tr>
<td>Hasilo Craig</td>
<td><a href="mailto:craig.hasilo@mail.mcgill.ca">craig.hasilo@mail.mcgill.ca</a></td>
<td>Experimental Surgery RI-MUHC</td>
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<tr>
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</tr>
<tr>
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<td>Sales, Waters</td>
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