

URMC Proteomics Symposium

Presented by the URMC Mass Spectrometry Resource Laboratory and

Thermo Fisher

Save the date! Wednesday, September 14, 2016

Class of '62 Auditorium, G-9425 10 am – 2:15 pm

Dr. Michael C. Fitzgerald

Department of Chemistry, Duke University "New Mass Spectrometry-Based Methods and Applications for Protein Folding Free Energy Measurements"

Dr. Jun Qu

Department of Pharmaceutical Sciences, SUNY Buffalo "Ionstar: Extensive and reproducible proteomic measurements in large biological cohorts with high accuracy, precision and extremely-low levels of missing data"

Dr. Dragony Fu

Department of Biology, University of Rochester "Proteomic analysis of human RNA modification enzymes"

Dr. Gwillym Williams

Tanz Centre for Research in Neurodegenerative Diseases, University of Toronto "Isobaric label-based proteomics on an Orbitrap Fusion Tribrid instrument reveals elusive role of the cellular prion protein"

Dr. Greg Tombline

Department of Biology, University of Rochester "Proteomics of Longevity"

Kevin Welle

URMC Mass Spectrometry Resource Laboratory, University of Rochester "Mass Spectrometry Resources at the University of Rochester"

Contact: Anne Reed, 273-1359, anne_reed@urmc.rochester.edu

Please RSVP at the following link: https://www.surveymonkey.com/r/Sept14ProteomicsSymp