

| Date | Time | Track | Presentation Title | Speaker |
|--------|------------------------|-------|---|--|
| 12-Sep | 08:05 - 08:10 AM | | Day 1 Introduction | Ryan Bomgarden, Ph.D. Senior R&D Manager, Protein & Cell Analysis, Thermo Fisher Scientific |
| 12-Sep | 08:10 - 09:10 AM | | Adapting TMT and EasyPep Sample Preparation Reagents for High-Throughput, Automated Workflows | Dustin Frost, Ph.D. R&D Scientist, Protein & Cell Analysis, Thermo Fisher Scientific |
| 12-Sep | 08:40 - 09:15 AM | | Derailed Protein Turnover in the Aging Mammalian Brain | Jeffrey Savas Assistant Professor of Neurology, Northwestern University, Feinberg School of Medicine |
| 12-Sep | 09:25 - 09:50 AM | | Impact of Cell Background on Small Compound Target Deconvolution by CETSA Coupled to Mass Spectrometry | Daniel Martinez Molina CSO, Pelago Bioscience Tomas Friman Senior Scientist, R&D Department at Pelago Bioscience AB, Solna, Sweden |
| 12-Sep | 10:25 - 10:25 AM | | Assessing the performance of the Automated Sample Preparation Platform AccelerOme | Alberto Paradela Proteomics Core Facility Fernando Corrales Senior Scientist and Head of the Proteomics Laboratory, Centro Nacional de Biotecnología |
| 12-Sep | 11:15 - 12:30 PM | | Targeted Pathway Proteomics Using TMT and Real-Time Analytics to Create GoDig Assays | Steven Gygi Faculty Director of two MS Core Facilities (Taplin Biological MS Facility and the Thermo Fisher Center for Multiplexed Proteomics) |
| 12-Sep | 11:50 - 12:30 PM | | Roundtable Discussion Day 1 | |
| 13-Sep | 08:05 - 08:10 AM | | Day 2 Introduction | Bhavin Patel, MD, MS Senior R&D Manager, Mass Spectrometry Reagents, Thermo Fisher Scientific |

| 13-Sep | 08:10 - 09:10 AM | Mapping Communication Dynamics Between Physically Interacting Cells Using Hybrid Quantitative Proteomics | Kelly Stecker, PhD Assistant Professor, Utrecht University |
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| 13-Sep | 08:50 - 09:20 AM | An Intelligent Data Acquisition Approach to Increase the Throughput of Multiplexed Proteome Analyses | Chris Rose Director of Discovery Proteomics within the Microchemistry, Proteomics, and Lipidomics Department |
| 13-Sep | 09:50 - 10:25 AM | Multilevel Statistical Modeling for Proteomic Experiments With Complex Designs and Isobaric Labeling | Olga Vitek Professor, Khoury College of Computer Sciences, Northeastern University |
| 13-Sep | 10:45 - 11:15 AM | HyperSCP: Combining Isotopic and Isobaric Labeling for Higher Throughput Single-Cell Proteomics | Ryan Kelly Professor of Chemistry, Brigham Young University |
| 13-Sep | 11:35 - 12:30 PM | Roundtable Discussion Day 2 | |