

Mass Spectrometry Symposium

Wednesday, September 18th, 2024

9:00 AM – 3:00 PM

University of North Carolina at Chapel Hill

Medical Biomolecular Research Building, Room G-202

111 Mason Farm Road, Chapel Hill, NC, 27599

*Paid parking available in the Dogwood Deck or the Ambulatory Care Center Visitor Lot



[Register free here](#)

9:00– 9:30 am	Registration, coffee & networking
9:30 – 9:50 am	Introduction to the UNC Metabolomics & Proteomics (MAP) Core and its expanding capabilities <i>Laura Herring, Director, UNC MAP Core;</i> <i>Whitney Stutts, Associate Director-Metabolomics, UNC MAP Core</i>
9:50 – 10:30 am	Topic: Novel ZipChip ACE™ metabolomics workflow <i>Will Thompson, Principal Scientist, R&D 908 Devices</i>
10:30 – 11:10 am	Topic: Targeted mass spectrometry workflows on Stellar MS <i>Scott Peterman, Product Marketing Manager, Mass Spectrometry, Thermo Fisher Scientific</i>
11:10 – 11:20 am	Break
11:20 – 12:00 pm	Topic: Proteomics applications <i>Lilian Heil, Application Scientist, Thermo Fisher Scientific</i>
12:00 – 1:00 pm	Catered lunch & networking
1:00 – 1:40 pm	Topic: Single cell proteomics and immunopeptidomics applications <i>Fernanda Salvato, Senior Application Scientist, Thermo Fisher Scientific</i>
1:40 – 2:20pm	Topic: A multi-omics approach to understanding resistance to targeted therapies in cancer <i>Clint Stalneck, Assistant Professor, Pharmacology, UNC School of Medicine</i>
2:20 – 3:00 pm	Q&A session & networking

