



Multi-Omics Seminar

Odense, Denmark | October 23 2018

09:00	Coffee and Registration
09:30	Welcome and Introduction – Willy Bjørklund, Thermo Fisher Scientific
09:45	Wide and Deep: New Reagents and Workflows for Multiplexed Quantification of the Phospho Proteome and Plasma Proteome – Segal Ndiaye, Thermo Fisher Scientific
10:15	Integrated Lipidomics and Proteomics Experiments to Study Lipid Biochemistry – Martin Hermansson, Syddansk University
10:45	Coffee Break
11:15	Unleash your Orbitrap™: Innovations in hardware and workflows for deep proteome profiling compounds – Claire Daully, Thermo Fisher Scientific
11:45	Computer-controlled Top-down MS Characterization of Intact Proteins – Pavel Shliaha, Syddansk University
12:15	Lunch
13:15	Improved Metabolome Coverage and Increased Confidence in Unknown Identification through Novel Automated Acquisition Strategy combining Sequential Injections and MSⁿ – Anas Kamleh, Thermo Fisher Scientific
13:45	Signaling Pathways at Your Fingertips – mIP-tMS Assay for the Akt Pathway – Sebastien Gallien, Thermo Fisher Scientific
14:15	Next Generation Software Tools for Compound Identification and Stable Isotope Tracer Analysis – Anas Kamleh, Thermo Fisher Scientific
14:45	Conclusions
15:00	Meeting Ends

Venue:

University of Southern Denmark
 BMB Seminar Room
 Campusvej 55
 5230 Odense