



Multi-Omics Seminar

Lund, Sweden | October 24 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 – Lars Kristensen, Thermo Fisher Scientific
10:15	Proteogenomics – Combining Genotype With Molecular Phenotype in Cancer – Janne Lehtiö, Science for Life Laboratory / Karolinska Institute
10:45	Coffee Break
11:15	Unleash your Orbitrap™: Innovations in hardware and workflows for deep proteome profiling compounds – Claire Dauly, Thermo Fisher Scientific
11:45	Deep Mining the Protein Expression in Melanoma Cancer Patients within the European Cancer Moonshot Center Lund – Gyorgy Marko-Varga, Lund 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:

Lund University
Astronomihuset, Lundmarksalen
Sölvegatan 27
223 63 Lund