



Multi-Omics Seminar

Lisbon, Portugal | October 9 2018

09:15	Coffee and Registration
9:45	Introduction to the Oneida Project – Monica Serrano -
09:50	Wide and Deep: New Reagents and Workflows for Multiplexed Quantification of the Phospho Proteome and Plasma Proteome – Lars Kristensen, Thermo Fisher Scientific
10:15	Multionomics application @UNIMS – Ana Catarina Guerreiro (Mass Spectrometry Unit ITQB-NOVA IBET)
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	Mass Spectrometry as a tool to explore food composition and biomarkers of food intake (Rosário Bronze, ITQB-NOVA/IBET)
12:15	The sweet and sour life of proteins: a tale of (glyco)proteomics at UPorto – Dr Andre Silva, Faculty of Sciences of Porto
12:45	Lunch
13:45	Improved Metabolome Coverage and Increased Confidence in Unknown Identification through Novel Automated Acquisition Strategy combining Sequential Injections and MSⁿ – Marie-Pierre Pavageau, Thermo Fisher Scientific
14:15	Signaling Pathways at Your Fingertips – mIP-tMS Assay for the Akt Pathway – Lars Kristensen, Thermo Fisher Scientific
14:45	Next Generation Software Tools for Compound Identification and Stable Isotope Tracer Analysis – Marie-Pierre Pavageau, Thermo Fisher Scientific
15:15	Conclusions
15:30	Meeting Ends

Venue:

ITQB Nova

Avenida da República, Estação Agronómica

2780-157 Oeiras