



# Multi-Omics Seminar

Heidelberg, Germany | October 16 2018

09:00	<b>Coffee and Registration</b>
09:30	<b>Welcome and Introduction</b> – Susanne Szotek, Thermo Fisher Scientific
09:45	<b>Wide and Deep: New Reagents and Workflows for Multiplexed Quantification of the Phospho Proteome and Plasma Proteome</b> – Rosa Viner, Thermo Fisher Scientific
10:15	<b>Spatial Single-cell Metabolomics by Integrative Microscopy and Imaging Mass Spectrometry</b> – Theodore Alexandrov, EMBL
10:45	<b>Coffee Break</b>
11:15	<b>Unleash your Orbitrap™: Innovations in hardware and workflows for deep proteome profiling compounds</b> – Samy Memmi, Thermo Fisher Scientific
11:45	<b>Exploring Global Protein-RNA Interactions Using XRNAX</b> – Jakob Trendel, EMBL
12:15	<b>Lunch</b>
13:15	<b>Improved Metabolome Coverage and Increased Confidence in Unknown Identification through Novel Automated Acquisition Strategy combining Sequential Injections and MS<sup>n</sup></b> – Anas Kamleh, Thermo Fisher Scientific
13:45	<b>Signaling Pathways at Your Fingertips – mIP-tMS Assay for the Akt Pathway</b> – Sega Ndiaye, Thermo Fisher Scientific
14:15	<b>Next Generation Software Tools for Compound Identification and Stable Isotope Tracer Analysis</b> – Anas Kamleh, Thermo Fisher Scientific
14:45	<b>Conclusions</b>
15:00	<b>Meeting Ends</b>

## Venue:

Best Western Plus Palatin Kongresshotel  
 Ringstrasse 17-19  
 69168 Wiesloch