





Day 1 – October 30

Time PDT	Time EDT	Time GMT	Session
08:00 – 08:50	11:00 – 11:50	15:00 – 15:50	<p>The Human Protein Atlas – Creating a Pan-Disease Resource Based on Next Generation Blood Profiling <i>Mathias Uhlén, PhD</i> KTH Royal Institute of Technology</p> <p style="text-align: right;">Keynote Speaker</p>
08:50 – 08:55	11:50 – 11:55	15:50 – 15:55	Networking Break
08:55 – 09:40	11:55 – 12:40	15:55 – 16:40	<p>Mitigating Sorter-Induced Cellular Stress with Cartridge, Microfluidics-Based Sorting <i>Crystal Montoya</i> Miltenyi Biotec</p> 
09:40 – 09:45	12:40 – 12:45	16:40 – 16:45	Networking Break
09:45 – 10:35	12:45 – 13:35	16:45 – 17:35	<p>Spatial Tissue Proteomics to Assess Health and Disease <i>Fabian Coscia, PhD</i> Max Delbrück Center</p>
10:35 – 10:40	13:35 – 13:40	17:35 – 17:40	Networking Break
10:40 – 11:00	13:40 – 14:00	17:40 – 18:00	<p>The Sulpho-Phospho-Vanillin Assay: Accurate Lipid Normalization in Quantitative Lipidomics <i>Karri Basso, PhD</i> University of Florida</p> 
11:00 – 11:05	14:00 – 14:05	18:00 – 18:05	Networking Break
11:05 – 11:55	14:05 – 14:55	18:05 – 18:55	<p>Proteomics for Cancer Precision Medicine <i>Janne Lehtiö, PhD</i> Karolinska Institutet</p>
11:55 – 12:00	14:55 – 15:00	18:55 – 19:00	Networking Break
12:00 – 12:20	15:00 – 15:20	19:00 – 19:20	<p>How Breakthroughs in Depth, Scale and Speed are Unlocking New Proteomic Discovery <i>Aaron Gajadhar, PhD</i> Seer Bio</p> 
12:20 – 12:25	15:20 – 15:25	19:20 – 19:25	Networking Break
12:25 – 12:55	15:25 – 15:55	19:25 – 19:55	<p>Phantom Metabolites: Unraveling Metabolic Shadows <i>Gary Siuzdak, PhD</i> Scripps Research Institute</p>
12:55	15:55	19:55	End of Day 1

Day 2 – October 31

Time PDT	Time EDT	Time GMT	Session
08:00 – 08:50	11:00 – 11:50	15:00 – 15:50	<p>A Novel Strategy for the Absolute Quantification of Human Proteins <i>Christoph Borchers, PhD</i> McGill University</p> <p style="text-align: right;">Keynote Speaker</p>
08:50 – 09:00	11:50 – 12:00	15:50 – 16:00	Networking Break
09:00 – 09:20	12:00 – 12:20	16:00 – 16:20	<p>From Single Cells to the Metaproteome: Latest Developments in timsTOF Technology for Proteomics <i>Matt Willetts, PhD</i> Bruker</p> 
09:20 – 09:30	12:20 – 12:30	16:20 – 16:30	Networking Break
09:30 – 10:20	12:30 – 13:20	16:30 – 17:20	<p>Single-Cell Chromatin Proteomics to Investigate the Biology of Exceptional Longevity <i>Simone Sidoli, PhD</i> Albert Einstein College of Medicine</p>
10:20 – 10:30	13:20 – 13:30	17:20 – 17:30	Networking Break
10:30 – 10:50	13:30 – 13:50	17:30 – 17:50	<p>Optimizing Metabolomics Workflows: From Sample Preparation to Multi-omics Integration <i>Hagen Magnus Gegner, PhD</i> CTC Analytics</p> 
10:50 – 11:00	13:50 – 14:00	17:50 – 18:00	Networking Break
11:00 – 11:30	14:00 – 14:30	18:00 – 18:30	<p>Analyzing Metabolic Responses to a Standardized Meal Challenge Using Exhaled Breath <i>Matteo Tardelli, PhD</i> Owlstone Medical</p>
11:30	14:30	18:30	Event Close



You may also be interested in our upcoming online event:
“The Spatial Biology Revolution 2024”
 December 4-5, 2024
 Register your free space today – scan or click the code!