



Metabolomic and proteomic applications with latest NMR- and (ultra) high resolution LC-IMS-MS and MRMS technologies

17.4.2019, Christian - Albrechts - Universität zu Kiel, Institut für Humanernährung und Lebensmittelkunde, - Lebensmitteltechnologie – Seminarraum 14 (EG), Heinrich-Hecht-Platz 10, 24118 Kiel

Learn more about recent workflow advances that allow for a better description of the proteome, metabolome and lipidome with Bruker's latest MS and NMR based omics solutions.

Program:

13:00 - 13:10

Welcome

Dr. Stephan Kühne, Bruker Daltonik / Dr. Tobias Demetrowitsch, Kiel Network for Analytical Spectroscopy and Mass Spectrometry, University Kiel

13:10 - 13:50

Overcoming challenges in shotgun proteomics with TIMS and PASEF technology

Dr. Scarlet Koch, Proteomics Business Development Manager, Bruker Daltonik

13.50 – 14:10

Snacks and Drinks

14:10 - 14:50

NMR metabolomics from food quality and authenticity to personalized nutrition

Dr. Manfred Spraul, CTO AIC Division, Bruker Biospin

14:50 - 15:30

MS Fingerprinting and the other half of the characterization Equation

Dr. Christopher Thompson, Global Business Development Manager MRMS, Bruker Daltonics

15:30 - 16:15

Novel TIMS and MRMS based solutions for advancing Metabolomics and large cohort Phenomics research

Dr. Florian Zubeil, Application Scientist Bruker Daltonik

We would be pleased to welcome you! Please register [here](#) or via mail at spectromics@email.uni-kiel.de