30th MassSpec-Forum Vienna

19th to 20th February 2019



University of Vienna, Faculty of Chemistry Währinger Str. 38, 1090 Vienna, Austria



Program Day 1 (Tuesday)

08:45-09:00 Gunda Köllensperger, Christopher Gerner University of Vienna, Vienna, Austria Welcome and Opening Session 1: Proteomics 09:00-09:45 **Sven Degroeve** Ghent University, Ghent, Belgium Ionbot: a novel, fully data-driven search engine for open modification and mutation searches 09:45-10:10 Claire Dauly Thermo Fisher Scientific, Vienna, Austria Comprehensive Proteome Mapping using LC-FAIMS Pro-MS/MS 10:10-10:30 Michael Przybylski Steinbeis Centre for Biopolymer Analysis and Biomedical MS, Rüsselsheim, Germany Molecular Epitope Determination of Aptamer Complexes of the Multi-domain Protein C-Met by Proteolytic Affinity- Mass Spectrometry 10:30-10:45 APMA Young Investigators 10:45-11:25 Coffee Break and Company Exhibition (Second Floor) **Session 2: Glycomics** 11:25-12:10 Manfred Wuhrer Leiden University, Leiden, Netherlands Mass spectrometry glycomics: increasing sensitivity, throughput and precision 12:10-12:30 Friedrich Altmann University of Natural Resources and Life Sciences, Vienna, Austria MALDI-MS of N-glycans from algae reveals nature's sense of humour 12:30-12:50 Katharina Paschinger University of Natural Resources and Life Sciences, Vienna, Austria Mass spectrometric and array analysis of highly modified and immunoactive N-glycans of the canine heartworm 12:50-13:50 Lunch Break and Company Exhibition (Second Floor) Session 3: Biomedical Applications I 13:50-14:35 Denise Wolrab University of Pardubice, Pardubice, Czech Republic UHPSFC/MS lipid profiling for the analysis of biological samples 14:35-15:00 Moritz Wagner Agilent Technologies, Basel, Switzerland Expanding the Lipidomics tool box – from ion mobility to lipid annotation and profiling 15:00-15:15 Tamara Tomin Medical University of Graz, Graz, Austria Altered glutathione redox status in failing hearts revealed by a novel assay based on two-step alkylation with N-ethylmaleimide isotopologues and mass spectrometry 15:15-15:30 Armin Sebastian Guntner Johannes Kepler University, Linz, Austria Development of a highly sensitive GC-MS method preceded by solid-phase microextraction for the analysis of Propofol in low-volume cerebral microdialysate samples 15:30-16:10 Coffee Break and Company Exhibition (Second Floor) **Session 4: Environmental Analysis** 16:10-16:35 **Sven Vedder** Shimadzu Europe GmbH, Duisburg, Germany High resolution mass spectrometry quantitation to help accelerate data review and deliver actionable data 16:35-16:50 Lisa Fischer University of Natural Resources and Life Sciences, Vienna, Austria ICP-MS based methodologies for the trace- and ultra-trace analysis of high mineralized ground waters 16:50-17:05 Elisabeth Varga University of Vienna, Vienna, Austria Deciphering the chemical diversity of prymnesins by MS/HRMS

Targeted analysis of emerging Alternaria toxins in plasma, urine and fecal samples of Sprague Dawley rats

Deoxynivalenol and A431 epidermoid squamous carcinoma cells: proteomic profiling reveals the pathways

connecting ribosomal inhibition with a loss of function of the cell membrane

17:05-17:20 Hannes Puntscher University of Vienna, Vienna, Austria

Informal Get Together/Dinner (Ground Floor)

17:20-17:35 Lukas Janker University of Vienna, Vienna, Austria

Posters & Drinks

17:35

19:00