32nd MassSpec-Forum Vienna

Program Day 1 (Tuesday)

08:45-09:00	Gunda Köllensperger, Christopher Gerner University of Vienna Welcome and Opening
	Session 1: Elemental Analysis
09:00-09:35	,
	Production and Characterisation of an isotopic spike for the analysis of environmental 237Np by (A)MS
09:35-10:10 <i>T2</i>	Johanna Irrgeher Montanuniversität Leoben Expanding the isotopic toolbox in support of the sustainable development goals
	Andreas Limbeck TU Wien
T3	Analysis of trace elements in microplastics – release and uptake during environmental degradation
10:30-10:45	Anna Schöberl University of Vienna
	The Uptake of Metal-Based Anticancer Drugs in Macrophage Subtypes at the Single-Cell Level by LA-ICP-TOFMS Imaging
10:45-11:25	Coffee Break and Company Exhibition (Second Floor)
	Session 2: Metabolomics I
11:25-11:45	André Müller Thermo Fisher
T5	
	Anya Sherman University of Vienna
	Orbitrap Mass Spectrometry to Elucidate Plant Uptake and Metabolism of Tire Leachate Compounds
12:00-12:15	
	A systematic study of lipids – from ESI analysis to MALDI Imaging using a 7T FT-ICR instrument
	Sarah Fleissner TU Wien
<i>T8</i> 12:30-13:30	Characterization of selected organometallic compounds monitored for catalytic activity Lunch Break and Company Exhibition (Second Floor)
12.30-13.30	Editor Break and Company Exhibition (Second Floor)
	Session 3: Lipidomics
13:30-14:05	Dominik Schwudke Forschungszentrum Borstel
<i>T9</i>	Tuberculostearic acid (TSA)-containing phosphatidylinositols as molecular tool to study Tuberculosis
14:05-14:25	
T10	3 / 3
14:25-14:45	Robert Ahrends University of Vienna
<i>T11</i> 14:45-15:00	Probing the Membrane Landscape and Identifying Key Lipids Critical for Platelet Activation by Lipidomics Ernst Pittenauer TU Wien
	High-energy CID of diether phospholipids: parallels to diester phospholipids and undescribed new fragmentation pathways
15:00-15:40	Coffee Break and Company Exhibition (Second Floor)
13.00 13.10	
	Session 4: Honory Session for Günter Allmaier
15:40-15:50	•
15:50-16:25	
	From Plasma Desorption to blood plasma analysis - a personal proteomics journey
16:25-17:00	Karoly Vekey Research Centre for Natural Sciences, Hungarian Academy of Sciences
T14	Importance of CID energy dependence
17:00-17:35	Jasna Peter-Katalinic Westfälische Wilhelms-Universität Münster
T15	Guenter Allmaier: Contributions to molecular characterization of glycoconjugates by mass spectrometry
17:35-17:55	Marcella Frauscher AC2T research GmbH
T16	
17:55	Poster Session
19:00	Informal Get Together/Dinner

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Program Day 2 (Wednesday)

	Session 5: Metabolomics II
09:00-09:35	Noé Karl Brasier ETH Zürich, University Hospital Basel
T17	Translating clinical sweat analysis – A promising outlook with remaining challenges
09:35-09:55	Marcel Kwiatkowski University of Innsbruck
T18	Combined Metabolic and chemical (cometchem) labeling - a strategy to quantify site-specific reaction rates []
09:55-10:15	J. Thomas Hannich CeMM Research Center for Molecular Medicine of the ÖAW
T19	Targeted Metabolomics Pipeline to Deorphanize Solute Carrier Transporters (SLCs)
10:15-10:55	Coffee Break and Company Exhibition (Second Floor)
	Session 6: MS Fundamentals
10:55-11:00	Laudatio Pregl-Award
11:00-11:35	Gérard Hopfgartner Université de Genève
T20	
11:35-11:55	
T21	Cyclic IMS, a new tool for R&D
11:55-12:15	Younes Valadbeigi University of Natural Resources and Life Sciences
T22	Ionization Mechanism in an Atmospheric Pressure Chemical Ionization Ion Source
12:15-12:35	Dietmar Hammerschmid King's College London
T23	Targeting protein dynamics in membranes: Novel avenues for hydrogen/deuterium exchange mass spectrometry
12:35-13:35	Lunch Break and Company Exhibition (Second Floor)
	Session 7: Data Interpretation
13:35-14:10	Tomáš Pluskal IOCB Prague
T24	MZmine 3 - a tool from and for the mass spectrometry community
14:10-14:30	Jan Bucek Plasmion
T25	SICRIT®- New Dimensions in Mass Spectrometry
14:30-14:50	Samuel Meier-Menches University of Vienna
T26	Causal inference to deconvolute metabolomics data from human plasma
14:50-15:05	Mathias Gotsmy University of Vienna
T27	Probabilistic quotient's work &pharmacokinetics' contribution: countering size effect in metabolic time series measurements
15:05-15:45	Coffee Break and Company Exhibition (Second Floor)
	Session 8: Proteomics
15:45-16:20	Manuel Matzinger IMP - Research Institute of Molecular Pathology
T28	Reproducible, sensitive and automated one-pot sample preparation for high throughput single cell proteomics
16:20-16:40	Christoph Krisp Bruker Daltonics
T29	Diversifying the proteomics toolbox []
16:40-17:00	Djuro Josić Brown University Providence RI, USA University of Pula
T30	Changes in the proteome of extracellular vesicles shed by rat liver after sub-toxic exposure to acetaminophen
17:00-17:20	Michael Przybylski AffyMSLifeChem Centre for Biochemistry & Biomedical Mass Spectrometry
T31	Molecular determination of survival motor neuron protein (SMN) in Spinal Muscular Atrophy by affinity- mass spectrometry
17:20	Awards & Closing Remarks