

CelebrateMS

Celebrating 50 Years of Mass Spectrometry Research at the University of Southern Denmark

Advances in Biological Mass Spectrometry

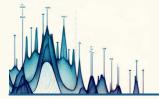
Program	DAY 1 - Tuesday, December 3, 2024	
08:30-09:30 Registration and coffee		
09:30	Welcome and introduction, Ole N. Jensen & Adelina Rogowska-Wrzesinska	
Session 1	Chair: Martin R. Larsen	
10:00	Ruedi Aebersold, ETH Zürich, Switzerland	
10.00	Navigating the proteomic landscape with a mass spectrometer.	
10:40	Amanda Campbell, Odense Univ. Hospital, Univ. of Southern Denmark A carrier-based quantitative proteomics method applied to biomarker discovery in pericardial fluid.	
10:55	Luke Gamon , Univ. Copenhagen Spatial proteomics of the human atherosclerotic microenvironment.	
11:10	Ileana Cristea, Princeton Univ. USA The Virus Microenvironment.	
11:50	Group Photo	
12:00	Lunch, poster session, exhibition	
Session 2	Chair: Thomas J.D. Jørgensen	
13:30	Neil Kelleher , Northwestern Univ., USA The case for levelling up compositional, spatial and single-cell biology.	
14:10	Cagla Sahin , Univ. of Southern Denmark Native and ion mobility MS of intrinsically disordered proteins and their complexes.	
14:25	Refreshments, poster session, exhibition	
Session 3	Chairs: Christer Stenby Ejsing	
15:00	Andrea Sinz, Univ. Halle-Wittenberg, Germany Cross-Linking Mass Spectrometry for Investigating Protein Conformations and Protein-Protein Interactions - A Method for All Seasons	
15:40	Anton Berg Hansen, Univ. Copenhagen Development of a thiol-ene microfluidic chip for sub-zero temperature hydrogen/deuterium exchange mass spectrometry (HDX-MS)	
15:55	Helle Bogetofte-Barnkob , Univ. of Southern Denmark Pathways to Parkinson's Disease: using Proteomics/PTMomics to understand disease development in patient iPSC-derived neurons	
16:10	Morten Thaysen-Andersen, Macquarie Univ, Australia MS unlocks the glycocode of the tumour microenvironment	
16:50	Closing remarks, End of Day 1	
17:15	Transport to Odense City/Odeon by tram ODEON, Odeons Kvarter 1, 5000 Odense C	
18:00-18:30 Symposium reception		



18:30-01:00 Symposium Dinner and Party







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Program	DAY 2 - Wednesday, December 4, 2024
Session 4	Chair: Adelina Rogowska-Wrzesinska
09:00	Judith Steen, Harvard Med. School, USA Unraveling the Mysteries of Misfolded Proteins by Mass spectrometry
09:40	Markku Varjosalo, Univ. Helsinki, Finland A proteocistromic atlas of human transcription factors
09:55	Coffee/Tea, posters, exhibition
Session 5	Chair: Finn Kirpekar
10:25	Klaus Dreisewerd, Univ. Münster, Germany Almost Forty - A fruitful Münster-Odense MALDI connection
11:05	Sudip Ghosh , Lund University, Sweden Delineation of antigen presentation modulation for tailored treatments of infant leukemia
11:20	Perdita Barran , Univ. Manchester, UK Adventures with Mass Spectrometry and Joy - Towards Development of a Quantitative MRM assay using Sebum to Diagnose Parkinson's Disease
12:00	Lunch, poster session, exhibition
Awards	Chairs: Nils J. Færgeman & Cagla Sahin
12:45	Poster awards
Session 6	Chair: Veit Schwämmle
12:55	Lennart Martens , Univ. Ghent, Belgium A recipe for novel discoveries: thousands of pinches of protein, with a spoonful of silicon
13:35	Ulises Guzman , Univ. Copenhagen Proteome-wide measurement of peptide Collisional Cross-Sections in an Orbitrap Mass Analyzer without using ion mobility separations
13:50	Philip Kaulich, Univ. Kiel, Germany Truncated Proteoforms in Top-Down Proteomics Studies
14:05	Bernhard Küster, Techn. Univ. München, Germany ProteomeTools, Understanding Cancer and Repurposing Drugs - All by Mass Spectrometry!
14:45	Closing remarks Ole N. Jensen & Adelina Rogowska-Wrzesinska
15:00	End of conference

15:00-18:00 Tour of MS labs, networking, refreshments

18:00 End of Day 2





