

Day 1			
10:30	Bus from Pasila station		
10:50	Bus from Leppävaara		
11:30	Registration Opens		
12:00	Coffee, sandwiches & exhibition		
12:30	Welcome words (10min)		
Session 1 chair: Dorota Muth-Pawlak			
12:40	Sajjalisa Kangasjärvi	University of Helsinki	Proteomics in plant stress research
13:20	Laura Wey	University of Turku	Omics approaches to identify the molecular players in the pathway of electricity generation from cyanobacteria
13:40	Jukka Kallijärvi	University of Helsinki	O-GlcNAc, an abundant and evolutionarily conserved temperature-dependent protein modification of unknown function
14:00	Jo Kirkpatrick	Thermo Fisher Scientific	TBA
14:20	Coffee break & exhibition		
Session 2 chair: Tuula Nyman			
14:50	Kirsi Savijoki	University of Helsinki	Revealing Hidden Mechanisms of Mycobacterial Infection Using Proteomics
15:30	Rajaram Venkatesan	Biocenter Oulu	Structural characterization of native Mce1 complex from Mycobacterium tuberculosis
15:50	Anam Hammid	University of Eastern Finland	Parallel Reaction Monitoring-Parallel Accumulation-Serial Fragmentation (prm-PASEF) for targeted proteomics analysis and, a correlation between prm-PASEF and DIA-PASEF
16:10	Ross Chawner	Waters Corporation	Combining sensitivity, selectivity, speed and resolution: Utilization of ToF-MRM for the quantification of peptide biomarkers in human glioblastoma
16:30	FinnProt AGM (1h)		
17:30	Posters, exhibition & refreshments		
18:00	Dinner		
18:30	Dinner		
19:00	Dinner		
22:00	Dinner		
Day 2			
Session 3 chair: Markku Varjosalo			
9:00	Lotta Happonen	Lund University	Decoding complement evasion: multimodal structural proteomics as a tool to study host-pathogen interactions
9:40	Oluwatosin Abe	Turku Bioscience Centre	Tumor cell invasion and therapy resistance in head and neck squamous cell carcinoma (HNSCC)
10:00	Dani Flinkman	Turku Bioscience Centre	MINING PARKINSON'S BLOOD CELL PROTEOMES REVEALS ENDOTYPE AND MECHANISTIC INFORMATION
10:20	Ifrat Jahan Tamanna	University of Turku	Decoding emerging microbial lipids and proteins in diet-liver-gut crosstalks
10:40	Tom Ruane	Sciex	TBA
11:00	Coffee break		
Session 4 chair: Otto Kauko			
11:30	Mohieddin Jafari	University of Helsinki	Preventing Proteomics Data Tombs Through Collective Responsibility and Community Engagement
11:50	Balázs Bálint	Turku Bioscience Centre	glaDIAtor-nf: a complete data analysis workflow for quantitative DIA mass spectrometry using nextflow
12:10	Juho Salonen	Agilent Technologies	Elevating Protein Sample Prep: Precision Meets Flexibility
12:30	Lunch		
Session 5 chair: Karoliina Hirvonen			
13:30	Otto Kauko	Turku Proteomics Facility	Advanced mass spectrometry -based proteomics services provided by Turku Proteomics Facility
13:45	Markku Varjosalo	Helsinki Proteomics Center	Helsinki Proteomics Center: State-of-the-Art Mass Spectrometry Infrastructure Enabling Deep, Sensitive, and Accurate Proteomics for Finnish and International Life Sciences Research

14:00	Lackson Kashobwe	Mass Spectrometry Facility Tampere	Advancing Research with Liquid Chromatography Mass Spectrometry (LC-MS/MS): Establishing a State-of-the-Art Facility at Tampere University supported by the Research Council of Finland (FIRI 2024 – 2026)
14:15	Ulrich Bergmann	Biocenter Oulu	Proteome and protein analysis in the core facility of the Biocenter Oulu
14:30	Award ceremony & Closing words		
14:45	Farewell drink		
15:00	Bus to Leppävaara & Pasila		
15:30	Bus in Leppävaara		
16:00	Bus in Pasila		

