SEPTEMBER 16-19 2024 SMAP SEPTEMBER 16-19 2024 LILLE / FRANCE

CONGRESS OF MASS SPECTROMETRY AND PROTEOMIC ANALYSIS

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Artificial Intelligence and Bioinformatics

- Spatial OMICs and MS Imaging
- Top-Down and Structural analysis
- Proteogenomics and metaproteomics
- New topics in OMICs
- Plant and Food OMICs
- Lipidomics, Metabolomics, and Small molecules
- Quantification
- Ion mobility spectrometry
- Instrumental development
- Environmental analysis
- Complex mixtures polymers microplastics
- Forensic and cultural heritage
- Health and biological sciences
- PTM (Post-Translational Modifications)
- Interactomic

Abstract submissions Close: 14 June 2024



Registrations End Early bird: 12 July 2024

Alexander Makarov,

Frontiers of high-resolution accurate mass





Karl Mechtler, 5300 protein per single-cell

Leila Birolo, Challenge of identification of ancient proteins



Sarah Cianferani,

My journey into structural mass spectrometry methods





Maarten Dhaenens, The metabolic projection hypothesis

Sebastian Böcker

Turning tandem mass spectra into metabolite structure





t our plenary Speakers















