IMSC 2024 PROGRAM

Saturday 17 August								Location
8.00 AM - 5.00 PM	Short Course Registration							Level 1 Foyer
Short Courses	**Short Course 1**	**Short Course 2**	**Short Course 3**	**Short Course 4**	**Short Course 5**	**Short Course 6**	**Short Course 7**	**Short Course 8**
9:00am - 5:00pm	**LC-MS Practical Method Development and Validation** Robert Trengrove - Curtin University Damien Callahan - Deakin University	**Tandem mass spectrometry** Vicki Wysocki - Ohio State University Lijilana Pasa-Tolic - Pacific Northwest National Laboratory	**Ion-mobility mass spectrometry** Erin Baker - University of North Carolina Valerie Gabelica - University of Geneva	**Proteomics and Post-Translational Modifications** Sean Humphrey - Murdoch Children's Research Institute Shabaz Mohammed - Rosalind Franklin Institute University of Oxford	**Lipidomics** Stephen Blanksby - Queensland University of Technology Anne Bendt - National University of Singapore	**Imaging mass spectrometry** Martina Marchetti Deschmann - Vienna University of Technology Shane Ellis - University of Wollongong	**Glycomics** Yehia Mechref - Texas Tech University Daniel Kolarich - Griffith University	**Computational and Statistical proteomic analysis** David Tabb - US National Cancer Institute Birgit Schilling - University of California San Francisco (UCSF) Jordan Burton - Buck Institute for Research on Aging
Short Courses to include Morning Tea 10:30am - 11:00am Lunch 12:30pm - 1:30pm Afternoon Tea 3:00pm - 3:30pm								

Sunday 18 August								Location
8.00 AM - 9.00 PM	Conference Registration							Main Foyer 3
Short Courses	**Short Course 1**	**Short Course 2**	**Short Course 3**	**Short Course 4**	**Short Course 5**	**Short Course 6**	**Short Course 7**	**Short Course 8**
9:00am - 5:00pm	**LC-MS Practical Method Development and Validation** Robert Trengrove - Curtin University Damien Callahan - Deakin University	**Tandem mass spectrometry** Vicki Wysocki - Ohio State University Lijilana Pasa-Tolic - Pacific Northwest National Laboratory	**Ion-mobility mass spectrometry** Erin Baker - University of North Carolina Valerie Gabelica - University of Geneva	**Proteomics and Post-Translational Modifications** Sean Humphrey - Murdoch Children's Research Institute Shabaz Mohammed - Rosalind Franklin Institute University of Oxford	**Lipidomics** Stephen Blanksby - Queensland University of Technology Anne Bendt - National University of Singapore	**Imaging mass spectrometry** Martina Marchetti Deschmann - Vienna University of Technology Shane Ellis - University of Wollongong	**Glycomics** Yehia Mechref - Texas Tech University Daniel Kolarich - Griffith University	**Computational and Statistical proteomic analysis** David Tabb - US National Cancer Institute Birgit Schilling - University of California San Francisco (UCSF) Jordan Burton - Buck Institute for Research on Aging
Short Courses to include Morning Tea 10:30am - 11:00am Lunch 12:00pm - 1:00pm Afternoon Tea 3:00pm - 3:30pm								
4.30 PM - 5.00 PM	**Opening Ceremony**							Plenary
5.00 PM - 6.00 PM	**IMSC Plenary Lecture 1** **Membrane protein complexes – from recombinant complexes to regions of the brain** - Professor Dame Carol Robinson							
6.00 PM - 9.00 PM	**Welcome Reception**							Exhibition Hall

IMSC 2024 PROGRAM

	Monday 19 August				Location
8.00 AM - 5.00 PM	Conference Registration				Main Foyer 3
8.00 AM - 9.30 AM	**Plenary - IMSF Thomas Medal** *Chair - Ron Heeren*				Plenary
9.30 AM - 10.00 AM	**Morning Tea** ***Innovation Stage Talks:*** Sciex 9:40am - 9:55am				Exhibition Hall
Location	**Plenary**	**Meeting Room 105**	**Meeting Room 106**	**Meeting Room 109**	**Meeting Room 110**
Concurrent Sessions	**Lipidomics: Applications** ***Session Sponsored by Avanti Polar Lipids***	**Clinical Chemistry**	**Mass Spectrometry in Structural Biology A**	**Earth, Space, Geoscience and Atmospheric Chemistry A**	**Ion Chemistry, Reactions and Structure A**
10.00 AM - 12.00 PM	**KEYNOTE: Yu Xia - Tsinghua University** **Michal Holčapek** (347 - CLIG interlaboratory study on the harmonization of lipid concentrations in human plasma) **Anthony Don** (396 - Quantifying brain lipid synthesis and turnover through deuterium labelling of endogenous brain lipids in vivo) **Amy Liang** (452 - Automated retention time calibration for complex, targeted reverse phase chromatography based lipidomics) **Stephanie Cologna** (168 - Probing fatty acid alterations linked to cholesterol dysregulation in Niemann-Pick Type C Disease) **Rachel Pryce** (176 - Lipidomic Alterations in the Retina of a mouse model of Zellweger Spectrum Disorder Investigated by Mass Spectrometry Imaging) *Chair* - Anne Bendt	**KEYNOTE: Ronda Greaves - Murdoch Childrens Research Institute** **Stefani Thomas** (590 - Adopting fundamental principles from the clinical laboratory to accelerate the clinical translation of targeted mass spectrometry-based proteomic assays) **Priscilla Yeung** (92 - Clonality Determination by Detecting Unmodified Monoclonal Serum Free Light Chains Using On-Probe Extraction Coupled with Liquid Chromatography-High-Resolution Mass Spectrometry) **Keziah Liebenberg** (569 - Alterations in glutaminolysis detected by direct mass spectrometry techniques enable diagnosis and molecular subtyping of breast and ovarian cancers) **Dan Lane** (87 - The Validation Processor: the development of a novel tool that that automates, standardises, and accelerates mass spectrometric assay validation) **Ruben Luo** (12 - Microprobe-Capture In-Emitter Elution Coupled with Mass Spectrometry for Structural Elucidation and Clinical Testing of β2-Transferrin) *Chair - Russell Grant*	**KEYNOTE: Michal Sharon Weizmann - Institute Of Science** **Cameron Fairweather** (426 RAMP it up! Exploring conformational dynamics of the amylin receptors using HDX-MS) **Tara Pukala** (597 - Extending the molecular view of snake venoms to higher order structure) **Weijing Liu** (266 - Automated molecular glues screening using native mass spectrometry) **Duong Bui** (50 - Deciphering Mechanisms and Thermodynamics of Protein Assembly using native mass spectrometry) **Ryan Julian** (76 - Isomerization of tau provides mechanistic insight into the underlying causes of Alzheimer's disease) *Chair - Meng-Qiu Dong*	**KEYNOTE: Roger Summons - Massachusetts Institute Of Technology** **Hyeon-Woo Lee** (56 -Study on the Determination Methods of the Natural Radionuclides(U-238, Th-232) in Building Materials and Consumer Products Using ED-XRF and ICP-MS) **Samir Damare** (500) -Mass spectrometry as a tool for understanding biological processes in Oceans **Clemens Walther** (589 - Nuclear forensics on single micrometer sized particles: recent developments of secondary neutral mass spectrometry for ultra-trace isotope analysis) **Oliver Jones** (447 - Combining contaminants of emerging concern with environmental isotopes to distinguish wastewater and agricultural impacts on groundwater systems) **Robert Kirkby** (634 - Using an automated soil incubation system coupled to online IRMS to resolve N2 and N2O emission pathways from agricultural soils) *Chair - Trevor Ireland*	**KEYNOTE: Evan Bieske - University of Melbourne** **Minsu Kim** (491 - Research on size dependent molecular behavior of ESI-generated charged droplets by using X-ray scattering) **Xianglei Kong** (594 - Generation and Identification of Free Radical Cations Using a UV/IR Double-Beam Laser System Combined with an FT ICR Mass Spectrometer) **Alexander Giuliani** (643 - UV action spectroscopy of polyalanine ions) **Ri Wu** (326 - Probe the Gas-Phase Structures of Biomolecular Ions with Native Ion Mobility-Mass Spectrometry and Transition Metal Ion Förster Resonance Energy Transfer) **Adam Trevitt** (667 - Laser photodissociation and ion reactivity of selected protonation-site isomers) *Chair - Isabelle Compagnon*

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12.00 PM - 1.00 PM	**Lunch (no catering)** *Sponsored Lunch Seminars:*				**Exhibition Hall**
Sponsored Lunch Seminars	**Meeting Room 105**		**Meeting Room 106**	**Meeting Room 109**	
1.00 PM - 2.00 PM	Lunch Seminar **- Bruker**		Lunch Seminar - **Waters**	Lunch Seminar - **Syft**	
1:00 PM - 3:00 PM	**Innovation Stage Talks-** Thermofisher 1:30pm - 1:45pm Shimadzu 1:45pm - 2:00pm Waters 2:00pm -2:15pm Tecan 2:15pm - 2:30pm		**Poster Session** - Monday Posters		Exhibition Hall
2:30 PM - 3:00 PM	**Afternoon Tea**				**Exhibition Hall**

Location	**Plenary**	**Meeting Room 105**	**Meeting Room 106**	**Meeting Room 109**	**Meeting Room 110**
Concurrent Sessions	**Metabolomics A**	**Proteomics: Top-Down**	**Food, Nutrition and Agriculture A**	**Forensics, Sports Doping, Homeland Security A**	**Ion Activation Methods**
3.00 PM - 5.00 PM	**KEYNOTE: Anne Bendt - National University of Singapore** **Breanna Dixon** (48 - Uncovering the resistance phenotype of carbapenemase-producing Enterobacteriaceae (CPE) using metabolomics) **Prabhu Rangabashyam** (438 - Elucidating novel fatty acid structures in vernix caseosa by combining ultraviolet photodissociation and ozone-induced dissociation mass spectrometry) **Islam Sk Ramiz** (205 - Identification of novel glucose-independent, and reversible metabolic pathways associated with anti-proliferative effect of metformin in HepG2 cells) **Agustinus Thomas Soerianto** (445 - Utilizing high resolution mass spectrometry and cheminformatic approaches to delineate the full metabolic capability of the parasitic protist Leishmania mexicana) **Patrik Spanel** (86 - Gas Phase Reactions of O-, OH-, O2- and NO2-with Volatile Fatty Acids for Quantitative SIFT-MS Breath Analyses) *Chair - Nicola Gray*	**KEYNOTE: Ying Ge - University of Wisconsin** **Jack Bennett** (559 - Uncovering hidden protein modifications with native top-down mass spectrometry) **Athanasios Smyrnakis** (624 - A new adaptation of the Omnitrap platform integrated into a trapped ion mobility time-of-flight mass spectrometer) **Tatiana Samgina** (144 - EThcD method as a unique tool for top-down de novo sequencing of intact amphibian skin peptides) **Christian Neusüß** (52 - Characterization of proteoforms of intact proteins by 2-dimensional CE-MS techniques) **Muhammad Zenaidee** (369 - Ion mobility curated internal fragments enhance on-line top-down proteomics experiments) *Chair - Helen Cooper*	**KEYNOTE: Melanie Downs - University of Nebraska-Lincoln** **Genc Heljiti** (61 - The Proteomes that Feed the World: Unveiling Plant Proteomics and Peptidomics) **Michelle Colgrave** (606 - Proteomics application to deliver healthy and sustainable foods of the future) **Peter Hoffmann** (495 - Diving Deep into the Faba Bean Proteome) **Larissa Buedenbender** (512 - chelOMICS for a holistic understanding of siderophore-mediated host-pathogen interactions in aquaculture infections) **Omar Mendoza-Porras** (453 - Understanding Thermal Stress in Salmon Tissues Using Proteomics and Metabolomics) *Chair - Subhra Chakraborty*	**KEYNOTE: Chris Gill - Vancouver Island University** **Maiken Ueland** (122 - When disaster strikes, locating victims using electronic nose technology) **Jeff Zonderman** (387 - Rapid Chromatography-Free Quantitative Screening of Synthetic Cannabinoids in Urine Using a Novel DART-TripleQuad MS) **Louise O'Grady** (25 - Identifying Psychedelics in Australian Acacia Species: Wattle We Do?) **Simon Ovenden** (392 - The identification of VX chemical attribution signatures from four different synthetic methods) **Lisa Scharrenbroch** (95 - High-Resolution and Isotope Ratio Mass Spectrometry based profiling of Ricinus communis - A forensic approach) *Chair - Nick Manicke*	**KEYNOTE: Shabaz Mohammad - Rosalind Franklin Institute University of Oxford** **Catherine Costello** (671 - ExD fragmentation and multistage MSn facilitate high-confidence characterization of intact glycopeptides and glycolipids) **Rachel Ogorzalek Loo** (246 - The Big Break Up—Understanding How a Subunit or Polypeptide Abandons its Noncovalent Complex in Native Top-Down MS) **Li Ding** (169 - Implementation of Electron Capture Dissociation in an RF Linear Ion Trap without Assistance of Magnetic Field) **Oliver Hale** (351 - Declustering protein complexes by infrared photoactivation for in situ native mass spectrometry) **Takahashi Hidenori** (434 - Ion Fragmentation for Detailed Lipid Structural Analysis using Atomic Hydrogen/Oxygen Irradiation (HAD/OAD)) *Chair - Hahn Bin Oh*

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Evening Workshops *Light Catering provided for evening workshop attendees from 5:00pm - 5:30pm*				
5.30 PM - 7.30 PM	**IMSF Focus Group: MS Imaging** Organisers: **Martina Marchetti-Deschmann** (Vienna University of Technology) and **Samuele Zorrato** (Medical University of Vienna)	**FeMS Workshop** *Empowering Women Mass Spectrometrists in a Traditionally Male-Dominated Workplace* Organisers: **Denise Tran** (University of Sydney)	**IMSC Spectroscopy-MS Workshop** *Uniting Mass spectrometry and laser spectroscopy* Organisers: **Caroline Dessent** (University of York) and Sarah Wilson (University of York)	
7.00 PM - 11.00 PM	**Affiliates Dinner - *Invitation only event***			Old Melbourne Gaol

	Tuesday 20 August				**Location**
8.00 AM - 5:00 PM	Conference Registration				Main Foyer 3
8.30 AM - 9.30 AM	**Plenary -** **Professor Jana Roithova - Radboud University Netherlands** *Chair - W. Alex Donald*				Plenary
9.30 AM - 10.00 AM	**Morning Tea**				**Exhibition Hall**
Location	**Plenary**	**Meeting Room 105**	**Meeting Room 106**	**Meeting Room 109**	**Meeting Room 110**
Concurrent Sessions	**Clinical 'omics**	**Mass Spectrometry in Structural Biology B**	**Data Science in Mass Spectrometry A**	**Ionization Methods**	**Environmental Mass Spectrometry A**
10.00 AM - 12.00 PM	**KEYNOTE: Thomas Kislinger - University of Toronto** **Nicola Gray** (496 - Signatures of future bronchiectasis risk in children with cystic fibrosis via lipid profiling of bronchoalveolar lavage fluid) **Tereza Kacerova** (231 - Unlocking Cognitive Impairment: Integrative Analysis Across Multiple Metabolomics and Lipidomics Platforms Reveals Promising Biomarkers for Diagnosis and Prognosis) **Anna Emilia Hoffman** (71 - Interplay between sphingolipid metabolism and the endocannabinoid system in schizophrenia: Insights into biomarker candidacy) **Thomas Meikle** (455 - Development and clinical translation of high-throughput lipidomic profiling for the assessment of individual cardiometabolic risk) **Elizabeth Want** (531 - Exploring novel metabolic changes in tissue injury: possible new treatment routes) *Chair - Stefani Thomas*	**KEYNOTE: Lisa Jones - University of California San Diego** **Adalet Memetimin** (582 - Improved identification of cross-linked peptide pairs by focused accurate inclusion mass screening (fAlMS)) **T. W. Dominic Chan** (203 - Millimeter Water-in-Oil Droplet as an Alternative Back Exchange Prevention Strategy for Hydrogen/Deuterium Exchange Mass Spectrometry of Peptide/Protein) **Joshua Sharp** (28 - Radical Protein Footprinting in Stabilized Whole Blood) **Kuang-Ting Kuo** (275 - Integration of Structural Proteomics and Computational Simulation for Elucidating Structural-Activity Relationships of Novel PPARγ Inverse Agonists in Metabolic Disease) **Terese Eisgruber** (233 - Investigating Modification-Specific Interactions of Linker Histone H1 by Mass Spectrometry-based Proteomics) *Chair - Michal Sharon*	**KEYNOTE: Paul Pigrim - La Trobe University** **Alexander Aksenov** (55 - Illuminating the Dark Matter of Metabolomics Through Molecular Community Networking) **Mengbo Li** (27 - Missingness-informed protein quantification and differential expression analysis) **Yuji Sekiguchi** (290- Genomically predicted protein mass database (GPMsDB) for rapid and broad-spectrum identification of bacterial and archaeal isolates by mass spectrometry) **Piotr Radziński** (245 - Contrastive learning encoding algorithm of MS images for memory management and segmentation enhancement) **Toan Phung** (349 - CHRONICLE, A Family of Tools and Softwares for Mass Spectrometry-based Proteomics Data Processing, Analysis, Visualization, and Exploration) *Chair - Mark Condina*	**KEYNOTE: Nathalie Agar - Brigham and Women's Hospital** **Lea Ledoux** (355 - Bacterioscore in 2D/3D MS imaging: pioneering in vivo cancer microbiome study and its link with diagnosis and patient survival) **Yoichi Otsuka** (34 - Mass spectrometry imaging of a single HeLa cell by tapping-mode scanning probe electrospray ionization) **Jae-Chul Pyun** (123 - Laser desorption/ionization (LDI) mass spectrometry based on nanomaterials for biomedical applications) **Garry Corthals** (572 - Electroless ionisation mass spectrometry (ELI-MS): a simple, speedy and soft ionisation method) **David Borts** (91 - A Coated Blade Spray Mass Spectrometry Workflow for Rapid Toxicology General Unknown Screening) *Chair - Yu Bai*	**KEYNOTE: Kevin Thomas - University of Queensland** **Alicia Macan Schonleben** (236 - Occurrence and Patterns of Emerging Organic Chemicals in Vegan and Vegetarian Products) **Michaela Lerch** (282 - Mass Spectrometric Profiling of Antarctic Sea Water for Expedited Chemical Regulation) **Elliot Price** (556 - Functional analysis of chemical exposure and health by applying molecular hallmark principles) **Yik-Sze Lau** (412 - The Coupling of a High-efficiency Aerosol Collector with Electrospray Ionisation/Orbitrap Mass Spectrometry for the Real-time Chemical Characterisation of Aerosol Particles) **Quan Cheng** (242 - Improving Cytotoxicity Study with Single Cell Lipid Profiling of Microalgae and Bacterial Cells by Microchip-MALDI-MS) *Chair - Emma Schymanski*

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12.00 PM - 1.00 PM	**Lunch (no catering)** ***Sponsored Lunch Seminars:***				**Exhibition Hall**
Sponsored Lunch Seminars	**Meeting Room 105**		**Meeting Room 106**		**Meeting Room 109**
1.00 PM - 2.00 PM	Lunch Seminar - **Bruker**		Lunch Seminar - **Waters**		Lunch Seminar - **Sciex**
1:00 PM - 3:00 PM	**Innovation Stage Talks:** IonOptics 1:15pm - 1:30pm Thermofisher 1:30pm - 14:45pm Agilent 1:45pm - 2:00pm Waters 2:00pm - 2:15pm		Poster Session **- Tuesday Posters**		Exhibition Hall
2:30 PM - 3:00 PM	**Afternoon Tea**				**Exhibition Hall**
Location	**Plenary**	**Meeting Room 105**	**Meeting Room 106**	**Meeting Room 109**	**Meeting Room 110**
Concurrent Sessions	**Metabolomics B**	**Proteomics: Method Development**	**Food, Nutrition and Agriculture B**	**Ion Mobility A**	**Ion Chemistry, Reactions and Structure B**
3.00 PM - 5.00 PM	**KEYNOTE: David Wishart - University of Alberta** **Rafael Montenegro Burke** (509 - The Human Metabolome Atlas unveils metabolic heterogeneity across cell types and stratifies cancer subtypes) **Darren Creek** (456 - Combining metabolite standards cocktails with IDEOM v24 to enable routine semi-targeted metabolomics) **Per Andrén** (681 - Spatial metabolomics reveals region-specific alterations induced by parkinsonism and L-DOPA-induced dyskinesia) **Fan Yang** (16 - Untargeted mass spectrometry-based metabolomics workflow optimization for chronic and autoimmune chronic pancreatitis biomarker discovery) **Soumen Manna** (714 - Effect of wearing face mask on cardiopulmonary parameters and salivary metabolome) *Chair - Elizabeth Want*	**KEYNOTE: Ruijun Tian - Southern University of Science and Technology** **Christopher Overall** (413 - Multiplex MS Profiling of SARS-CoV-2 3CLpro/Main Protease Cleavage Kinetics for Ranking Interactors as Substrates and Designing Optimal Peptide Assay Formats) **Haoyun Fang** (448 - Deciphering subcellular proteomic niches of mouse heart using label-free DIA-MS and machine learning) **Bente Siebles** (343 - Spatial resolved mass spectrometric proteomics analysis facilitated by infrared laser-based sampling of infected murine bladders) **Kermit Murray** (563 - Laser Ablation Mass Spectrometry of Native Proteins) **Coleen Maxwell** (13 - The Edge Effect in High-Throughput Proteomics: A Cautionary Tale) *Chair - Lin Qingsong*	**KEYNOTE: Bruno Fedrizzi -The University of Auckland** **Oscar Nunez** (35 - High-throughput LC-LRMS and FIA-LRMS fingerprinting and polyphenolic profiling for the geographical characterization and authentication of honey) **Yada Novalchai** (165 - Enhancing food safety and authenticity: GC–MS analysis of adulterated edible oils and early detection of foodborne pathogens) **Jessica Prenni** (573 - The Periodic Table of Food Initiative) **Rebecca Jelley** (270 - Using HPLC-MS/MS to uncover novel thiols in hops) **Arundhati Singh** (447 - Mass spectrometry to unveil the foliar distribution of fluxapyroxad within fungicide seed-treated barley) *Chair - Zhongpin Yao*	**KEYNOTE: Perdita Barran -The University of Manchester** **Alex Shvartsburg** (568 - Novel Differential Ion Mobility Approaches Based on the Macromolecular Dipole Alignment) **Tim Causon** (545 - The role of unsaturation in defining the 3-dimensional structure of ionised lipids in the gas phase) **Christopher Wootton** (66 - A prototype TIMS-FT-ICR MS instrument capable of deep characterisation of complex samples) **Yiming Wang** (298 - Simultaneous Polyphenol Profiling and Quantification with LC-TIMS-TOF-MS: An Application to Different Apple Matrices) **Jackie Mosely** (565 - Structures for ion mobility resolved radical cations of benzocaine, and consequences for dissociation.) *Chair - Valarie Gabelica*	**KEYNOTE: Isabelle Compagnon - University of Lyon** **Andy Chi Kit Siu** (706 - Impact of Solvent Reorganization on Disulfide Bond Cleavage in Hydrated Electron Clusters) **Frantisek Turecek** (89 - Nitrile Imines as Photochemical Crosslinkers in Gas-Phase Peptide Ions: Reactivity and Action Spectroscopy) **Han Bin Oh** (166 - A combination of genetic code expansion and free radical-initiated peptide sequencing mass spectrometry) **Nicole Rijs** (190 - Directing Molecular Trams on the Picoscale! Perturbing Structural Outcomes of Self-Assembly Monitored by Ion Mobility Mass Spectrometry) **Caroline Dessent** (574 - Mapping the Photodegradation Products of Antibiotics Using Laser Interfaced Mass Spectrometry) *Chair - Chris Hansen*
Evening Workshops *Light Catering provided for evening workshop attendees from 5:00pm - 5:30pm*					
5.30 PM - 7.30 PM	**IMSF Focus Group: Native MS** *Integrating native and structural MS methods to solve open questions in structural/mechanistic biology and medicine* *Organisers: **Anton Calabrese** (University of Leeds)*	**IMSC Sustainability Workshop** *Sustainability in Mass Spectrometry: An International Perspective* Organisers**: Perdita Barran** (University of Manchester), **Gordon Kearney** (Shimadzu Research Laboratory, Europe) and **Chris Bowen** (Shimadzu Scientific Australasia)	**IMSC Paper Writing Workshop** *Write Right: How to Improve Your Scientific Writing* Organisers: **John Langley** (University of Southampton)		

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	Wednesday 21 August	Location
8.00 AM - 5:00 PM	Conference Registration	Main Foyer 3
8.30 AM - 9.30 AM	**Plenary - IMSF Curt Brunnee and Jochen Franzen awards**	Plenary
9.30 AM - 10.00 AM	**Morning Tea**	**Exhibition Hall**

Location	Plenary	Meeting Room 105	Meeting Room 106	Meeting Room 109	Meeting Room 110
Concurrent Sessions	**Imaging Mass Spectrometry: Applications**	**Biopharmaceuticals and Vaccines**	**Protein-Interactions**	**Forensics, Sports Doping, Homeland Security B**	**High Throughput Sampling, Screening and Analysis**
10.00 AM - 12.00 PM	**KEYNOTE: Kristine Glunde - Johns Hopkins Medical Institutions** **Isabelle Fornier** (372 - Click and Detect: A New Generation of Tagged Probes for Customed Targeted Multiplex MS Imaging) **Esther Cheow** (547 - Spatial Multi Omics Strategy to Advance Target Biology and Biomarker Discovery for Pulmonary Fibrosis) **Charles Schurman** (358 - Spatial Proteomics via Extracellular Matrix Imaging of Bone Fracture Callus Reveals Delayed Transition of Osteochondral Remodeling with Age) **Jayden Mckinnon** (260 - Unveiling Single Cell Small Metabolite Distributions via an Oversampling approach enabled by MALDI-2-MSI) **Haruki Ochino** (672 - Sex-dependent changes in renal spatial lipidome revealed by MALDI-2-TIMS-MS imaging) *Chair - Ron Heeren*	**KEYNOTE: Da Ren - BioTherapeutics Solutions** **Daniele Fabris** (340 - A Mid-Down Strategy for the Characterization of Non-Coding and mRNAs) **Phillip Bittner** (208 - In-depth Characterization of DNA-Encoded Chemical Libraries using Native Mass Spectrometry: The Impact of DNA-tags on Binding Affinities) **Devin Makey** (235 - Cyclic Ion Mobility-Mass Spectrometry for Rapid Protein Structure and Stability Assessment During the Development of Next-Generation Antibody Therapeutics) **Laura van der Vloet** (540 - Visualizing antisense oligonucleotides and its biological impact in brain tissue using a multi-omics mass spectrometry imaging approach) **Ilse Marialuce De Salve** (117 - Hybrid Immunocapture (IC)/LC-MS methods for the quantification of New Biological entities (NBE) and ADC (Antibody Drug Conjugates):perspective on different approaches.) *Chair - Matthias Pelzing*	**KEYNOTE: Lan Huang - University of California** **Ashleigh Dale** 633 - Membrane and flagellar enrichment increase the depth of large-scale bacterial interactomics studies using ion mobility and cross-linking mass spectrometry (XL-MS)) **Liuyu Peng** (196 - A Co-Fractionation Mass Spectrometry-Based Method for Investigating Disease Associated Alterations in Lipid-Protein Interactomes) **Venita Sitahal** (363 - Elucidating the Structural Dynamics of Binding Interactions in Regulator of G-Protein Signaling-1 (RGS1) using Hydrogen Deuterium Exchange Mass Spectrometry (HDX-MS)) **Adam Cahill** (322 - Development of photoactivatable lysine reactive crosslinking reagents) **Debashmita Ghosh** (337 - Mass Shifts Induction by Protein-Protein Interactions: A Novel Direct-MS Method) *Chair - Dezerae Cox*	**KEYNOTE: Adam Cawley - Racing Analytical Services Ltd** **Cailtin Jenkins** (10 - Chemical Analysis of Electronic Cigarettes in Australian Schools) **Madysen Elbourne** (543 - The indirect detection of dopaminergic manipulation in equine urine via an optimised routine and metabolomic-based LC-HRMS method.) **Kin-sing Wong** (47 - Screening and confirmation of recombinant human follistatin in equine plasma for doping control purposes) **Conor Jenkins** (570 - Soldier-on-a-Chip: Biomarker Identification through Multiomic Analyses of Chemical and Biological Threat Exposures to Model Organ-on-a-Chip Systems) **Bruce Pui-nam** (645 - Gene doping control analysis of human erythropoietin transgene in equine plasma by PCR-liquid chromatography high resolution tandem mass spectrometry) *Chair - Mario Thevis*	**KEYNOTE: Yashushi Ishihama - Kyoto University** **Laura Sanchez** (68 - Trapped ion mobility spectrometry for high-throughput directed evolution screening of α-ketoglutarate dependent dioxygenases) **Tin Cham Mak** (197 - Discovery of inhibitor against Mycobacterium tuberculosis Leucyl-tRNA synthetase (LeuRS) via Mass Spectrometry-based screening) **Rachel Smith** (229 - Development of automated high-throughput mass spectrometry methods for biotechnology and biomedical targets using desorption electrospray ionisation) **Xiaobo Tian** (45 - Differentiating specific and non-specific protein-metabolite interactions using gradient open port probe electrospray ionization mass spectrometry) **Chengyi Xie** (429 - Cellular-level resolution DESI-MS imaging) *Chair - Philippe Schmitt-Kopplin*

12.00 PM - 1.00 PM	**Lunch (no catering)** ***Sponsored Lunch Sessions:***	**Exhibition Hall**

Sponsored Lunch Seminars	**Meeting Room 105**	**Meeting Room 106**	**Meeting Room 109**
1.00 PM - 2.00 PM	Lunch Seminar - **Thermofisher**	Lunch Seminar - **Shimadzu**	Lunch Seminar - **Evosep**

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1:00 PM - 3:00 PM	**Innovation Stage Talks:** Merck 1:15pm -1:30pm Syft 1:30pm - 1:45pm Agilent 1:45pm - 2:00pm Trajan 2:00pm - 2:15pm Bruker 2:15pm - 2:30pm		Poster Session - **Wednesday Posters**		Exhibition Hall
2:30 PM - 3:00 PM	**Afternoon Tea**				**Exhibition Hall**
Location	**Plenary**	**Meeting Room 105**	**Meeting Room 106**	**Meeting Room 109**	**Meeting Room 110**
Concurrent Sessions	**Lipidomics: Technology**	**Proteomics: Quantitative**	**Polymers and Nanomaterials**	**Hyphenation, Separations, Lab-on-a-Chip**	**Ion Mobility B**
3.00 PM - 5.00 PM	**KEYNOTE: Makoto Arita - Keio University** **Takeshi Bamba** (126 - Solid phase extraction and LC/MS/MS methods for comprehensive targeted profiling of bioactive lipids) **Myeong Hee Moon** (72 - Optimization of skin sampling method for lipidomic analysis by nanoflow nUHPLC-ESI-MS/MS) **Gerard Hopfgartner** (367 - Enhanced Mass Spectrometry Workflows using ESI and APPI with Multi Ion Activation Methods for Characterization of Lipids in Plasma Samples) **Huong Giang Vo** (336 - Comprehensive analysis of structural glycosphingolipids in clinical samples using trapped ion mobility spectrometry mass spectrometry) **Huaqi Su** (54 - Multi-omics characterization of highly enriched human plasma extracellular vesicles) *Chair - Yu Xia*	**KEYNOTE: Jesper Olsen -University of Copenhagen** **Nathan Burke** (248 - Phosphoproteomic analysis of human sperm capacitation reveals novel, druggable kinases offering new non-hormonal male contraceptive targets) **Molly Talbot** (617 - Reversibly oxidised cysteine post-translational modifications in diabetic cardiomyopathy following antioxidant N-propionylglycine, identified using quantitative mass spectrometry.) **Subra Chakraborty** (712 - Climate Change & Food Security: Organellar Crosstalk and Post-translational Control Shaping Plant Immunity) **Scott Peterman** (518 - Novel Hybrid Nominal Mass Instrument Enables Rapid Development of Large-Scale Targeted Plasma Proteomics Assays) **Li Zhong** (11 - Mass Spectrometry Proteomics Reveals PLEK as a Biomarker for the Early Phase of Severe COVID-19) *Chair - Mark Malloy*	**KEYNOTE: Xiaopeng Li -Shenzhen University** **Scott McLuckey** (629 - Ion/Ion Reactions for Complex Mixture Analysis: Precursor Resolution via Ion Z-state Manipulation (PRIZM)) **Takaya Satoh** (128 - High Mass Resolution Mass Spectrometry for Assessing Polyethylene terephthalate Degradation: A Comprehensive Study using MALDI-TOFMS and GC-TOFMS) **Callan Littlejohn** (554 - Tools for Polymer Identification using Ultra-high Resolution Mass Spectrometry) **Elvis Okoffo** (548 - Quantitative analysis of micro- and nano-plastics in environmental samples by pyrolysis gas chromatography mass spectrometry) **Laura Puente-De La Cruz** (379 - Micro and nanoplastic migration from plastic breast milk storage bags and storage bottles) *Chair - Laurent Charles*	**KEYNOTE: Jin-Ming Lin -Tsinghua University** **Hartmut Schluter** (232 - Towards original compositions of proteoforms in tissues via soft laser ablation) **Andrei Fedorov** (67 - MicroTAS (Total Analysis System) for ESI-MS Dynamic Monitoring of Extracellular Proteome and Intracellular Metabolome in Cell and Gene Therapy Biomanufacturing) **Nick Manicke** (365 - On-Paper Electrophoretic Stacking and Separations Coupled to Paper Spray Mass Spectrometry) **John Langley** (26 - Sustainable fuels and the need for different hyphenated solutions) **Russell Grant** (562 - High Throughput Quantitative Amino Acid Analysis for The Masses) *Chair - Myeong Hee Moon*	**KEYNOTE: Kevin Pagel - Freie Universität Berlin** **Felicia Hansen** (497 - Direct observation of phospholipase A activity via isomer-resolved mass spectrometry.) **Bram van de Put** (325 - Towards De Novo Sequencing of Oligosaccharides Using Cyclic Ion Mobility Spectrometry) **David Marshall** (281 - Separation and characterisation of isomeric coordination complexes with high-resolution ion mobility mass spectrometry) **Madellen Wooding** (74 - The Story of a Sorptive Sampler: From Fairy Circles to the Detection of Tuberculosis-Associated Compounds using GC×GC-TOFMS and UPLC-IMS-HRMS) **Patricia Skowronek** (312 - Optimal trapped ion mobility workflows coupled with dia-PASEF and synchro-PASEF for high throughput and sensitivity) *Chair - Tim Causon*
Evening Workshops *Light Catering provided for evening workshop attendees from 5:00pm - 5:30pm*					
5.30 PM - 7.30 PM	**IMSF Focus Group: MS Instrumentation** *The journey from instrument concept, prototype development, to market* Organisers: **Shane Ellis** (University of Wollongong)		**IMSC Career Workshop** *Navigating career pathways for the mass spectrometrist* Organisers: **Jackie Mosely** (University of York)		

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Thursday 22 August						Location
8.00 AM - 5:00 PM	Conference Registration					Main Foyer 3
8.30 AM - 9.30 AM	**Plenary** **Professor Yu-ju Chen - Academia Sinica Taiwan** *Chair - Melanie White*					Plenary
9.30 AM - 10.00 AM	**Morning Tea** ***Innovation Stage Talks:*** Sciex 9:40am - 9:55am					**Exhibition Hall**
Location	**Plenary**	**Meeting Room 105**	**Meeting Room 106**	**Meeting Room 109**	**Meeting Room 110**	
Concurrent Sessions	**PTMs and Cross Talk, Cellular Signalling and Systems Biology**	**Cancer and Immunology**	**Environmental Mass Spectrometry B**	**Chemistry at the Solution/Gas Phase Interface**	**Ion Chemistry, Reactions and Structure C**	
10.00 AM - 12.00 PM	**KEYNOTE: Rena Robinson - Vanderbilt University** **Marco Jochem** (444 - Analysing Non-Proteinaceous Ubiquitination by Mass Spectrometry) **Guangcan Shao** (503 - Mapping the Modification Sites of Ubiquitin-like Proteins (UBLs)) **Siqi Li** (189 - Development of highly sensitive mass spectrometry methods for phosphorylated protein analysis) **Lin Zhu** (130 - Real-world PM2.5 Exposure Causes Liver Metabolic Reprogramming via Oxidation of Specific Cysteine Residues on MDH2 and CPT2 in Mice) **Maria Tanzer** (44 - Proteome profiling of macrophage reprogramming upon dead cell clearance) *Chair - Ben Parker*	**KEYNOTE: Bernd Wollscheid - Swiss Federal Institute of Technology** **Arash Zarrine-afsar** (14 - 10-second Classification of Lung Cancer Subtypes by Picosecond Infrared Laser Mass Spectrometry: Evaluation of Diagnostic Power Across Various Tissue Models) **Michelle Hill** (611 - Ovarian cancer serum glycoform biomarker panel discovery to lectin magnetic array-mass spec (LeMBA-MS) clinical assay) **Hong Yan** (80 - Machine Learning-Driven Identification of sex and KRAS specific features for Ferroptosis-Targeted Drug Repurposing in Colorectal Cancer) **Paula Nissen** (120 - Quantitative mass spectrometric proteome analysis of colorectal carcinoma liver metastasis reveals distinct phenotypes associated with specific signalling pathways and survival) **Nicole Brace** (680 - A Study of Altered B Cell Responses to PAMP-Activation in Schizophrenia) *Chair - Peter Hoffmann*	**KEYNOTE: Jean Armengaud - CEA** **Yuanjiang Pan** (188 - Wide-energy digital programmable microwave plasma for high coverage mass spectrometry analysis of various compounds) **Hang-kin Kong** (96 - Identifying Morphologically Similar Toxic Microalgal Strains by Proteomic Approaches) **Zhu Yang** (468 - Inducible energy source shift alleviates geo-specific PM2.5 components causing imbalances in energy metabolism) **Siobhan Peters** (460 - Bioaccumulation of the environmental neurotoxin BMAA in mussels exposed to cyanobacteria) **Albert Lebedev** (140 - Formation of Specific Disinfection By-products in Water treatment by Aqueous Chlorination) *Chair - Sutin Kingtong*	**KEYNOTE: Xinxing Zhang - Nankai University** **Jongcheol Seo** (359 - Investigating Protein Conformations Using Protein-Iodine Interactions) **Lukas Benzenberg** (29 - Microsolvation of charged residues prevents backbone collapse and aids in retention of native-like structural features after desolvation) **Valerie Gabelica** (307 - To unfold, or not to unfold, that is the question. On the preservation of solution phase structures upon electrospray ionization) **Sangwon Cha** (637 - Derivatization of Single Cell Saccharides and Analysis by Induced ESI MS) **Qianhao Min** (652 - Monitoring the Dynamic Fate of Interfacial Electrochemical Intermediates by Ambient Mass Spectrometry) *Chair - Shibdas Banerjee*	**KEYNOTE: Zheng Ouyang - Tsinghua University** **Howard Ma**(224 - Photoelectron Spectroscopy of Some Copper Borohydride and Acetylide Anions) **Salome Poyer** (507 - Copper-based mass spectrometry and ion mobility to resolve isomeric barriers of phosphatidylcholines) **Oscar Lloyd Williams** (462 - Of Cryptophanes and Cations: Unravelling structural changes induced by encapsulation or complexation.) **Shibdas Banerjee** (708 - Stabilizing Reactive Intermediates in Aqueous Microdroplets) **Takemichi Nakamura** (441 - Collision-Induced Isomerization of Cyclic Peptide Ions Probed by Energy-Resolved Ion Mobility/Tandem Mass Spectrometry: A Case of Anabaenopeptins) *Chair - Nicole Rijs*	
12.00 PM - 1.00 PM	**Lunch (no catering)** ***Sponsored Lunch Seminars:***					**Exhibition Hall**
Sponsored Lunch Seminars	**Meeting Room 105**					
1.00 PM - 2.00 PM	Lunch Seminar - **Thermofisher**					

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1:00 PM - 3:00 PM	***Innovation Stage Talks:*** Bruker 2:15pm - 2:30pm		Poster Session - **Thursday Posters**		Exhibition Hall
2:30 PM - 3:00 PM	**Afternoon Tea**				**Exhibition Hall**
Location	**Plenary**	**Meeting Room 105**	**Meeting Room 106**	**Meeting Room 109**	**Meeting Room 110**
Concurrent Sessions	**Neurodegeneration and Aging**	**Glycomics and Glycoproteomics**	**Single Cell Mass Spectrometry**	**Imaging Mass Spectrometry: Development/Technology**	**Data Science in Mass Spectrometry B**
3.00 PM - 5.00 PM	**KEYNOTE: Birgit Schilling - University of California San Francisco** **Martina Marchetti-deschmann** (481 - Spatially resolved analyses reveals significant changes of the N-glycome and transcriptome in aging skin) **Gaurav Chopra** (591 - Artificial Intelligent Agents for Automating Deep Lipidomics Workflows to Investigate Alzheimer's Disease and Aging-Related Lipid Droplets) **Michael Gotsbacher** (636 - Biodistribution of a Copper-Delivering Agent in Mouse Brains) **Emily Byrd** (324 - Understanding how ALS-associated mutations alter the structure and phase separation propensity of the TDP-43 C-terminal domain using structural mass spectrometry) **Emma Schymanski** (173 - Can Small Molecules Provide Clues on Disease Progression in Cerebrospinal Fluid from Mild Cognitive Impairment and Alzheimer's Disease Patients?) *Chair - Rena Robinson*	**KEYNOTE: Nicki Packer - Macquarie University** **Xue Sun** (616 -Large-Scale pattern analysis of N- and O-glycoproteomics using ion-mobility assisted mass spectrometry) **Ronghu Wu** (118 - Effective Mass Spectrometry-Based Methods for Comprehensive and Site-Specific Analysis of Surface Glycoproteins and Their Dynamics) **Abarna Murugan** (674 - Phyloglycomics: Mapping the serum glycome in vertebrates to understand the evolution of vertebrate glycosylation) **Mark Larance** (22 - Unbiased analysis of the human platelet proteome identifies a new form of domain-specific O-fucosylation generated by FUT10 and FUT11) **Kristian Tkalec** (83 - Protein Aggregation Capture enabled carboxylate group derivatisation allows proteome scale peptide supercharging for O-glycoproteomic analysis) *Chair - Cathy Costello*	**KEYNOTE: Yu Bai - Peking University** **Andrew Ewing** (346 - Correlation of Cellular Measurements of Partial Exocytosis and Nano Vesicular Subcompartments Using NanoSIMS) **David Bishop** (70 - Quantitative, Multiplexed, Immuno-Mass Spectrometry Imaging Of The Dystrophin-Glycoprotein Complex) **Reuben Young** (401 - Exploring different post-ionisation techniques coupled with MALDI-MSI for single cell lipidomics of neurons and astrocytes) **Mariachiara Squillaci** (366 - Single cell proteomics (SCP) analysis to study crosstalk dynamics of signaling complexes in extrinsic and intrinsic cell death pathways.) **Nhu Phan** (370 - Targeted molecular imaging with correlative NanoSIMS: Applications to study protein organization and turnover in neuronal cells.) *Chair - Lingjun Li*	**KEYNOTE: Helen Cooper - University of Birmingham** **Martin Sala** (601 - The Achilles heel of LA-ICP-MS (calibration) and how to solve it) **Jianing Wang** (127 - Subcellular Resolution MALDI Mass Spectrometry Imaging of Lipids) **Alice Ly** (212 - Weave: A software package for integrated spatial multi omics visualization and data analysis) **Shane Ellis** (383 - A MALDI-2-MSI source with transmission and reflective mode capabilities that exploits laminar gas flows of novel intermediate pressure ion guides) **Jan Preisler** (41 - Digital Mass Spectrometry Imaging using Nanoparticle Tags) *Chair - Isabelle Fornier*	**KEYNOTE: Magnus Palmblad - Leiden University Medical Center** **Emma Palm** (283 - Identifying dead-end pesticide transformation products of potential concern) **Sarah Bamford** (277 - Self-Organizing Maps with Relational Perspective Mapping (SOM-RPM) Applied to Time-of- Flight Secondary Ion Mass Spectrometry (ToF-SIMS) Data **Felix Servant** (642 - Statistical separation of simultaneously activated precursors) **Natan Horacek** (113 - An automatic and unsupervised artificial peak detection approach for preprocessing GC-MS and GC×GC-MS metabolomic data.) **Ove Johan Ragnar Gustafsson** (384 - Proteomics Lab: streamlining computational proteomics for life scientists) *Chair - Paul Pigrim*
6.00 PM - 10.00 PM	Conference Dinner - Showtime Events Centre				

Friday 23 August					**Location**
8.00 AM - 5:00 PM	Conference Registration				Main Foyer 3
8.30 AM - 9.30 AM	**Distinguished Keynote Speaker** **Koichi Tanaka - Executive Research Fellow, Shimadzu Corporation, Japan** *Chair - Gavin Reid*				Plenary
Location	**Plenary**	**Meeting Room 105**	**Meeting Room 106**	**Meeting Room 109**	**Meeting Room 110**
Concurrent Sessions	**Mass Analysers**	**Plant 'omics**	**Earth, Space, Geoscience and Atmospheric Chemistry B**	**Recent Advances in Mass Spectrometry, and Beyond**	**Ion Chemistry, Reactions and Structure D**

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9.30 AM - 11.00 AM	**KEYNOTE: Evan Williams - University of California, Berkeley** **Hamish Stewart** (62 - Behind the Curtain: The Long Development of Next Generation HRAM Analyzers.) **Yi-Sheng Wang** (TBC) **Taous Abar** (20 - Analysis of VOCs in aqueous solutions using the coupling of an oven with a mobile FT-ICR-MS associated to chemical ionization) *Chair - Zheng Ouyang*	**KEYNOTE: Joachim Kopka - Max-Planck-Institute of Molecular Plant Physiology (MPIMP)** **River Pachulicz** (42 - Acid-catalysed esterification of anthocyanin glucosyl units in Brassica oleracea) **Melanie Odenkirk** (546 - Juicy Insights: A Standardized, Nontargeted Metabolomics Approach to Facilitate the Comparison of Apples to Apples Across the Globe) **Laurent Bigler** (297 - Structure elucidation of iron chelators produced by microorganisms)*Chair - Ute Rosener*	**KEYNOTE: Haofei Zhang - University of California, Riverside** **Ralf Zimmermann** (335 - Application of a Novel, Field-Deployable Single Particle Mass Spectrometer for Detection of Toxicants and Climate-Relevant Parameters in Wildfire Aerosols) **Philippe Schmitt-Kopplin** (100 -Molecular atlas of meteorite soluble organic matter using non-targeted ultra)high resolution organic spectroscopy reveals hydrothermal history of asteroid Ryugu samples) **Trevor Ireland** (431 - Highs and Lows in Analysis of Water with SIMS) *Chair - Branka Miljevic*	**KEYNOTE: Yujia Qing - University of Oxford** **Niklas Geue** (98 - Visualizing Conformational Dynamics of Biomacromolecules by Coupling Ion Mobility Mass Spectrometry to Electron Microscopy) **Lar Gruber** (213 - Guiding imaging mass spectrometry by mid-infrared vibrational spectroscopy for deep lipidomic profiling) **Yanis Zirem** (350 - Spatial multi-omics informed by SVD k-means++ clustering and statistical assessment of heterogeneity: Advance dry proteomic guided by lipids MALDI MSI) *Chair - Justin Benesch*	**KEYNOTE: Vaughn Langford - Syft Technologies Limited** **Berwyck Poad** (284 - Hyphenation of chromatography with ion-ion charge inversion chemistries for the resolution and structure elucidation of branched-chain lipids) **Weiguo Wang** (341 - Miniature Continuous Atmospheric Pressure Interfaced Ion Trap Mass Spectrometer with Radial Electric Field Driven Collision-Induced Dissociation and its applications) **Hua Lei** (267 - Photoionization/photoionization-induced chemical ionization mass spectrometry for operando characterization of catalytic reaction processes: instrumentation and applications) *Chair - Scott McLuckey*
11:00 AM - 11:30 AM	**Morning Tea**				**Exhibition Hall**
Location	**Plenary**	**Meeting Room 105**	**Meeting Room 106**	**Meeting Room 109**	**Meeting Room 110**
Concurrent Sessions	**Toxicology and Metabolism**	**Biosimilars, Biobetters and Glycoengineering**	**Cultural Heritage, Conservation and Archaeology**	**JMS Awardee Session**	**High Resolution Mass Spectrometry**
11.30 AM - 1.00 PM	**KEYNOTE: Zongwei Cai - Hong Kong Baptist University** **Maya Cameron** (393 - Comprehensive analysis of Endocrine Disrupting Chemicals using High Resolution Mass Spectrometry) **Denise Tran** (592 - Method Development for High-throughput Quantification of PFAS in Plasma for Correlation to the Omics Profile of Patients with Cardiovascular Disease) **Gwendolyn Cooper** (38 - Untargeted metabolomic analysis of acute lead exposure to Danio rerio embryos using LC-MS/MS) *Chair - Laura Sanchez*	**KEYNOTE: Alain Beck - Laboratoires Pierre Fabre** **Magdelina Biesaga** (37 - Identification of surfactants with antifungal activity produced by antarctic bacteria Bacillus subitlis strain) **Troy Wood** (230 - Quality Assurance Using Mass Spectrometry to Analyze Structural Fidelity of Monoclonal Antibodies in HIV-1 Therapeutics) **Adam Pruska** (345 - Temperature-Controlled Mass Spectrometry as a Tool for Structural Characterization of Enzymes and Antibodies) *Chair - Hiroshi Hinou*	**KEYNOTE: Asher Newsome - Smithsonian Museum Conservation Institute** **Renaud Joannes-Boyau** (TBC) **Teodora Raicu** (506 - MeV SIMS Approach for Identifying Colorants in Artists' Modern Inks) **Paul Haynes** (TBC) *Chair - Paul Haynes*	*Chair - Miaomiao Liu*	**KEYNOTE: Vicki Wysocki - Ohio State University** **Lee Gethings** (216 - Comprehensive discovery lipidomic workflow which utilizes a prototype, multi-reflecting ToF with integrated informatics, providing highly confident lipid characterization and quantification) **Jordan Partington** (459 - Comparison of high-resolution mass spectrometry acquisition methods for the simultaneous quantification and identification of per- and polyfluoroalkyl substances (PFAS)) **Benedict Gannon** (560 - Effects of sustainable rejuvenator on aged British roads compared with commercial bitumen binder rejuvenator by FT-ICR MS) *Chair - Ljiljana Pasa-Tolic*
1.00 PM - 2.00 PM	**Lunch (no catering)**				
2.00 PM - 3.00 PM	**Closing Plenary** **Professor Veena Sahajwalla - University of New South Wales** *Chair - Sarah Hancock*				Plenary
3.00 PM - 4.00 PM	Closing Ceremony, Award Presentations and Presentation of IMSC2026				Plenary