



15th Annual Conference of the Metabolomics Society

METABOLOMICS 2019

JUNE 23–27 | THE HAGUE

SCHEDULE OF ORAL PRESENTATIONS



AGENDA AT A GLANCE

	Biomedical		Technology
	Plant, Food, Environmental and Microbial		New Frontiers

SUNDAY, JUNE 23				
	Atlantic	Alexia	Ariane	Oceania Foyer
11:00 a.m.	REGISTRATION OPEN			
12:30 p.m. – 2:15 p.m.	W1: EMN – Data Fusion	W2: Mining the Metabolome	W3: Multi-Omics Integration & Systems Metabolomics	Intro to the Field 1: Data Acquisition
2:30 p.m. – 4:15 p.m.	W4: Application of Graphical Models to Metabolomics	W5: Plant Metabolomes: Natural & Generated Variability	W6: How to Link Metabolome & Genome Mining	Intro to the Field 1: Data Acquisition
4:30 p.m. – 6:15 p.m.	W7: EMN – Professional Career Development	W8: Putting Metabolomic Data into Context	W9: An Open-Source Pipeline for Appraisal of NMR Datasets	Intro to the Field 1: Data Acquisition
6:30 p.m. – 8:30 p.m.	Career Night – Pacific			
MONDAY, JUNE 24				
	Atlantic	Alexia	Ariane	Oceania Foyer
7:30 a.m.	REGISTRATION / INFO DESK OPEN			
8:30 a.m. – 10:15 a.m.	W10: The Importance of Quality Assurance & Quality Control	W11: Towards FAIR Spectral Libraries	W12: EMN – Stable Isotope-Resolved Metabolomics	Intro to the Field 2: Data (pre) Processing and Biostatistics
10:30 a.m. – 12:15 p.m.	W13: Beyond pathway mapping	W14: Standardizing the Fluxomics Workflows	W15: Tools to Study the Microbiome-Metabolome Interplay	Intro to the Field 2: Data (pre) Processing and Biostatistics
12:15 p.m. – 1:30 p.m.	LUNCH BREAK – ON YOUR OWN			
1:30 p.m. – 3:15 p.m.	W16: Application Metabolomics in Industry	W17: Dynamic Modeling of Human Metabolism	W18: EMN – Volatomics in Human Health	Intro to the Field 2: Data (pre) Processing and Biostatistics
3:30 p.m. – 5:00 p.m.	Opening Ceremony Plenary Session 1 – Joshua Rabinowitz – King Willem-Alexander Hall			
5:15 p.m. – 6:45 p.m.	Welcome Reception – Poster Session 1 – Odd Numbers – Exhibit Foyer			
7:00 p.m. – 8:00 p.m.	Metabolomics Society Town Hall Meeting – Atlantic			
TUESDAY, JUNE 25				
	Atlantic	King Willem-Alexander	Alexia	Ariane
7:45 a.m.	REGISTRATION / INFO DESK OPEN			
8:30 a.m. – 9:30 a.m.	Plenary Session 2 – Dorret Boomsma – King Willem-Alexander Hall			
9:30 a.m. – 10:15 a.m.	BREAK – EXHIBIT FOYER			
10:15 a.m. – 12:00 p.m.	1. Cancer	2. Plant Applications 1	3. Data Integration & Data Basing 1	4. Novel Technologies
12:00 p.m. – 1:30 p.m.	LUNCH – IN FOYER WITH EXHIBITS – PLATINUM SPONSOR PRESENTATIONS			
12:20 p.m. – 1:20 p.m.			Sponsor Pres: SCIEX	Sponsor Pres: Waters Corporation
1:30 p.m. – 3:15 p.m.	5. Metabolic Disease	6. Food Applications 1	7. Flux Studies	8. Novel Instruments, Tools and Services
3:15 p.m. – 3:45 p.m.	BREAK – EXHIBIT FOYER			
3:45 p.m. – 5:30 p.m.	9. Epidemiology	10. Microbial Applications	11. Metabolite Identification 1	12. Stem Cells, Organoids
5:30 p.m. – 7:00 p.m.	Poster Session 2 – Odd Numbers – Exhibit Foyer			
7:00 p.m. – 8:30 p.m.	EMN Reception – Pacific			
WEDNESDAY, JUNE 26				
	Atlantic	King Willem-Alexander	Alexia	Ariane
8:00 a.m.	REGISTRATION / INFO DESK OPEN			
8:30 a.m. – 9:30 a.m.	Plenary Session 3 – Cathie Martin – King Willem-Alexander Hall			
9:30 a.m. – 10:15 a.m.	BREAK – EXHIBIT FOYER			
10:15 a.m. – 12:00 p.m.	13. Lipidomics and Cardiovascular Disease	14. Plant Defense	15. Data Analysis & Statistics	16. Single Cell
12:00 p.m. – 1:30 p.m.	LUNCH – IN FOYER WITH EXHIBITS – PLATINUM SPONSOR PRESENTATIONS			
12:20 p.m. – 1:20 p.m.			Sponsor Pres: Thermo Fisher Scientific	Sponsor Pres: Shimadzu Europa GmbH
1:30 p.m. – 3:15 p.m.	17. Ageing and Disease	18. Food Applications 2	19. Data Integration & Data Basing 2	20. Regulatory Session
3:15 p.m. – 3:45 p.m.	BREAK – EXHIBIT FOYER			
3:45 p.m. – 5:30 p.m.	21. Infection and Immunity	22. Environment & Toxicology	23. New Instrumentation	24. Novel Applications
5:30 p.m. – 7:00 p.m.	Poster Session 3 – Even Numbers – Exhibit Foyer			
7:30 p.m. – 10:30 p.m.	Conference Dinner – Xiringuito			
THURSDAY, JUNE 27				
	Atlantic	King Willem-Alexander	Alexia	Ariane
8:30 a.m.	REGISTRATION / INFO DESK OPEN			
8:45 a.m. – 10:00 a.m.	25. Respiratory Diseases			26. Model Organisms
10:00 a.m. – 10:45 a.m.	Poster Session 4 – Even Numbers – Exhibit Foyer			
10:45 a.m. – 12:00 p.m.	28. Microbiome			29. Plant Applications 2
12:00 p.m. – 1:30 p.m.	LUNCH – IN FOYER WITH EXHIBITS – PLATINUM SPONSOR PRESENTATIONS			
12:20 p.m. – 1:20 p.m			Sponsor Pres: Bruker Daltonics/Bruker Biospin	Sponsor Pres: Agilent Technologies
1:30 p.m. – 3:30 p.m	Plenary Session 4 – Jean-Charles Portais – Closing Ceremony – King Willem-Alexander Hall			

BIOMEDICAL
PLANT, FOOD, ENVIRONMENTAL AND MICROBIAL

***AWARD WINNERS**

Monday, June 24		
Time	Session	Location
3:30 p.m. – 5:00 p.m.	Opening Ceremony and Plenary Session 1 <i>Joshua Rabinowitz, Lewis-Sigler Institute, United States</i>	
Tuesday, June 25		
Time	Session	Location
8:30 a.m.– 9:30 a.m.	Plenary Session 2 <i>Dorret I. Boomsma, Netherlands Twin Register, Vrije Universiteit, Netherlands</i>	
10:15 a.m. – 12:00 p.m.	Session 1: Cancer <i>(Session Chairs: Hunter Moseley & Maria Eugenia Monge)</i>	
10:15 a.m. – 10:45 a.m.	1A SESSION KEYNOTE Metabolomics Applications for Oncology and Immunology in Drug Discovery <i>Thomas Roddy, Agios Pharmaceuticals, United States</i>	Atlantic
10:45 a.m. – 11:05 a.m.	1B Multi-omic discovery of metabolic rewiring in triple-negative breast cancer following mitochondrial folate transport ablation <i>Steven Gross, Weill Cornell Medicine, United States</i>	
11:05 a.m. – 11:20 a.m.	1C Investigating the metabolic response of cancer cells to indisulam <i>Lili Herendi, PhD student, United Kingdom</i>	
11:20 a.m. – 11:40 a.m.	1D Rewiring of energy metabolism drives resistance to the proteasome inhibitor bortezomib <i>Esther Zaai, Biomolecular Mass Spectrometry and Proteomics, Utrecht University, Netherlands</i>	
11:40 a.m. – 11:55 a.m.	1E* Understanding metabolite heterogeneity in pancreatic ductal adenocarcinoma, and the role of adaptive metabolic reprogramming in chemotherapy resistance <i>Sarah Hancock, University of New South Wales, Australia</i>	
10:15 a.m. – 12:00 p.m.	Session 2: Plant Applications 1 <i>(Session Chairs: Sanjay Swarup & Jasen Finch)</i>	
10:15 a.m. – 10:45 a.m.	2A SESSION KEYNOTE The volatile flavor composition network in tomato and its modification <i>Antonio Granell, CSIC, Spain</i>	King Willem Alexander
10:45 a.m. – 11:05 a.m.	2B Developing Advanced and Integrated Metabolomics Technologies to Address the Grand Challenges of Metabolite Identification and Depth of Coverage <i>Lloyd Sumner, University of Missouri, United States</i>	
11:05 a.m. – 11:20 a.m.	2C* From MS peak to unambiguous metabolite identification using the WeizMass spectral library and LC-MS-SPE-NMR system <i>Adam Jozwiak, Weizmann Institute of Science, Israel</i>	
11:20 a.m. – 11:40 a.m.	2D Chocolate metabolomics <i>Robert Hall, Wageningen UR, Netherlands</i>	
11:40 a.m. – 11:55 a.m.	2E A tissue specific metabolomic study in hybrid aspen <i>Ilara Budzinski, Swedish University of Agricultural Sciences (SLU), Sweden</i>	

***AWARD WINNERS**

Tuesday, June 25		
Time	Session	Location
10:15 a.m. – 12:00 p.m.	Session 3: Data Integration & Data Basing 1 (Session Chairs: Fabien Jourdan & Natasa Giallourou)	
10:15 a.m. – 10:45 a.m.	3A SESSION KEYNOTE The Internet for Social Machines <i>Barend Mons, GO FAIR, Netherlands</i>	Princess Alexia
10:45 a.m. – 11:05 a.m.	3B Matching Untargeted Liquid Chromatography - Mass Spectrometry Features Across Multiple Cohorts: Finding the Same Needles in Several Haystacks Via Networks <i>Rui Climaco Pinto, Imperial College London, United Kingdom</i>	
11:05 a.m. – 11:20 a.m.	3C* Visualizing metabolomics data in directed biological networks <i>Denise Slenter, Maastricht University, Netherlands</i>	
11:20 a.m. – 11:40 a.m.	3D High-throughput metabolomics identifies substrate-enzyme relationships in a metabolism-wide CRISPR interference library <i>Hannes Link, Max Planck Institute for Terrestrial Microbiology, Germany</i>	
11:40 a.m. – 11:55 a.m.	3E* Weighting strategies for the analysis of secondary outcomes in nested case-control metabolomics data <i>Gerard Gonzales, Ghent University, Belgium</i>	
10:15 a.m. – 12:00 p.m.	Session 4: Novel Technologies (Session Chairs: Rob Vreeken & Lieven van Meulebroek)	
10:15 a.m. – 10:45 a.m.	4A SESSION KEYNOTE Improving Technologies for High Throughput and Miniaturized Metabolomics for Precision Medicine <i>Thomas Hankemeier, Leiden University, Netherlands</i>	Princess Ariane
10:45 a.m. – 11:05 a.m.	4B Investigation of host-microbiota co-metabolism as a new strategy for biomarker discovery – New Chemical Biology tools for Metabolomics analysis <i>Daniel Globisch, Uppsala University, Sweden</i>	
11:05 a.m. – 11:20 a.m.	4C* GC×GC-TOFMS and SIFT-MS approaches for clinical breath-based asthma phenotyping <i>Pierre-Hugues Stefanuto, Liège University, Belgium</i>	
11:20 a.m. – 11:40 a.m.	4D Mathematical modelling of metabolism: a driver for developing personalized and precision medicine <i>Natal van Riel, Eindhoven University of Technology, Netherlands</i>	
11:40 a.m. – 11:55 a.m.	4E TIMS and PASEF multiply speed and sensitivity in lipidomics <i>Catherine G. Vasilopoulou, Max Planck Institute of Biochemistry, Germany</i>	

*AWARD WINNERS

Tuesday, June 25		
Time	Session	Location
1:30 p.m. – 3:15 p.m.	Session 5: Metabolic Disease (Session Chairs: Christophe Junot & Katharina Herzog)	
1:30 p.m. – 2:00 p.m.	5A SESSION KEYNOTE Integrating epidemiologic, pharmacologic, genetic and gut microbiome data in the BBMRI-NL drug-metabolome atlas Cornelia van Duijn, University of Oxford, United Kingdom	Atlantic
2:00 p.m. – 2:20 p.m.	5B Metabolic profiling of tissue-specific insulin resistance in human obesity: Results from the Diogenes Study and The Maastricht Study Ilja Arts, Epidemiology & MaCSBio, Maastricht University, Netherlands	
2:20 p.m. – 2:35 p.m.	5C Identification of novel metabolites in alkaptonuria by LC-QTOF-MS profiling and flux analysis of a targeted HGD-/- mouse model Brendan Norman, Institute of Ageing & Chronic Disease, University of Liverpool, United Kingdom	
2:35 p.m. – 2:55 p.m.	5D* Combining untargeted metabolomics, human genetics, causal inference, and pathway enrichment to define the obesity metabolome Yu-Han Hsu, Harvard Medical School, United States	
2:55 p.m. – 3:10 p.m.	5E* Metabolomics of Ndufs4-/- skeletal muscle: adaptive mechanisms converge at the ubiquinone-cycle Karin Terburgh, North-West University, South Africa	
1:30 p.m. – 3:15 p.m.	Session 6: Food Applications 1 (Session Chairs: Cristina Andres Lacueva & Jessica Cooperstone)	
1:30 p.m. – 2:00 p.m.	6A SESSION KEYNOTE A metabolomic study on coffee – From coffee brews to liver cancer risk Augustin Scalbert, International Agency for Research on Cancer (IARC), France	King Willem Alexander
2:00 p.m. – 2:20 p.m.	6B Integrated analysis of metabolomics and microbiome data showing additional dietary effects Jildau Bouwman, TNO, Netherlands	
2:20 p.m. – 2:35 p.m.	6C Quantitative Dietary Fingerprinting (QDF)—A Novel Tool for Comprehensive Dietary Assessment Based on Urinary Nutrimetabolomics Raúl González-Domínguez, Biomarkers and Nutrimetabolomics Laboratory, University of Barcelona, Spain	
2:35 p.m. – 2:55 p.m.	6D Grass, beef and human. Unlocking the value of the food chain in New Zealand Arvind Subbaraj, AgResearch Ltd., New Zealand	
2:55 p.m. – 3:10 p.m.	6E NMR metabolomics revealed different impact of non-acylated and acylated anthocyanins on plasma metabolic profiles in obese diabetic Zucker rats Kang Chen, University of Turku, Finland	

*AWARD WINNERS

Tuesday, June 25

Time	Session	Location
1:30 p.m. – 3:15 p.m.	Session 7: Flux Studies (Session Chairs: Barbara Bakker & Bekzod Khakimov)	
1:30 p.m. – 2:00 p.m.	7A SESSION KEYNOTE A systems medicine approach to identify new drug targets: Model-driven discovery of metabolic reprogramming in metastatic prostate cancer <i>Marta Cascante, University of Barcelona, Spain</i>	Princess Alexia
2:00 p.m. – 2:20 p.m.	7B Using stable-isotope labelled metabolomics to explore cell-cell communication between malaria parasites and host cells <i>Darren Creek, Monash University, Australia</i>	
2:20 p.m. – 2:35 p.m.	7C* Using deuterium labelled glucose to quantify redox metabolism in plants <i>Edward Smith, University of Oxford, United Kingdom</i>	
2:35 p.m. – 2:55 p.m.	7D* Modelling Cancer Lipogenesis using REIMS Metabolic Flux analysis in breast cancer cell lines <i>Seyma Turkseven, Imperial College London, United Kingdom</i>	
2:55 p.m. – 3:10 p.m.	7E* Exploring the use of GC-MS for stable isotope labeling in metabolomics <i>Jordi Capellades Tomàs, IISPV, Spain</i>	
1:30 p.m. – 3:15 p.m.	Session 8: Novel Instrumentations, Tools and Services (Presented by Platinum and Gold Sponsors) (Session Chairs: Thomas Hankemeier & David Wishart)	
1:30 p.m. – 2:30 p.m.	PLATINUM PRESENTERS AND PANEL DISCUSSION Agilent Technologies: Christine Miller, Omics Market Manager, USA Bruker Daltonics/Bruker BioSpin: Lucy Woods, PhD, Product Manager QTOF, Germany SCIEX: Baljit Ubhi, Market Manager - Metabolomics Business, USA Shimadzu: Emily Armitage, Research Scientist, UK Thermo Fisher Scientific: Amanda Souza, Manager, Product Marketing, USA Waters Corporation: David Heywood, Senior Manager Omics Business Development, UK	Princess Ariane
2:30 p.m. – 3:15 p.m.	GOLD PRESENTERS AND PANEL DISCUSSION Biocrates Life Sciences AG: Therese Koal, PhD, Head of Research & Development, Austria Cambridge Isotope Laboratories, Inc: Krista Backiel, Marketing and Metabolomics Manager, USA Human Metabolome Technologies: Tom Hoshiba, Managing Director, Netherlands Metabolon: Alex Forrest-Hay, VP Population Health, USA	

*AWARD WINNERS

Tuesday, June 25		
Time	Session	Location
3:45 p.m. – 5:30 p.m.	Session 9: Epidemiology (Session Chairs: Krista Zanetti & Julie Schmidt)	
3:45 p.m. – 4:15 p.m.	9A SESSION KEYNOTE Integrating metabolomics with genomics, proteomics, and other omics for health and drug research <i>Karsten Suhre, Weill Cornell Medicine - Qatar, Qatar</i>	Atlantic
4:15 p.m. – 4:35 p.m.	9B Application of 1H NMR metabolomics in ~7,000 people to investigate potential molecular mechanisms of genetic risk variants for coronary artery diseases <i>Ibrahim Karaman, Imperial College London, United Kingdom</i>	
4:35 p.m. – 4:50 p.m.	9C Untargeted metabolomics in a prospective cohort to identify diet-related metabolites associated with age-related cognitive decline <i>Dorrain Low, Lee Kong Chian School of Medicine, Nanyang Technological University, Singapore</i>	
4:50 p.m. – 5:10 p.m.	9D Metabolic profile and the risk of developing prostate cancer - A nested-case control study <i>Ali Moazzami, Department of Molecular Sciences, SLU, Sweden</i>	
5:10 p.m. – 5:25 p.m.	9E The Metabolome of BMI: A Consortium of METabolomics Studies (COMETS) Meta-analysis of 85,000 adults <i>Rachel Kelly, HMS and BWH, United States</i>	
3:45 p.m. – 5:30 p.m.	Session 10: Microbial Applications (Session Chairs: Adriana Carvalho de Souza & Douglas McCloskey)	
3:45 p.m. – 4:15 p.m.	10A SESSION KEYNOTE Functional characterization of Escherichia coli lipid genes <i>Jos Brouwers, Utrecht University, Netherlands</i>	King Willem Alexander
4:15 p.m. – 4:35 p.m.	10B Metabolomics-based engineering biology of microorganisms <i>Tomohisa Hasunuma, Engineering Biology Research Center, Kobe University, Japan</i>	
4:35 p.m. – 4:50 p.m.	10C* Expanding the Applications of Rapid Evaporative Ionisation Mass Spectrometry (REIMS) to the Pharmaceutical Product Development Workflow <i>Toma Ramonaite, Imperial College London, United Kingdom</i>	
4:50 p.m. – 5:10 p.m.	10D Linking metabolic function to member taxa in complex microbial communities using metabolome-guided multi-omics <i>Nay Min Min Thaw Saw, Singapore Center for Environmental Life Science Engineering, Singapore</i>	
5:10 p.m. – 5:25 p.m.	10E* Microbial metabolic networks: the hidden key to resilience of coral algal endosymbionts <i>Jennifer Matthews, University of Technology Sydney, Australia</i>	

***AWARD WINNERS**

Tuesday, June 25		
Time	Session	Location
3:45 p.m. – 5:30 p.m.	Session 11: Metabolite Identification 1 (Session Chairs: Justin van der Hooft & Sonia Liggi)	
3:45 p.m. – 4:15 p.m.	11A SESSION KEYNOTE Infrared ion spectroscopy: new opportunities for molecular structure identification in MS-based metabolomics <i>Jos Oomens, Radboud University, Netherlands</i>	Princess Alexia
4:15 p.m. – 4:35 p.m.	11B mFam – prediction of metabolite families based on spectral data <i>Hendrik Treutler, Dept. of Biochemistry of Plant Interactions, Leibniz Institute of Plant Biochemistry, Germany</i>	
4:35 p.m. – 4:50 p.m.	11C* Supercharging Comparative Metabolomics with METABOseek <i>Maximilian Helf, Cornell University / Boyce Thompson Institute, United States</i>	
4:50 p.m. – 5:10 p.m.	11D Ion mobility – mass spectrometry based multi-dimensional metabolite annotation <i>Zhiwei Zhou, IRCBC, Shanghai Institute of Organic Chemistry, Chinese Academy of Sciences, China</i>	
5:10 p.m. – 5:25 p.m.	11E* Recommending substructures for unknown tandem mass spectra <i>Youzhong LIU, University of Antwerp, Belgium</i>	
3:45 p.m. – 5:30 p.m.	Session 12: Stem Cells/Organoids/Organ-on-a-Chip (Session Chairs: Alain van Gool & Isabelle Kohler)	
3:45 p.m. – 4:15 p.m.	12A SESSION KEYNOTE No transporters means no transport – and assessment of the ‘real’ (natural) substrates of xenobiotic transporters <i>Douglas Kell, University of Liverpool, United Kingdom</i>	Princess Ariane
4:15 p.m. – 4:35 p.m.	12B Spatial Isotope tracer Metabolomics to study 13C labelled Metabolite Distribution in 3D Tumor Spheroid Cell culture <i>Prasad Phapale, EMBL, Germany</i>	
4:35 p.m. – 4:50 p.m.	12C* Microengineered Human Blood Vessel for Next Generation Drug Discovery <i>Abidemi Junaid, Systems Biomedicine and Pharmacology, LACDR, Leiden University, Netherlands</i>	
4:50 p.m. – 5:10 p.m.	12D* Stable Isotope-Resolved Metabolomics (SIRM) Defines DEK Oncogene Driven Metabolic Reprogramming in 3D Epidermal Organoids <i>Sara Vicente-Muñoz, NMR-Based Metabolomics Core Facility, Division of Pathology and Laboratory Medicine, United States</i>	
5:10 p.m. – 5:25 p.m.	12E Metabolomics as a quality control for the production of chondrogenic microtissues towards characterized endochondral bone regeneration <i>Niki Loverdou, Prometheus, Division of Skeletal Tissue Engineering, KULeuven, Belgium</i>	

Wednesday, June 26

Time	Session	Location
8:30 a.m. – 9:30 a.m.	Plenary Session 2 <i>Cathie Martin, John Innes Centre, Colney Research Park, UK</i>	
10:15 a.m. – 12:00 p.m.	Session 13: Lipidomics and Cardiovascular Disease <i>(Session Chairs: Jules Griffin & Aline Martins)</i>	
10:15 a.m. – 10:45 a.m.	13A SESSION KEYNOTE High-throughput lipidomic quantitation of human blood in cancer screening <i>Michal Holcapek, University of Pardubice, Czech Republic</i>	Atlantic
10:45 a.m. – 11:05 a.m.	13B Changes in plasma lipids predict pravastatin efficacy in secondary prevention <i>Peter Meikle, Baker Heart and Diabetes Institute, Australia</i>	
11:05 a.m. – 11:20 a.m.	13C* Metabolomics analysis of human atherosclerotic plaques reveals a potential novel pathway of macrophage foam cell apoptosis in advanced atherosclerosis <i>Panagiotis Vorkas, Imperial College London, United Kingdom</i>	
11:20 a.m. – 11:40 a.m.	13D Plasma metabolomics profiles associated with endothelial health and dysfunction impose changes to endothelial glycan biosynthesis and reflect endothelial catecholamine response <i>Óttar Rolfsson, University of Iceland, Iceland</i>	
11:40 a.m. – 11:55 a.m.	13E Sphingolipidomics investigation of the ischemic brain injury <i>Ching-Hua Kuo, National Taiwan University, Taiwan</i>	
10:15 a.m. – 12:00 p.m.	Session 14: Plant Defense <i>(Session Chairs: Annick Moing & Halford Dace)</i>	
10:15 a.m. – 10:45 a.m.	14A SESSION KEYNOTE Elucidating insect resistance in tomato through genetical metabolomics <i>Ric de Vos, Bioscience, Wageningen Plant Research, Wageningen University and Research, Netherlands</i>	King Willem Alexander
10:45 a.m. – 11:05 a.m.	14B Dissecting the Genetic Basis of Variation in Tomato Fruit Metabolism and Pathogen Resistance through Multimodal Investigation of Wild Species Introgressions <i>Jedrzej Szymanski, Leibniz Institute of Plant Genetics and Crop Plant Research, Germany</i>	
11:05 a.m. – 11:20 a.m.	14C Metabolomics application to unravel the biochemistry underlying enhancement of drought stress tolerance in crops by plant growth promoting rhizobacteria <i>Fidele Tugizimana, University of Johannesburg, South Africa</i>	
11:20 a.m. – 11:40 a.m.	14D A strategy for the discovery and characterisation of health and decline biomarkers in British Oak trees using non-targeted metabolomics <i>Jasen Finch, Aberystwyth University, United Kingdom</i>	
11:40 a.m. – 11:55 a.m.	14E* Linking the volatilome and metabolome via plant defenses: stimulation of the salicylic acid pathway recruits natural enemies below ground <i>Camila Filgueiras, Cornell University, United States</i>	

***AWARD WINNERS**

Wednesday, June 26		
Time	Session	Location
10:15 a.m. – 12:00 p.m.	Session 15: Data Analysis & Statistics (Session Chairs: Age Smilde & Isabel Orf)	
10:15 a.m. – 10:45 a.m.	15A SESSION KEYNOTE Is Metabolomics ready for the return of Artificial Neural Networks? <i>David Broadhurst, Edith Cowan University, Australia</i>	Princess Alexia
10:45 a.m. – 11:05 a.m.	15B MS-DIAL 4.0: a computational workflow for ion mobility tandem mass spectrometry data in metabolomics <i>Hiroshi Tsugawa, RIKEN, Japan</i>	
11:05 a.m. – 11:20 a.m.	15C* Chemically informed distance metrics for tandem mass spectrometry data <i>Madeleine Ernst, Department of Congenital Disorders, Statens Serum Institut, Denmark</i>	
11:20 a.m. – 11:40 a.m.	15D Identifying biologically relevant modules in metabolomics and lipidomics data with Differential Network-based Enrichment Analysis (DNEA) <i>Alla Karnovsky, University of Michigan, United States</i>	
11:40 a.m. – 11:55 a.m.	15E Signature Mapping (SigMa): A new automatic tool for rapid processing of complex urine NMR spectra <i>Bekzod Khakimov, University of Copenhagen, Denmark</i>	
10:15 a.m. – 12:00 p.m.	Session 16: Single Cell (Session Chairs: Jildau Bouwman & Michel van Weeghel)	
10:15 a.m. – 10:45 a.m.	16A SESSION KEYNOTE Spatial metabolomics in tissues and single cells <i>Theodore Alexandrov, EMBL, Germany</i>	Princess Ariane
10:45 a.m. – 11:05 a.m.	16B Comprehensive Single Cell Multi Omics <i>Christian Berchtold, Institute for Chemistry and Bioanalytics, Switzerland</i>	
11:05 a.m. – 11:20 a.m.	16C Quantifying heterogeneity in drug uptake, metabolism and effect using Raman spectroscopy and mass spectrometry on the single-cell level <i>Ahmed Ali, Leiden University, The Netherlands RIKEN, Japan, Japan</i>	
11:20 a.m. – 11:40 a.m.	16D* Relationships between cellular metabolite and drug uptake and transporter expression profiles <i>Marina Muelas, University of Liverpool, United Kingdom</i>	
11:40 a.m. – 11:55 a.m.	16E* A Single-Cell Look at Biological Nitrogen Fixation: Direct Determination of Metabolite Formulas from Isotopic Fine Structures in Heterogeneous Cell Populations <i>Tina Tran, The George Washington University, United States</i>	

*AWARD WINNERS

Wednesday, June 26		
Time	Session	Location
1:30 p.m. – 3:15 p.m.	Session 17: Ageing and Disease (Session Chairs: <i>Matej Oresic & Candice Ulmer</i>)	
1:30 p.m. – 2:00 p.m.	17A SESSION KEYNOTE Multiomic data integration using machine learning and data-driven inverse metabolic modelling - from diabetes to immune system modulation <i>Wolfram Weckwerth, University of Vienna, Austria</i>	Atlantic
2:00 p.m. – 2:20 p.m.	17B Mass spectrometric analysis of sebum contents for classification of Parkinson's disease <i>Drupad Trivedi, University of Manchester, United Kingdom</i>	
2:20 p.m. – 2:35 p.m.	17C Mechanistic model-driven exometabolomic characterisation of human dopaminergic neuronal metabolism <i>Ronan Fleming, Leiden University, Netherlands</i>	
2:35 p.m. – 2:55 p.m.	17D Interleaving metabolic effects of sleep and aging <i>Arjun Sengupta, University of Pennsylvania, United States</i>	
2:55 p.m. – 3:10 p.m.	17E Analysis of changes in the eye lens and aqueous humor under cataract development using quantitative metabolomics <i>Vadim Yanshole, ITC SB RAS / NSU, Russian Federation</i>	
1:30 p.m. – 3:15 p.m.	Session 18: Food Applications 2 (Session Chairs: <i>Sastia Putri & Johanna Jokioja</i>)	
1:30 p.m. – 2:00 p.m.	18A SESSION KEYNOTE Rapid Evaporative Ionisation Mass Spectrometry for detecting compounds related to consumer liking of meat <i>Alastair Ross, AgResearch, New Zealand</i>	King Willem Alexander
2:00 p.m. – 2:20 p.m.	18B Fiber mix supplementation in wheat-based flatbreads delays the exogenous appearance of glucose and its downstream metabolites <i>Lisa Schlicker, BRICS, TU Braunschweig, Germany</i>	
2:20 p.m. – 2:35 p.m.	18C* A metabolomic approach to the identification and validation of biomarkers of apple intake <i>Aoife McNamara, Institute of Food and Health, Ireland</i>	
2:35 p.m. – 2:55 p.m.	18D Brainfood metabolomics study (Icebreaker, Brave sub-study) <i>Kati Hanhineva, Afekta Technologies Ltd., Finland</i>	
2:55 p.m. – 3:10 p.m.	18E* Mass spectrometry based non-targeted metabolomics enables the identification of Amadori products in feces of formula-fed infants <i>Alesia Walker, Helmholtz Zentrum München, Research Unit Analytical BioGeochemistry, Germany</i>	

Wednesday, June 26

Time	Session	Location
1:30 p.m. – 3:15 p.m.	Session 19: Data Integration & Data Basing 2 (Session Chairs: Dan Raftery & Dara Daygon)	
1:30 p.m. – 2:00 p.m.	19A SESSION KEYNOTE The UniProt, Rhea, and SwissLipids knowledge resources for metabolomics and lipidomics <i>Alan Bridge, Swiss-Prot group, SIB Swiss Institute of Bioinformatics, Switzerland</i>	Princess Alexia
2:00 p.m. – 2:20 p.m.	19B RIKEN Plant Metabolome MetaDatabase: an integrated plant metabolome data repository based on the semantic web <i>Atsushi Fukushima, RIKEN Center for Sustainable Resource Science, Japan</i>	
2:20 p.m. – 2:35 p.m.	19C Contextualizing Metabolomics Data by Integrating Text Mining and Machine Learning <i>Magnus Palmblad, Leiden University Medical Center, Netherlands</i>	
2:35 p.m. – 2:55 p.m.	19D Metabolic Reaction Network-based Recursive Metabolite Annotation for Untargeted Metabolomics <i>Zheng-Jiang Zhu, Chinese Academy of Sciences (CAS), China</i>	
2:55 p.m. – 3:10 p.m.	19E Sparse Multi-block PLS for Selection of Related Signals in Multi-platform Metabolomics Data <i>Timothy Ebbels, Imperial College London, United Kingdom</i>	
1:30 p.m. – 3:15 p.m.	Session 20: Regulatory Session Translating metabolomics from academic science into regulatory practice: challenges and progress in pharmacology and toxicology (Session Chair: Pim Leonards)	
1:30 p.m. – 1:50 p.m.	An Overview of the 21st Century Cures Act: Opportunities for Biomarkers and Precision Medicine <i>Dr. Rick Beger, National Center for Toxicological Research, US FDA, United States</i>	Princess Ariane
1:50 p.m. – 2:10 p.m.	Big Data “Omics” – Challenges and Opportunities in Regulation <i>Dr. Renate König, Paul-Ehrlich-Institut, Federal Institute for Vaccines and Biomedicines, Germany</i>	
2:10 p.m. – 2:30 p.m.	Various Applications of Metabolomics in Toxicology <i>Prof Hennie Kamp, BASF, Germany</i>	
2:30 p.m. – 2:50 p.m.	Best Practice Guidelines and Reporting Standards for Applications of Metabolomics in Regulatory Toxicology: the International MERIT Project <i>Prof. Mark Viant, University of Birmingham, UK</i>	
2:50 p.m. – 3:15 p.m.	Panel Discussion with all presenters, moderated by Pim Leonards	

*AWARD WINNERS

Wednesday, June 26

Time	Session	Location
3:45 p.m. – 5:30 p.m.	Session 21: Infection and Immunity (Session Chairs: Darren Creek & Xinzhu Wang)	
3:45 p.m. – 4:15 p.m.	21A SESSION KEYNOTE Understanding the metabolic perturbation in severe fever with thrombocytopenia syndrome <i>Zeping Hu, Tsinghua University, China</i>	Atlantic
4:15 p.m. – 4:35 p.m.	21B Hijacking of host cellular function by opportunistic pathogens during intracellular infections <i>Volker Behrends, University of Roehampton, United Kingdom</i>	
4:35 p.m. – 4:50 p.m.	21C* Diagnosis of Aspergillosis based on metabolomics data – application of relative expression analysis-RXA to minimize interindividual metabolic differences <i>Joanna Godzien, Medical University of Bialystok, Clinical Research Centre, Poland</i>	
4:50 p.m. – 5:10 p.m.	21D Alterations of systematic metabolism upon viral infection: from metabolite profiling to isotope tracing. <i>Kristaps Klavins, CeMM - Research Center for Molecular Medicine, Austria</i>	
5:10 p.m. – 5:25 p.m.	21E* Characterisation of the human milk lipidome using liquid chromatography-ion mobility spectroscopy-mass spectrometry reveals novel lipids <i>Alexandra George, The University of Western Australia, School of Molecular Sciences, Australia</i>	
3:45 p.m. – 5:30 p.m.	Session 22: Environment & Toxicology (Session Chairs: Melissa Fitzgerald and Ralf Weber)	
3:45 p.m. – 4:15 p.m.	22A SESSION KEYNOTE Metabolomics and toxicity studies of zebrafish embryos exposed to environmental contaminants <i>Pim Leonards, Vrije Universiteit, Netherlands</i>	King Willem Alexander
4:15 p.m. – 4:35 p.m.	22B Gestational and post-natal exposure to ambient air pollution: a metabolomics approach <i>Marina Tavares, University of Sao Paulo, Brazil</i>	
4:35 p.m. – 4:50 p.m.	22C Association between coffee consumption, plasma metabolites and hypertension risk <i>Wei Jie Seow, National University of Singapore, Singapore</i>	
4:50 p.m. – 5:10 p.m.	22D Small molecules and lipids preserved over thousands of years in mummified humans <i>Kevin Quinn, University of Colorado Anschutz, United States</i>	
5:10 p.m. – 5:25 p.m.	22E Food preservative zinc oxide nanoparticles (ZnONPs) induces neurobehavioral deficits in mice with altered gut microbiome and dysregulated serum and hippocampus metabolome <i>Chang Chen, Chongqing Medical University, China</i>	

***AWARD WINNERS**

Wednesday, June 26		
Time	Session	Location
3:45 p.m. – 5:30 p.m.	Session 23: New Instrumentation (Session Chairs: Rawi Ramautar & Charmion Cruickshank-Quinn)	
3:45 p.m. – 4:15 p.m.	23A SESSION KEYNOTE Integration of MS imaging and LC-MS techniques – discovering tissue heterogeneity with the accuracy and depth of traditional profiling tools Matthew Lewis, Imperial College London, United Kingdom	Princess Alexia
4:15 p.m. – 4:35 p.m.	23B Mapping Hydrophilic Intermediates of the Central Carbon and Energy Metabolism (CCEM) by Ion Pairing and Ion Exchange Chromatography – Tandem MS Gerd Balcke, Leibniz-Institute of Plant Biochemistry, Germany	
4:35 p.m. – 4:50 p.m.	23C* High-Resolution μ-scaled Magic-Angle Spinning NMR Mapping of diseased rat brains Covadonga Lucas-Torres, CEA Saclay, France	
4:50 p.m. – 5:10 p.m.	23D Electroextraction-based sample preparation strategies for enrichment of low-abundant metabolites Peter Lindenburg, Analytical Biosciences & Metabolomics, Leiden Centre for Applied Bioscience, Leiden University, Netherlands	
5:10 p.m. – 5:25 p.m.	23E How to automate boring lipidomic extraction? Justine Bertrand-Michel, MetaToul-Lipidomic, France	
3:45 p.m. – 5:30 p.m.	Session 24: Novel Applications (Session Chairs: Lorraine Brennan & Bert Wouters)	
3:45 p.m. – 4:15 p.m.	24A SESSION KEYNOTE Metabolomics: Central Approach for Study of the Exposome Dean Jones, Emory University, United States	Princess Ariane
4:15 p.m. – 4:35 p.m.	24B Metabolomes of mitochondrial diseases and inclusion body myositis patients: treatment targets and biomarkers Vidya Velagapudi, FIMM, University of Helsinki, Finland	
4:35 p.m. – 4:50 p.m.	24C Boosting anti-doping screening through metabolomics Luca Narduzzi, Oniris - Laberca, France	
4:50 p.m. – 5:10 p.m.	24D A Novel Metabolomics Method for Quantitative Analysis of over 1,000 Metabolites Guowang Xu, Dalian Institute of Chemical Physics, Chinese Academy of Sciences, China	
5:10 p.m. – 5:25 p.m.	24E Post-mortem changes in the metabolomic profiles of animal and human tissues Ekaterina Zelentsova, International Tomography Center SB RAS, Novosibirsk, Russia; Novosibirsk State University, Russian Federation	

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***AWARD WINNERS**

Thursday, June 27		
Time	Session	Location
8:45 a.m. – 10:00 a.m.	Session 25: Respiratory Diseases (Session Chairs: Nichole Reisdorph & Priyanka Choudhury)	
8:45 a.m. – 9:15 a.m.	25A SESSION KEYNOTE Omega-6 polyunsaturated fatty acid metabolites are associated with vitamin D levels in early life <i>Jessica Lasky-Su, Brigham and Women's Hospital, United States</i>	Atlantic
9:15 a.m. – 9:40 a.m.	25B Comprehensive metabolomic and immunological profiling of Asthma COPD Overlap (ACO) <i>Nilanjana Ghosh, School of Medical Science and Technology, India</i>	
9:40 a.m. – 9:55 a.m.	25C* The relationship between the maternal metabolome and childhood asthma: results from the Vitamin D Antenatal Asthma Reduction Trial (VDAART) <i>Mengna Huang, Channing Division of Network Medicine, Brigham and Women's Hospital and Harvard Medical School, United States</i>	
8:45 a.m. – 10:00 a.m.	Session 26: Model Organisms (Session Chairs: Oliver Jones & Lyudmila Yanshole)	
8:45 a.m. – 9:15 a.m.	26A SESSION KEYNOTE Can Stage-Specific Metabolites be Recovered from Mixed-Stage <i>C. elegans</i> Cultures? <i>Arthur Edison, University of Georgia, United States</i>	Princess Alexia
9:15 a.m. – 9:40 a.m.	26B Bringing together what belongs together – Bridging the <i>Caenorhabditis elegans</i> model organism database, genome scale model and metabolite databases <i>Michael Witting, Research Unit Analytical BioGeoChemistry, Helmholtz Zentrum München, Germany</i>	
9:40 a.m. – 9:55 a.m.	26C* The use of mitochondrial metabolomics in discovering the molecular function of a mitochondrial membrane protein <i>Daqiang Pan, Centre for Biological Systems Analysis, University of Freiburg, Germany</i>	
8:45 a.m. – 10:00 a.m.	Session 27: Genome-Scale Modeling (Session Chairs: Gabi Kastenmüller & Patrick Trainor)	
8:45 a.m. – 9:15 a.m.	27A SESSION KEYNOTE Computational modelling of host-microbiome co-metabolism <i>Ines Thiele, National University of Ireland, Ireland</i>	Princess Ariane
9:15 a.m. – 9:40 a.m.	27B Metabolic footprint of Parkinson's Disease: integration of patient-derived X-omics data with a human genome-scale model. <i>Agnieszka Wegrzyn, LACDR, Leiden University, Netherlands</i>	
9:40 a.m. – 9:55 a.m.	27C Genome-wide metabolic modeling of human CD4+ T-helper cells differentiation unraveled the relative importance of ceramides <i>Partho Sen, Postdoctoral Researcher, Finland</i>	

***AWARD WINNERS**

Thursday, June 27		
Time	Session	Location
10:45 a.m. – 12:00 p.m.	Session 28: Microbiome (Session Chairs: Guowang Xu & Madeleine Ernst)	
10:45 a.m. – 11:15 a.m.	28A SESSION KEYNOTE Global Chemical Impacts of the Microbiome Include Novel Conjugated Bile Acids that Stimulate FXR <i>Robert Quinn, Michigan State University, United States</i>	Atlantic
11:15 a.m. – 11:40 a.m.	28B* Exploring the microbiota-host epigenetics axis in female and male germ-free and conventional mice <i>Joan Miró Blanch, Metabolomics Platform, IISPV & Department of Electronic Engineering, Universitat Rovira i Virgili, Spain</i>	
11:40 a.m. – 11:55 a.m.	28C* Individual Variations in Plasma Metabolites are Driven by Diet, Genetics and Gut Microbiome <i>Lianmin Chen, University Medical Centre Groningen, Netherlands</i>	
10:45 a.m. – 12:00 p.m.	Session 29: Plant Applications 2 (Session Chairs: Lloyd Sumner & Adam Jozwiak)	
10:45 a.m. – 11:15 a.m.	29A SESSION KEYNOTE A novel plant-microbiome co-culturing system reveals key associations of specific metabolites with plant growth and stress tolerance <i>Sanjay Swarup, National University of Singapore, Singapore</i>	Princess Alexia
11:15 a.m. – 11:40 a.m.	29B An untargeted LC-MS based work-flow for the structural characterization of biological polyesters <i>Rebecca Dauwe, Université de Picardie Jules Verne, France</i>	
11:40 a.m. – 11:55 a.m.	29C NMR-MS metabolomics reveals sulfonation in the Salicaceae <i>Clarice Noletto-Dias, Rothamsted Research, United Kingdom</i>	
10:45 a.m. – 12:00 p.m.	Session 30: Metabolite Identification 2 (Session Chairs: Fidele Tugizimana & Christina Jones)	
10:45 a.m. – 11:15 a.m.	30A SESSION KEYNOTE Correlation-based deconvolution (CorrDec) method for data-independent acquisition mass spectrometry <i>Ipputa Tada, Department of Genetics, SOKENDAI (Graduate University for Advanced Studies), Japan</i>	Princess Ariane
11:15 a.m. – 11:40 a.m.	30B Evaluation of molecular ionization propensities in different ionization modes: providing evidence for the presence of small molecules in synthetic blinded samples <i>Jamie Nunez, Pacific Northwest National Laboratory, United States</i>	
11:40 a.m. – 11:55 a.m.	30C* Improved oxylipins and fatty acids identification by using LC-DTIM-MS <i>Sonia Liggi, Department of Biochemistry and Cambridge Systems Biology Centre, University of Cambridge, United Kingdom</i>	
1:30 p.m. – 3:30 p.m.	Plenary Session 4 and Closing Ceremony <i>Jean-Charles Portais, University Toulouse, France</i>	