## PROGRAM

## 10th Theodor Escherich Symposium GRAZ AND FRIENDS!

Time	Title	Speaker	Chairs
08:15-09:00 09:00-09:15	Registration and setup Welcome note: 10 years Theodor Escherich Symposium	Caroline Schober	
09:15-09:50	SESSION 1 (Uni Graz): Keynote Talk		Christine
	Commensal Genotoxins: Microbial Metabolites Damage DNA	Ellen Zechner	Moissl-Eichinger
09:50-10:30	SESSION 2 (Med Uni Graz): Microbiome & Infection		1
	Antigen discovery and spatial microbiome sequencing tools dive deep in the mucosa Abstract 1: Virulome analysis of Enterobacteriaceae from colorectal cancer patients at a tertiary hospital in Ghana	Martin Strazar Vincent Amarh	Martin Zacharias &
10:30-11:15	Abstract 2: Role of Methanogenic Archaea in Gastrointestinal Disorders Coffee break/Industry exhibition	Christina Kumpitsch	Maria Durdevic
11:15-12:45	SESSION 3a (TU Graz): Plant/Exposome/Food Axis		1
	The central role of the soil microbiome in One Health policy Harnessing the potential of the plant microbiota to control prevalent phytopathogens Soil and plant microbiomes under changing environment The Agri Microbiomes in a Food Security Context Abstract 1: Indigenous seed endophytic bacteria of millets improve seedlings developments and protect from fungal diseases	Ahmed Abdelfattah Tomislav Cernava Magdalena Frąc Lise Korsten Satish Kumar Verma	Nina Bziuk & Isabella Kögl
12:45-13:45	Abstract 2: The Ecological Niche of Entomopathogenic Fungi: Global and Local Aspects Lunch break	Adrian Wolfgang	
13:45-14:20	SESSION 3b (TU Graz): Plant/Exposome/Food Axis		1
	Plant growth promoting or detrimental: The role of of the flagellar transcriptional regulator flhC of Acidovorax for plant-microbe interactions The edible plant microbiome and its applications to develop medical foods for women	Michael Schloter Gerardo V Toledo	Valeria Castaldi & Matevz Zlatnar
	in menopause and people with rheumatoid arthritis		ł

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	14:20-15:30	SESSION 4 (VAAM): Innovative Methods in Microbiome research		1
	15:30-16:00	Expanding the microbiome toolkit in genomics and microscopy Exploring links between the human gut microbiota and health using large multi-omics population cohorts Abstract 1: Development of a genome-scale metabolic model for the murine gut microbiome (McMurGut) Abstract 2: Ecological differentiation of organic acid utilizers in the animal gut microbiome Abstract 3: Genome-wide sweeps are fundamental adaptive units in the human gut microbiome Coffee break/Industry exhibition	Alexander Probst Guillaume Meric Torben Kühnast Cameron Strachan Xiaoqian Yu	Birgit Wasserman & Alexander Mahnert
	16:00-17:20	Session 5 (BioTechMed): Microbiome Dynamics		1
	17:20-19:00	Abstract 1: Age-Related Dynamics of Methanogenic Archaea in the Human Gut Microbiome: Implications for Longevity and Health Abstract 2: Microbiome research tools beyond 16S rRNA and shotgun sequencing Abstract 3: Decoding FMT Dynamics in UC: Early Assessment of Functional, Metabolic, and Taxonomic Predictors The intra- and peritumoral microbiome associated with treatment responses in pancreatic and colon cancer Parsing and interpreting the various dynamical regimes of the human gut microbiota Postersession/Industry exhibition	Rokhsareh Mohammadzadeh Christa Nöhammer Marija Durdevic Christoph Stein-Thöringer Sean Gibbons	Christina Kumpitsch & Alexander Mahnert
	19:00	Evening event: Mixer: Music, Party for all participants	ÖH BigBand and DJ	
	08:15-09:00	Registration and setup Session 6 (ÖGHMP): Microbial Interplay and Vesicles		•
	09.00-10.00	The development of the gut-brain axis in premature neonates vesicles Immunoglobulin A coated bacterial membrane vesicles drive pro-inflammatory responses in ulcerative colitis Gut bacteria nanovesicles: their potential role in mediating cross-kingdom communication within and beyond the gut	David Berry Himadri B. Thapa Regis Stentz	Roksareh Mohammadzadeh & Torben Kühnast
	10:00-11:00	Session 7 (CoE Microplanet): We're living on a micro-planet! Next generation chemical imaging of the gut microbiome-drug interaction network Understanding emergent phenomena of microbial decomposer communities using individual-based modelling Personalized engraftment prediction with metabolic models of the human gut microbiota	Michael Wagner Christina Kaiser Christian Diener	Charlotte Neumann & Tejus Shinde

19.1.2024

Time	Title	Speaker	Chairs
11:00-11:30	Coffee break/Industry exhibition		
11:30-13:00	Session 8 (ÖGGH, ÖGPATH and ÖGIT): Pathogens and Solutions		1
13:00-14:00	C. difficile: Breaking the Cycle of Spore to Vegetative to Spore in 2024 (online) The gut-lung microbiome: a new organ to ICU-revive? Abstract 1: Exploring microbial competition and warfare to therapeutically utilize the microbiome Abstract 2: Identification of the bacterial metabolite aerugine as potential trigger of human dopaminergic neurodegeneration Lunch break	Paul Feuerstadt Sebastian Imbert Lisa Osbelt Nathalie Christine Wörz	Carolina Lobato & Jasper Schierstaedt
14:00-16:00	Session 9 (AMICI): Microbiomes and Human Surfaces		1
	Abstract 1: Microbiota-associated metabolites: methylphenols and the blood-brain barrier Abstract 2: Widespread gut commensals of the family Sutterellaceae perform fumarate and nitrate respiration Abstract 3: Nanomechanical mechanisms of <i>Borrelia</i> interactions with extracellular matrix Abstract 4: Immediate targeting of host ribosomes by jumbo phage encoded proteins Abstract 5: iNKT cells immunomodulation and mucosal healing by microbiota-derived lactate Abstract 6: <i>Methanobrevibacter smithii</i> and <i>Cand</i> . M. intestini - Similar but different Taste the difference: cell biology of a multicellular bacterium inhabiting the human mouth T cell regulation by bacterial metabolites 10 years of the human skin archaeome	Lesley Hoyles Nataliia Solntseva Yoojin Oh Milan Gerovac Francesco Strati Viktoria Weinberger Silvia Bulgheresi Clarissa Campbell Alexander Mahnert	Kristina Michl & Adrian Wolfgang
16:00-16:10	Awards and Goodbye		
16:10	Goodbye reception		



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