

THE PROGRAM

We are delighted to welcome a fantastic panel of renowned speakers in order to make ICCMg3 a great scientific experience again. The list is provisional and should be completed soon.

THURSDAY, OCTOBER 18

Day	Session	Start	Stop	Talk	Speaker
Thursday 18	Welcome speech	10h45	11h	Welcome speech: welcome to ICCMg3!	Jacques Schrenzel
Thursday 18	Bioinformatics	11h	11h30	A year in clinical bioinformatics	Helena Seth Smith
Thursday 18	Bioinformatics	11h30	11h45	Selected abstract: Extensive viral nucleic acid contamination in clinical samples	Speaker abstract 37
Thursday 18	Bioinformatics	11h45	12h	Selected abstract: Culture-Free Metagenomic Deconvolution, Plasmid-Host Association, and Strain Genome Separation Using Hi-C Proximity-Ligation Technology	Speaker abstract 8
Thursday 18	Bioinformatics	12h	12h30	What's new with Nanopore sequencing	Liana Kafetzopoulos
Thursday 18	Lunch break	12h30	14h		
Thursday 18	Microbiota and personalized medicine	14h	14h30	Personalizing treatments using microbiome and clinical data	Eran Segal
Thursday 18	Microbiota and personalized medicine	14h30	15h	Microbiota and cancer	Bertrand Routy
Thursday 18	Microbiota and personalized medicine	15h	15h30	Now moving to the vaginal microbiota and personalized medicine?	Jacques Ravel
Thursday 18	Microbiota and personalized medicine	15h30	15h45	Selected abstract: Caesarean section stunts the gut microbiota and predisposes pathogen colonisation during very early life	Speaker abstract 41
Thursday 18	Coffee break	15h45	16h30		
Thursday 18	Clinical metagenomics	16h30	17h	Free-circulating DNA in clinical samples	Silke Grumaz
Thursday 18	Clinical metagenomics	17h	17h30	Cell-free metagenomics	Wei Gu
Thursday 18	Clinical metagenomics	17h30	17h45	Selected abstract: Microbiological Diagnostic Performance of Metagenomic Next-Generation Sequencing when Applied to Clinical Practice	Speaker abstract 15
Thursday 18	Clinical metagenomics	17h45	17h55	ICCMg prizes	TBD
Thursday 18	Prizes	17h55	18h10	Selected abstract: ICCMg3 award lecture: Investigation of viruses in human cancers by application of multiple pre-sequencing enrichment methods	Speaker abstract 36
Thursday 18	Poster session	18h10	19h	Poster viewing and discussion around a welcome drink	
Thursday 18	Walk	19h	19h15	Walk to the Vieux Bois restaurant.	
Thursday 18	Dinner	19h15	23h	Dinner at the Vieux Bois restaurant.	

FRIDAY, OCTOBER 19

Day	Session	Start	Stop	Talk	Speaker
Friday 19	Clinical genomics	9h	9h30	Practical issues	John Rossen
Friday 19	Clinical genomics	9h30	10h	Challenges and opportunities for whole-genome sequencing–based surveillance of antibiotic resistance	Willem van Schaik
Friday 19	Clinical genomics	10h	10h30	How can WGS help in outbreaks? The point of view of the infection prevention and control	Stephan Harbarth
Friday 19	Coffee break	10h30	11h15		
Friday 19	Beyond clinical metagenomics	11h15	11h45	Lost in translation? Considering user needs in reporting clinical microbial genomics results.	Jennifer Gardy
Friday 19	Beyond clinical metagenomics	11h45	12h15	Ethics in clinical metagenomics	Beatrice de Montera
Friday 19	Lunch break	12h15	13h45		
Friday 19	Clinical metagenomics	13h45	14h15	Emerging pathogens: new insights with WGS	Richard Neher
Friday 19	Clinical metagenomics	14h15	14h45	Clinical genomics in tuberculosis	Sébastien Gagneux
Friday 19	Clinical metagenomics	14h45	15h	Selected abstract: Diversity and composition of the microbiome in the lower respiratory tract is associated with lung damage in tuberculosis patients from Tanzania	Speaker abstract 39
Friday 19	Conclusive speech	15h	15h15	Conclusive speech: see you in ICCMg4!	Etienne Ruppé