

Lausanne Genomics Days 2020

Dorigny Campus, Génopode Building, Auditorium C

Attendance is free but [registration](#) required

Monday February 10, 2020

Tuesday February 11, 2020

9:30-10:00	Welcome and coffee in the Génopode hall	9:30-10:00	Welcome and coffee in the Génopode hall
10:00-11:45	TRANSLATIONAL REGULATION <i>Chair Prof. Sven Bergmann</i> - UNIL (Department of Computational Biology)	10:00-11:45	MICROBIOMES <i>Chair Prof. Philipp Engel</i> - UNIL (Department of Fundamental microbiology)
	10:00 Keynote presentation <i>Prof. Tamir Tuller</i> - Tel Aviv University, Israel Computational deciphering and modelling of novel gene expression rules encoded in the mRNA		10:00 Keynote presentation <i>Prof. Colin Hill</i> - University College Cork, Ireland A voyage through the gut virome
	10:45 <i>Dr. Anneke Brümmer (Prof. Sven Bergmann)</i> - UNIL (Department of Computational Biology) LncRNA sequences are biased to counteract their translation		10:45 <i>Dr. Kirsten Ellegaard (Prof. Philipp Engel)</i> - UNIL (Department of Fundamental Microbiology) Getting quantitative on the Microbiome: vast differences in strain-level diversity among host-associated bacterial communities
	11:05 <i>Dr. Cedric Gobet (Prof. Felix Naef)</i> - EPFL Modeling ribosome dwell times and relationships with aminoacyl-tRNAs in mouse liver		11:05 <i>Paddy Gibson (Prof. Jan-Willem Veenking)</i> - UNIL (Department of Fundamental Microbiology) Deciphering the origins of amoxicillin resistance in <i>Streptococcus pneumoniae</i> through experimental evolution and NGS
	11:25 <i>Prof. David Gatfield</i> - UNIL (Center for Integrative Genomics) Transcriptome-wide sites of collided ribosomes reveal sequence determinants of translational pausing		11:25 <i>Dr. Guillermo Salazar Guiral</i> - ETH (Microbiology) Disentangling the effects of gene expression changes and community turnover in the global ocean
11:45-13:00	Lunch in the Génopode hall	11:45-13:30	Lunch in the Génopode hall
13:00-14:45	RNA PROCESSING <i>Chair Dr. Julien Marquis</i> - UNIL (Genomic Technologies Facility)	13:30-15:15	EVOLUTION <i>Chair Prof. Tanja Schwander</i> - UNIL (Department of Ecology and Evolution)
	13:00 Keynote presentation <i>Prof. Stefan Ameres</i> - Institute of Molecular Biotechnology, Austria SLAMseq – Dissecting gene regulatory mechanisms by time-resolved RNA sequencing		13:30 <i>Prof. Anne Roulin</i> - UZH (Department of Plant and Microbial Biology) Transposable element evolution in small genomes What can we learn from population genomics in <i>B. distachyon</i> ?
	13:45 <i>Dr. Flora Brozzi (Prof. Romano Regazzi)</i> - UNIL (Department of Fundamental Neurosciences) RNA-tagging to unveil beta-cell/immune cell crosstalk in diabetes pathogenesis		13:50 <i>Jelisaveta Djordjevic (Prof. Tanja Schwander)</i> - UNIL (Department of Ecology and Evolution) Sexual conflict and the ontogeny of sex-biased gene expression in stick insects
	14:05 <i>Dr. Rodrigo Siqueira Reis (Prof. Yves Poirier)</i> - UNIL (Department of Plant Molecular Biology) Complex mRNA-antisense lncRNA interaction revealed by SHAPE-MaP		14:10 <i>Tomas Kay (Prof. Laurent Keller)</i> - UNIL (Department of Ecology and Evolution) Position in the social network is a central driver of behavior and gene expression in ants
	14:25 <i>Lina Worpenberg (Prof. Jean-Yves Roignant)</i> - UNIL (Center for Integrative Genomics) Shaping the nervous system by m6A mRNA modification		14:30 Keynote presentation <i>Prof. Detlev Arendt</i> - EMBL Heidelberg Evolution of neurons and of the nervous system
14:45-15:15	Coffee break in the Génopode hall		
15:15-16:55	GENE EDITING <i>Chair Prof. Julia Santiago Cuellar</i> - UNIL (Department of Plant Molecular Biology)		
	15:15 Keynote presentation <i>Prof. Martin Jinek</i> - University of Zurich, Switzerland Molecular mechanisms of genome editors		
	16:00 <i>Dr. Robertas Ursache (Prof. Niko Geldner)</i> - UNIL (Department of Plant Molecular Biology) CRISPR and the new age of genetic analysis in <i>Arabidopsis</i>		
	16:25 <i>Dr. Lotte Spel (Prof. Fabio Martinon)</i> - UNIL (Department of Biochemistry) A functional genetics approach to dissect inflammasome activation		
16:50	Apero in the Génopode hall		

Organization committee: Sven Bergmann, Philipp Engel,
Julien Marquis, Julia Santiago Cuellar and Tanja Schwander

Link to registration



Sponsorship

