

Workshop Program Day 1 (March 13)

11.00 - 13.00 Workshops 1 - 4

WS 1 Potassium channels: molecular biology, pathophysiology, and computational aspects

Room: Sydney

- 11.00 – 11.05 **Alex J. Baertschi**, CMU, University of Geneva, Switzerland
Introduction
- 11.05 – 11.35 **Ivet Bahar**, School of Medicine, University of Pittsburgh, USA
Gating mechanisms in membrane protein: computational study of the dynamics of potassium channels and glutamate transporters
- 11.35 – 12.00 **Richard Warth**, University of Regensburg, Germany
Pathophysiology of epithelial potassium channels in the gastrointestinal tract
- 12.00 – 12.20 **Hugues Abriel**, University of Lausanne
IKCNQ1 potassium channel is down-regulated by ubiquitylating enzymes of the Nedd4/Nedd4-like family
- 12.20 – 12.45 **Marcus C. Schaub**, University of Zurich, Switzerland
Role of cardiac KATP channels in calcium handling and metabolism
- 12.45 – 13.00 General Discussion and Conclusion

WS 2 Computational Biology:

Room: Montreal

- 11.00 – 11.40 **David Harel**, Weizmann Institute of Science, Rehovot, Israel
In Silico Biology, or On Comprehensive and Realistic Modeling
- 11.40 – 12.20 **Shalev Itzkovitz**, Weizmann Institute of Science, Rehovot, Israel
Using genomic variability to reconstruct cell lineage
- 12.20 – 13.00 **Alfio Quarteroni**, EPFL, Lausanne, Switzerland
On the mathematical modeling of the human cardiovascular system
- 13.00 – 13.10 **Christian von Mering**, University of Zurich, Switzerland
Phylogenetic Assessment of microbial communities through environmental sequence data
- 13.10 – 13.20 **Sven Bergmann**, SIB & University of Lausanne, Switzerland
Pre-steady-state decoding of the bicoid morphogen gradient
- 13.20 – 13.30 **Marco Scarsi**, SIB & Biozentrum University of Basel, Switzerland
A Combined Virtual Screening and Biological Assay Approach

WS 3 Stem cells and cancer:

Room: Singapore

- 11.00 – 11.15 **Stephen Ryser**, Geneva University Hospital, Switzerland
Defining the niche of male germinal stem cell in new born rat by expression profiling
- 11.15 – 11.40 **Neil J. Harrison**, University of Sheffield, UK
Culture adaptation of embryonic stem cells: models for germ cell tumour progression
- 11.40 – 12.05 **Cathrin Brisken**, ISREC, Lausanne, Switzerland
Hormones and paracrine signaling in breast development and breast cancer
- 12.05 – 12.20 **Martin Buess**, University of Basel, Switzerland
In vitro analysis of heterotypic cell-cell interaction effects to identify the contribution of tumor-stroma interaction on global gene expression profiles in breast cancer
- 12.20 – 12.45 **Andreas Trumpp**, ISREC, Lausanne, Switzerland
Cooperative control of hematopoietic stem cell function by c-Myc and N-Myc.
- 12.45 – 13.00 **Damaris Bausch-Fluck**, Swiss Federal Institute of Technology (ETH) Zurich, Switzerland
Cell Surface Capturing as a method of choice for the identification and quantification of relevant plasma membrane antigens of cancer stem cells.

WS 4 Statistics in animal research

Room: Rio

- 11.00 – 11.45 **Michael F.W. Festing**, NC3Rs, London, UK
We need to improve the design and statistical analysis of animal experiments.
- 11.45 – 12.05 **Robert Greif**, Inselspital Bern, Switzerland
Missing data: what can I do with it?
- 12.05 – 12.25 **Javier Fandino**, Inselspital Bern, Switzerland
So what? How do I define relevant outcome parameters in experimental research?
- 12.25 – 12.45 **Eva Waiblinger**, Tierversuchskommission Basel, Universität Zürich, Switzerland
Statistics and relevance – How can I convince the local animal welfare committee?
- 12.45 - 13.00 Discussion

Workshop Program Day 2 (March 14)

10.30 - 12.30 Workshops 5 - 7

WS 5 Small RNAs

Room: Sydney

- 10.30 – 11.00 **Mihaela Zavolan**, Biozentrum University of Basel, Switzerland
Computational prediction of ncRNA genes and targets
- 11.00 – 11.30 **Elisa Izaurralde**, Max Planck-Institute for Developmental Biology, Tübingen, Germany
Mechanisms of miRNA-mediated gene silencing
- 11.30 – 11.45 **Xavier Ding**, Friedrich Miescher Institute, Basel Switzerland
Let-7 microRNA mode of action in vivo: translational effect and role of eif-3
- 11.45 – 12.15 **Markus Stoffel**, Institute for Molecular Systems Biology, ETH Zurich, Switzerland
Approaches to study miRNA function in mammals
- 12.15 – 12.30 **Jurgi Camblong**, University of Geneva, Switzerland
*Anti-sense RNA stabilization in *S. cerevisiae* induces gene silencing by Hda1/2/3 histone deacetylase complex*

WS 6 Imaging and image analysis

Room: Rio

- 10.30 – 11.00 **Thomas Vetter**, University of Basel, Switzerland
Statistical shape models for the analysis of human morphology
- 11.00 – 11.30 **Jean-Yves Chatton**, University of Lausanne, Switzerland
Image analysis and processing for quantitative cellular imaging
- 11.30 – 12.00 **Graham Knott**, University of Lausanne, Switzerland
Exploring brain circuits with serial section electron microscopy; current technology and future directions
- 12.00 – 12.15 **Matthias Ochs**, University of Bern, Switzerland
Quantitation of lung structure by stereology: Leonhard Euler and the number of alveoli
- 12.15 – 12.30 **Tilman Vogt**, University of Fribourg, Switzerland
The three dimensional reconstruction of motor endplates of the vertebrate skeletal muscle fibres.

WS 7 Infectious Diseases

Room: Singapore

- 10.30 – 10.50 **Isabel Sorg**, Biozentrum University of Basel, Switzerland
*The *Yersinia* basal body protein YscU plays a role in recognition of Type III export*
- 10.50 – 11.10 **Nicolas Maire**, Swiss Tropical Institute, Basel, Switzerland
Stochastic simulation of Malaria epidemiology and control
- 11.10 – 11.30 **Laurence Neff**, Geneva University Hospital, Switzerland
*Molecular characterization and subcellular localization of macrophage infectivity potentiator, a *Chlamydia trachomatis* lipoprotein*
- 11.30 – 11.50 **Christian L. Althaus**, Utrecht University, The Netherlands
Modeling CTL evasion during SIV/HIV infection. Is there a great escape?

- 11.50 – 12.10 **Sabine Kuettel**, University of Geneva, Switzerland
Adenosine kinase and glyceraldehyde-3-phosphate dehydrogenase identified as putative CD12001 target(S) in Trypanosoma brucei rhodesiense
- 12.10 – 12.30 **Christian J. Burckhardt**, Institute of Zoology, University of Zürich, Zürich, Switzerland,
Computational dissection of adenovirus cell surface motion reveals receptor mediated virus drifting on filopodia

15.30 - 17.30 Workshops 8 - 11

WS 8 Cells to Organs

Room: Singapore

- 15.30 – 16.00 **Lilianna Solnica-Krezel**, Vanderbilt University, Nashville, USA
Apelin and its Receptor Control Heart Field Formation during Zebrafish Gastrulation
- 16.00 – 16.30 **Bernhard Herrmann**, Max-Planck-Institute for Molecular Genetics, Berlin, Germany
Towards systems biology of organogenesis: deciphering regulatory networks controlling mesoderm development in the mouse
- 16.30 – 17.00 **Andy McMahon**, Harvard University, Cambridge, MA, USA
From precursor to product: organogenesis of the mammalian kidney.
- 17.00 – 17.15 **Anne Grapin-Botton**, ISREC/EPFL, Epalinges/Lausanne, Switzerland
Waves of endocrine-cell differentiation in the pancreas
- 17.15 – 17.30 **Johannes Jaeger**, University of Cambridge, UK
Dynamics control of positional information by the Drosophila gap gene network

WS 9 Novel approaches in anti-inflammatory drug therapy

Room: Montreal

- 15.30 – 16.00 **Peter Mertens**, University of Aachen, Germany
From inflammation to fibrosis: the Y-box protein-1 as extracellular mediator of fibrogenesis
- 16.00 – 16.30 **Jürgen Stein**, University Hospital Frankfurt am Main, Germany
Anti-inflammatory mechanisms of histone deacetylase inhibitors
- 16.30 – 16.45 **Bruno Schnyder**, CNRS, Molecular Immunology and Embryology, Orléans, France
Interleukin-17 is a negative regulator of established allergic asthma
- 16.45 – 17.00 **Frauke Döll**, University of Bern, Switzerland
Hypoxia regulates the sphingosine kinase-1 activity and expression in the endothelial cell line EA.hy926
- 17.00 – 17.30 **Josef Pfeilschifter**, University of Frankfurt am Main, Germany
Sphingolipid signaling in inflammation

WS 10 Computation in heart disease

Room Rio

- 15.30 – 15.50 **Pascal Giat**, Europe GE Healthcare
Cardiac CT image reconstruction
- 15.55 – 16.15 **Francesco Faletta**, Cardiocentro Ticino, Lugano, Switzerland
New approaches in learning cardiac anatomy
- 16.20 – 16.40 **Giovanni Pedrazzini**, Cardiocentro Ticino, Lugano, Switzerland
Cardiac plaque imaging
- 16.45 – 17.05 **Stefanos Demertzis**,
From Leonardo da Vinci to the operating room via computational simulation: surgical reconstruction of the aortic root

WS 11 From stem cells to trees

Room: Sydney

- 15.30 – 16.00 **Henrik Jönsson**, Lund University, Sweden
Computational modelling and live imaging of plant development
- 16.00 – 16.30 **Cris Kuhlemeier**, University of Bern, Switzerland
Mathematical models of phyllotaxis
- 16.30 – 17.00 **Philipp Zimmermann**, ETH Zürich, Switzerland
Studying transcriptomes using gene and condition meta-profiles
- 17.00 – 17.30 **Christian Körner**, University of Basel, Switzerland
From assimilation to growth: carbon limitation in plants