



PRELIMINARY SCIENTIFIC PROGRAM

21 July 2014

14:00 Registration starts

16:00 Opening ceremony

Opening keynote lecture:

Douglas Lauffenburger, MIT, USA

Dynamical feedback circuits in cell biology: Network analysis of proteolytic shedding effects on receptor/ligand signaling

- 17:30 Welcome reception and dinner
- 20:30 Informal evening program (bowling, table soccer, swimming, etc)



Organizers:







Sponsors:















Structural biology and high-throughput experimental techniques

9:00 Keynote lecture:

Giulio Superti-Furga, CeMM, Austria

Genes, protein, protein complexes and environmental exchange: Understanding drug action in the context of environmental complexity

9:45 Expert talk:

Rune Linding, Technical University of Denmark, Denmark Decoding network dynamics in cancer

10:15 Expert talk:

Anne-Claude Gavin, EMBL, Germany Lipid transport in budding yeast

10:45 Tea/coffee break

11:15 Expert talk:

Gabor Balogh, BRC, Hungary

Lipidomics: Role of lipids in signalling networks

11:45 Contributed talks:

Laszlo Buday, MTA-TTK, Hungary

Tks scaffold proteins in tyrosine kinase signalling

Ivana Vonkova, EMBL, Germany

A systematic analysis of the PH domain recruitment to membranes using novel liposome microarray-based assay

Christin Zasada and Raphaela Fritsche, Max-Delbrueck-Center Berlin, Germany From signaling to metabolism and vice versa

12:30 Lunch

14:00 Teamwork - part 1

14:00 Getting know each other

Presenting the problem to be solved

16:45 Tea/coffee break

Medieval team building competition, informal discussions in the meantime 17:15

19:00 **Dinner**

21:30 Poster session 1

22:30 Informal evening program

Organizers:



















Network modelling

9:00 Keynote lecture:

Boris Kholodenko, University College Dublin, Ireland Catching signals surfing the net

9:45 Expert talk:

Laurence Calzone, INSERM at Institut Curie, France

Mathematical modelling of bladder tumorigenesis to explain co-occurrence and exclusivity of genetic alterations

10:15 Expert talk:

Julio Saez-Rodriguez, EMBL-EBI, United Kingdom Logic modelling of signalling networks in disease

10:45 Tea/coffee break

Contributed talks: 11:15

Attila Csikasz-Nagy, Kings College London, United Kingdom Epithelial topology dynamics in tissue homeostasis and tumorigenesis

Abulikemu Abudukelimu, VU University Amsterdam, The Netherlands

One stone, two birds: a therapy to both inflammation and tumorigenesis in a complex model of tumor incited inflammation

Aaron Meyer, MIT, USA

The AXL receptor is a sensor of ligand spatial heterogeneity

Orsolya Kapuy, Semmelweis University, Hungary

A systems-level study of mTOR-regulated autophagy induction during endoplasmic reticulum stress

Alexey Kolodkin, Luxembourg Centre for Systems Biomedicine, Luxemburg ROS-activated signaling network: dynamic modelling and design principles study

12:30 Lunch

14:00 Teamwork - part 2

14:00 Discussing the problem to be solved

15:30 Catamaran boat trip in the Danube Bend (each team in each catamaran)

17:00 Tea/coffee break

17:30 Summary activity on the problem, creating a presentation

19:00 **Dinner**

21:30 Poster session 2

22:30 Informal evening program

Organizers:





Sponsors:













Signaling systems & network medicine

9:00 Keynote lecture:

Hans Westerhoff, BioCentrum Amsterdam, The Netherlands Signalling principles

9:45 Expert talk:

Peter Csermely, Semmelweis University, Hungary

Adaptation of molecular networks and their use in drug design: a signaling perspective

10:15 Expert talk:

Paul Bates, Cancer Research, United Kingdom Modelling tumour heterogeneity and metastasis

10:45 Tea/coffee break

11:15 Contributed talks:

Sakshi Gulati, Cancer Research, UK - LRI, United Kingdom

Comprehensive analysis of molecular markers associated with survival and the impact of intratumour heterogeneity in clear cell renal cell carcinoma

Ferenc Gallyas, University of Pecs, Hungary

Mitochondrial network dynamics, tumour malignancy and small heat-shock protein expression: Is there a connection?

Alba Gutiérrez, Universitat Pompeu Fabra, Hospital del Mar Medical Research Institute, Spain A network medicine approach to explore comorbiditiy patterns in Catalonia

Nadezhda Doncheva, Max Planck Institute for Informatics, Germany

Dynamic residue interaction networks for understanding signaling protein interactions

András Zeke, MTA-TTK, Hungary

Identifying novel mitogen-activated protein kinase partners by a combined structural, sequence-based & evolutionary analysis

12:30 Lunch

Organizers:



The Genome Analysis Centre



















Further signaling network topics

Katalin Toth, University of Missouri, USA

Phosphorylation events involved in early perception of soybean rhizobial symbiont B. Japonicum

Bhaswar Ghosh, ZMBH, University of Heidelberg, Germany

Evolutionary flexibility achieved through phosphatase sharing in multilayer MAPK cascade

Nilgun Yilmaz, VU University Amsterdam, The Netherlands

Low stress weekends promote adaptation to stressful weeks, or do they?

14:45 Maria Victoria Schneider, TGAC, United Kingdom

How to be an interdisciplinary scientist - integrated discussion on the needs and requests

15:30 Presentation of teams

(10 minutes/team)

- 16:45 Tea/coffee break
- 17:15 Discussing group presentations, novel ideas, and concrete actions (e.g. collaborations)
- 18:00 Closing Keynote:

Toby Gibson, *EMBL*, *Germany* Regulation by molecular switching

19:00 Renaissance Farewell Feast,

Award presention for teams and poster presenters

Closing remarks

22:00 Informal evening program with music

25 July 2014

Departure's day

Organizers:





Sponsors:











