SYMPOSIUM

Associate Professor for Computational Biology

Max Delbrück Communications Center, Axon 1, Campus Berlin-Buch

Thursday, June 21, 2012

09:00 Welcome by Nikolaus Rajewsky

09:10 Zoran Nikoloski
Max Planck Institute of Molecular Plant Physiology, Potsdam

Coupling environments and models via data integration: The case of the TCA cycle

09:45 Philipp Khaitovich
CAS-MPG Partner Institute for Computational Biology, Shanghai

Systematic analysis of molecular changes accompanying human evolution

10:20 coffee break

10:40 Sven Bergmann
University of Lausanne, Swiss Institute of Bioinformatics, Lausanne

Computational analysis of biological systems - From developmental patterning to genome-wide association studies

11:15 João Pedro de Magalhães
University of Liverpool, Institute of Integrative Biology, Liverpool

Building haystacks and finding needles in the genomics of ageing

For further information, please contact: nadine.ewald@mdc-berlin.de