

LECTURE MEET THE SPEAKER*

4.00 - 5.00 pm 5.00 - 6.30 pm

*Registration is mandatory by sending an email to florence.henry@lih.lu

- (A) LECTURE**
University of Luxembourg (LCSB)
House of Biomedicine II
Ground Floor, Room 15
MEET THE SPEAKER*
Light snacks provided
University of Luxembourg (LCSB)
House of Biomedicine II
Ground Floor, Room 15
- (B) LECTURE**
Maison des Sciences Humaines
"Blackbox room"
MEET THE SPEAKER*
Light snacks provided
Maison des Sciences Humaines
Room N° 0015050, N° 0015040 or N° 2.07
- (C) LECTURE**
Maison du Savoir
Room N° 2.230, 4.340 or 4.510
MEET THE SPEAKER*
Light snacks provided
Maison du Savoir
Room N° 2.220, 4.390 or Espace A

LOCATIONS

Date	Title	Speaker	Invited by	Location
19 June	Hunt for the tipping points of diseases by dynamic network biomarkers	Prof Luonan Chen Chinese Academy of Sciences, Shanghai, China	Invited by : Feng He	(A)
11 Sept.	Biological interrogation of integrated resources of high-throughput chemical and phenotypic data of perturbations	Prof Mónica Campillos Institute of Bioinformatics and Systems Biology, Helmholtz Zentrum München, Germany	Invited by : Francisco Azuaje	(B)
07 Nov.	Analysing mammalian signal transduction and gene regulation using computational models	Dr Jana Wolf The Max Delbrück Center for Molecular Medicine in the Helmholtz Association, Berlin, Germany	Invited by : Gunnar Dittmar	(B)
01 Feb.	Translational bioinformatics in the precision medicine era	Prof Jessica Tenenbaum Department of Biostatistics and Bioinformatics, Duke University School of Medicine, Durham, NC, USA	Invited by : Francisco Azuaje Léon-Charles Tranchevent	(B)
14 Mar.	Systems Analysis of Cancer Metabolism for Identification of Cancer Drug Targets and Biomarkers	Prof Jens Nielsen Systems and Synthetic Biology, Chalmers University of Technology, Göteborg, Sweden	Invited by : Thomas Pfau Thomas Sauter	(C)
22 Mar.	Patient-past based precision medicine	Prof Søren Brunak Center for Biological Sequence Analysis (CBS), Lyngby, Denmark	Invited by : Reinhard Schneider	(B)
30 Nov.	Systems-level modeling between biocuration, predictions and validations	Dr Ioannis Xenarios Swiss Institute of Bioinformatics (SIB), Lausanne, Switzerland	Invited by : Francisco Azuaje	(C)
05 Dec.	Systematic integration of models and data for yeast growth and division	Prof Edda Klipp Institute of Biology, Humboldt-Universität zu Berlin, Berlin, Germany	Invited by : Gunnar Dittmar	(C)
03 May	Integrative approaches and workflows to identify disease specific biomarkers and therapeutic candidates	Dr Shailendra Gupta University of Rostock, Germany CSIR - Indian Institute of Toxicology Research, Lucknow, India	Invited by : Reinhard Schneider	(C)
26 June	Quantifying intratumoral heterogeneity at single cell level	Dr Andrei Zinovyev Institut Curie, Paris, France	Invited by : Francisco Azuaje Petr Nazarov	(A)
19 July	Title TBC			(B)
19 July		Prof Jochen Prehn Royal College of Surgeons in Ireland	Invited by : Reinhard Schneider	(B)
03 Dec.	Title TBC			(B)
03 Dec.		Prof Ursula Kummer University of Heidelberg, Germany	Invited by : Thomas Sauter Philippe Lucarelli	(B)

Coordinated by Department of Oncology (Bioinformatics and Modelling, Principal Investigator: Dr Francisco Azuaje)
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