

Current Challenges in Integrated Structural Biology

Program

June 19-20 2014

Day 1: Thursday, June 19th

12.00 - 14.00	Lunch buffet and Registration	
14.00	Welcome	
14:00 - 14:35	Opportunities for future multi-resolution integration at the CBI.	Bruno Klaholz
14.45 - 15.20	Imaging cells, viruses and protein complexes with 3D electron microscopy: From structure to mechanism.	Sriram Subramaniam
15.30 - 16.05	Geometry of the nucleus and gene regulation A single-molecule and super-resolution microscopy approach.	Ignacio Izeddin Aguirre
16.15 – 16.45	Coffee break	
16.45 - 17.20	Latest developments in correlative and cryo-TEM workflow solutions.	Sacha de Carlo
17.30 - 18.05	Large volume imaging of cellular ultrastructure using (cryo-) FIB-SEM microscopy.	Andreas Schertel
18.15 - 19.30	Poster session & visit of the CBI facilities	

Day 2: Friday, June 20th

9.00 – 9.35	Dynamic macromolecular complexes at high resolution by single particle cryo-EM.	Holger Stark
9.45 – 10.20	Using NMR and EPR to decipher the molecular mechanisms of translation regulation.	Frédéric Allain
10.30 - 11.00	Coffee break	
11.00 - 11.35	New developments in ITC to address biological problems.	Philippe Dumas
11.45 - 12.20	Hybrid pixel X-ray detectors for advanced life science applications.	Clemens Schulze- Briese
12.30 - 14.00	Lunch buffet	
14.00 - 14.35	In situ, multi-crystal, high frame-rate: New trends in X-ray crystallography data collection.	Gwyndaf Evans
14.45 - 15.20	Structural basis for transcriptional pausing in bacteria.	Albert Weixlbaumer
15.20 - 16.20	Selected posters: 12min + 3min discussion	
	Transient states in heme peroxidases: a multidisciplinary approach	Cecilia Casadei
	Structure and mechanism of the bacterial amyloid secretion channel CsgG	Imke Van den Broeck
	High-precision correlative fluorescence and electron cryo microscopy using two independent alignment markers	Pascale Schellenberger

16.20 – 18.00 Coffee break, specific discussions, posters, visits