PSDI 2018 - 26th Protein Structure Determination in Industry Meeting

11th - 13th November 2018

Hôtel Le LOUIS, Versailles Château - France

Programme

Sunday, November 11 th		
14:30 - 15:15	Registration	
15:15 - 16:45	1 st session of workshops, In parallel	
	Molecular Dimension/ Anatrace Montespan room Chair: James Gordon & Edward Pryor	Proteros Maintenon room Chair: Lars Neumann, Stephan Krapp & Claire Donat
16:45 - 17:00	Coffee break	
17:00 - 18:30	2 nd session of workshops, In parallel	
	Nanotemper Montespan room Chair: Pierrick Daniel & Pierre Soule	Global phasing Maintenon room Chair: Gerard Bricogne
18:30 - 19:00	Registration	
19:00 - 21:00	Welcome reception at the Hotel LE LOUIS	
21:00 - 22:00	Keynote speaker: A Decade of GPCR Structure-Based Drug I control of GPCR activity Andrew S. Doré - Heptares Therapeutics Ltd	Design - Structural insights into the allosteric



Monday, November 12 th		
07:45 - 08:15	Registration	
08:15 - 08:25	Welcome adress	
Session #1 – Biologics Conference room Chair: Djordje Musil		
08:25 - 08:45	Structural biology for antibody design at Roche *Armin Ruf - Roche Innovation Center Basel, Switzerland*	
08:45 - 09:05	Structural biology in the design of self-assembling protein nanoparticle vaccine antigens Matthew J. Bottomley - GSK, Rockville, USA	
09:05 - 09:25	Anti-biopharmaceutical immunization: Analysis of anti-drug antibodies interactions Thomas Bertrand - Sanofi R&D, Vitry-sur-Seine, France	
09:25 - 09:35	Sponsor talk : An analytical revolution: Introducing the next generation optima AUC Anthony Curran - Beckman Coulter, High Wycombe, U.K.	
09:35 - 09:45	Sponsor talk: New direct in-situ & non-invasive technique for the monitoring of bio-pharma product in injectable syringes David Jacob - Cordouan Technologies, Pessac, France	
09:45 - 09:55	Sponsor talk: Characterization of a novel ATP-cone driven activity regulation of ribonucleotide reductase by several techniques Stéphane Rouquette - Malvern Panalytical, Solna, Sweden	
09:55 - 10:30	Coffee break, Exhibitors & Networking	
Session #2 – CryoEM Conference room Chair: Wolfgang Baumeister		
10:30 - 11:15	Keynote speaker : Cryo-electron tomography: The promise and challenges of doing structural biology in situ Wolfgang Baumeister - Max Plank Institute of Biochemistry, Martinsried, Germany	
11:15 - 11:35	Instruct-NL: A focus on high resolution cryo electron microscopy at NeCEN Ludovic Renault - The Netherlands Centre for Electron Nanoscopy, Leiden, The Netherlands	



11:35 - 11:55	How Cryo-EM can push your structural biology boundaries for drug discovery? Hervé Remigy - ThermoFisher Scientific, Eindhoven, The Netherlands	
11:55 - 12:05	Sponsor talk: Detergent-free membrane protein sample preparation for cryo-electron microscopy Barbara Maertens - Cube biotech, Monheim, Germany	
12:05 - 12:15	Sponsor talk: Chameleon: A pico-litre dispense, self-wicking system for automated plunge freezing Paul Thaw - TTP Labtech, Melbourn, UK	
12:15 - 12:25	Sponsor talk : Pushing the boundaries of laboratory small angle X-ray scattering instrumentation Soren Skou - Xenocs Nordic, Hoersholm, Denmark	
12:25 - 13:30	Lunch break (buffet) - Exhibitors & Networking	
13:30	Departure for Synchrotron SOLEIL	
Session #3 - Cutting-edge technologies SOLEIL amphitheatre Chair: Andrew Thompson		
14:20 - 14:40	The Salipro® system for stabilization of membrane proteins *André Heuer - Salipro Biotech AB, Stockholm, Sweden*	
14:40 - 15:00	X-ray free electron laser: Opportunities for drug discovery Michael Hennig - LeadXpro AG, Villigen, Switzerland	
15:00 - 15:10	Sponsor talk: EIGER collects best data at the Synchrotron and in the laboratory Marcus Mueller - DECTRIS Ltd., Baden-Dättwil, Switzerland	
15:10 - 15:20	Sponsor talk: Latest innovations from Rigaku Oxford diffraction Mark Benson - Rigaku Europe, Oxford, U.K.	
15:20 - 15:30	Sponsor talk: Structural biology research center in KEK/high energy accelerator research organization Mikio Tanabe - KEK Tsukuba, Ibaraki, Japan	
15:30 - 16:10	Coffee break - poster session	
16:10 - 17:20	Synchrotron SOLEIL visit	
17:20 - 18:30	Round table	
18:45	Departure for Versailles	
19:30 - 22:30	Meeting Dinner	



Tuesday, November 13 th		
Session #4 - Membrane proteins Conference room Chair: Vadim Cherezov		
08:30 - 09:15	Keynote speaker: Dissecting GPCR structure and dynamics with X-ray lasers Vadim Cherezov - University of Southern California, Los Angeles, USA	
09:15 - 09:35	The high-resolution structure of a metabolite sensing GPCR bound to an intracellular nanobody Matthias Haffke - Novartis Pharma AG, Basel, Switzerland	
09:35 - 09:55	Application of microscale thermophoresis to GPCRs Alexey Mishin - Moscow Institute of Physics and Technology, Dolgoprudny, Russia	
Session #5 - Biophysical methods Conference room Chair: Alexey Rak		
09:55 - 10:15	¹⁹ F-NMR fragment screening at SANOFI <i>Magali Mathieu</i> - Sanofi R&D, Vitry-sur-Seine, France	
10:15 - 10:35	Evaluation of SPR based screening campaigns: Implications to define performance parameters and hit selection criteria Florian Krieger - Evotec AG, Hamburg, Germany	
10:35 - 11:05	Coffee break, Exhibitors and Networking	
Session #6 - Drug design case studies Conference room Chair: Armin Ruf		
11:05 - 11:25	A potent and orally bioavailable ERK1/2 series identified through fragment screening which modulates the phosphorylation and catalytic activity of ERK1/2 Puja Pathuri - Astex Pharmaceuticals, Cambridge, U.K.	
11:25 - 11:45	Fragment screening of lipoprotein-associated phospholipase A ₂ (Lp-PLA ₂) and the structure-based development of potent and selective lead molecules Philip Day - Astex Pharmaceuticals, Cambridge, U.K.	
11:45 - 12:05	Allosteric activation of striatal-enriched protein tyrosine phosphatase (STEP, PTPN5) by a fragment-like molecule Dennis Fiegen - Boehringer Ingelheim Pharma GmbH, Biberach an der Riss, Germany	



Session #7 - Computational method & crystallization Conference room Chair: Gérard Bricogne		
12:05 - 12:25	Memproc: Tools for semi-automated processing of data from serial crystallography experiments Matthias Haffke – Novartis Pharma AG, Basel, Switzerland	
12:25 - 12:45	Crystallophore, a nucleating and phasing agent Olivier Maury - École Normale Supérieure, Lyon, France	
12:45 - 13:05	Anticipated biased condition method for macromolecular crystallization Fabrice Gorrec - MRC Laboratory of Molecular Biology, Cambridge, U.K.	
13:05 - 13:15	Concluding remarks	
13:15 - 14:00	Lunch break (buffet)	
15:00	Visit of the Versailles castle for one who are registered (Duration 1h30)	