



National Hellenic Research Foundation (NHRF)
Athens, April 5th -10th 2014

Strategic pipeline planning: from sample preparation to 3D structure determination with bio SAXS and other biophysical technique

PROGRAMME

SATURDAY 05.04.2014

	Structural Biology in Europe 2014 onwards State-of-the-art facilities for cutting-edge research in Europe <ul style="list-style-type: none">○ ESFRI actions Instruct – a success story (more info in →)○ Large scale facilities in Europe &○ Opportunities for transnational access
08:45 – 09:30	Registration
09:30 – 10:00	Welcome remarks / Introduction NHRF, HeCrA
10:00 – 10:40	Integrated structural biology. The Instruct vision and some examples Prof. DAVID STUART , FRS, INSTRUMENT coordinator Professor of Structural Biology, Director of the Division of Structural Biology, MRC Research Professor <i>The Nuffield Department of Clinical Medicine, University of Oxford, UK</i>
10:40 – 11:20	Cryo-EM structural analysis of macromolecules: The emergence of a new universe of possibilities thanks to automation and new electron detectors. Prof. JOSE-MARIA CARAZO , Director of INSTRUMENT image processing centre Head of Biocomputing Unit, CNB and Node Director of Spanish National Bioinformatics Institute <i>National Centre for Biotechnology (CNB)-CSIC, Madrid, Spain</i>
11:20 – 12:00	Integrative use of synchrotron X-ray scattering with other structural techniques to characterize biological macromolecules and complexes. Dr DMITRI SVERGUN , BIOSTRUCT-X coordinator, SAXS Group leader <i>EMBL-Hamburg Unit, Germany</i>
12:00 – 12:30	Coffee break
12:30 – 13:10	Structural Biology at Diamond Light Source –An integrated approach Dr MARTIN WALSH , Life Science coordinator <i>Diamond Light Source, Oxford, UK</i>
13:10 – 13:50	Facilities for macromolecular crystallography at HZB-BESSY II in Berlin Dr UWE MÜLLER , Team Leader, Macromolecular Crystallography Group <i>Institute Soft Matter and Functional Materials HZB-Bessy II, Berlin, Germany</i>

13:50 – 14:15 **Big Data management in Structural Biology**
Prof YANNIS IOANNIDIS, President & General Director
“Athena”- Research and Innovation Centre in Information Communication and Knowledge technology, Athens, Greece

14:15 – 15:10 **Lunch break**

Perspectives in structural biology (part I)

An overview of additional main stream methods for structural biology
○ NMR, X-ray imaging, X-ray crystallography

15:10 – 15:50 **Cryo Soft X-ray tomography: an overview**
Dr EVA PEREIRO, Scientist Responsible of the Soft X-ray microscopy beamline at ALBA Light Source
ALBA Synchrotron facility, Barcelona, Spain

15:50 – 16:30 **From Single Proteins to Cellular Function: the Contribution of CERM-NMR Infrastructure**
Dr SIMONE CIOFI-BAFFONI, Faculty member of Magnetic Resonance Center and Department of Chemistry
CERM, Florence, Italy

16:30 – 17:10 **Synchrotron based fragment screening**
Dr UWE MUELLER, Team Leader, Macromolecular Crystallography Group
Institute Soft Matter and Functional Materials
HZB-Bessy II, Berlin, Germany

17:10 – 17:30 **Coffee break**

Career development session: Lessons learnt from mobility of young scientists & prospects in the frame of Horizon 2020

17:30 – 17:45 **Bottlenecks and opportunities of researchers' mobility**
Dr ALEXANDROS SAVVAIDIS, President of MCFA-Hellas
MCFA-Hellas & Institute of Engineering Seismology and Earthquake Engineering, Thessaloniki, Greece

Funding opportunities under Horizon 2020

17:45 – 18:00 **MARIA SAMARA**, Project officer National Contact Point for Horizon-2020 on Marie Skłodowska-Curie Actions

18:00 – 18:15 **GEORGIA TZENOU**, Coordinator of Enterprise Europe Network-Hellas, National Contact Point for Horizon 2020-RIs & Access to Finance
Coordinator of EU projects & networks

18:15 – 18:30 **CRISTINA PASCUAL**, National Contact Point for Horizon 2020 on the ERC Programme
National Documentation Center/NHRF, Athens Greece

SUNDAY 06.04.2014

Perspectives in Structural Biology (part II)

An overview of additional main stream methods for structural biology

10:00 – 10:45 **High throughput cloning and expression of recombinant proteins for structural biology**
Dr RAY OWENS, Head of the Oxford Protein Production Facility (OPPF) & Nuffield Department of Medicine Senior Fellow
OPPF- University of Oxford, UK

10:45 – 11:30 **3D Protein Structure Determination and protein-protein interaction by NMR**
Dr SIMONE CIOFI-BAFFONI, Faculty member of Magnetic Resonance Center and Department of Chemistry
CERM, Florence, Italy

11:30 – 12:15	Macromolecular Crystallography on the low-emittance synchrotron PETRA III Dr THOMAS SCHNEIDER , Coordinator MX@PETRA3, Group Leader <i>EMBL-Hamburg Unit, Germany</i>
12:15 – 12:45	Coffee break
12:45 – 13:15	Investigating the effect of structural changes <i>in vivo</i> through metabolic flux analysis and metabolomics Dr MARIA KLAPA , Head of Metabolic Engineering & Systems Biology Lab <i>Institute of Chemical Engineering Sciences (ICE-HT), FORTH-Patras, Greece</i>
13:15 – 13:45	Human and Animal Model Tissue Samples as High Quality Protein Sources for Structural and Proteomics Studies Dr LASKARINA-MARIA KOROU , Doctor of Veterinary Medicine <i>Laboratory for Experimental Surgery and Surgical Research "N.S. Christeas", Medical School University of Athens, Greece</i>
13:45 – 14:40	Lunch break
	Users' perspective
14:40 – 15:00	Structural Biology by NMR: Knowledge transfer from EU RIs to Greece Assoc. Prof. GEORGIOS SPYROULIAS <i>Department of Pharmacy, University of Patras, Greece</i>
15:00 – 17:00	Presentations by the participants on their work and motivation to attend the workshop
17:00 – 17:15	Coffee break
17:15 – 19:00	<i>Case studies – participants will form groups and discuss Brain storming and problem solving along with the tutors</i>
20:30	Dinner

MONDAY 07.04.2014

	Sample preparation (part I) What is needed for a "successful" sample preparation <ul style="list-style-type: none"> ○ Key objectives and how to benchmark the success of a strategy ○ Bottlenecks and trouble shooting ○ Decision making on method selection for 3D structure determination
09:15 – 10:00	Hybrid Structural Biology: recent examples and future perspectives Dr MATTHIAS WILMANN S, Head of EMBL-Hamburg Unit <i>EMBL-Hamburg Unit, Germany</i>
10:00 – 10:45	Construct design and protein expression Dr ROB MEIJERS , High Throughput Crystallization facility, Team Leader <i>EMBL-Hamburg Unit, Germany</i>
10:45 – 11:15	Coffee break
11:15 – 12:00	Quality control and protein optimization Dr ROB MEIJERS , High Throughput Crystallization facility, Team Leader <i>EMBL-Hamburg Unit, Germany</i>
12:00 – 12:45	The power of structural hybridomics Prof. Dr SAVVAS SAVVIDES , Head of Unit for Structural Biology and Biophysics <i>Lab. for Protein Biochemistry & Biomolecular Engineering (L-ProBE), Ghent University, Belgium</i>
12:45 – 13:05	Users' perspective

12:45 – 13:05	Integrated molecular and structural biology approaches towards unraveling the unique role of desmin's head domain in intermediate filament structure and function Dr MANOLIS MAVROIDIS Division of Cell Biology, <i>Biomedical Research Foundation, Academy of Athens</i>
13:05 – 13:25	Sensing biomolecular interactions in real time by SPR: principles, examples and caveats". Assist. Prof. PAVLOS (BOGOS) AGIANIAN, <i>Molecular biology & Genetics Department, Demokritus University of Thrace, Alexandroupolis, Greece</i>
13:25 – 14:15	Lunch break
14:15 – 15:45	Presentations by the participants on their work and motivation to attend the workshop
15:45 – 16:05	Coffee break
16:05 – 18:00	<i>Case studies – participants will form groups and discuss Brain storming and problem solving along with the tutors</i>

TUESDAY 08.04.2014

	Bio Small Angle X-ray Scattering (SAXS) (part I) <ul style="list-style-type: none"> ○ Bio SAXS at the forefront of structural biology ○ Basic principles of small angle X-ray scattering ○ Supporting biophysical techniques (e.g. Dynamic Light Scattering) ○ SAXS data analysis software <i>In vitro and in silico SAXS challenges</i>
09:00 – 09:45	Basics of SAXS by macromolecular solutions Dr DMITRI SVERGUN, BIOSTRUCT-X coordinator, SAXS Group leader <i>EMBL-Hamburg Unit, Germany</i>
09:45 – 10:30	It is worth the investment! Strategies to significantly improve SAXS data quality with a few additional biochemical/ biophysical steps. Dr MELISSA (Ann) GRAEWERT <i>EMBL-Hamburg Unit, Germany</i>
10:30 – 11:00	Coffee break
11:00 – 11:45	Ab initio shape determination Dr DMITRI SVERGUN, BIOSTRUCT-X coordinator, SAXS Group leader <i>EMBL-Hamburg Unit, Germany</i>
	Users' perspective
	Protein aggregation
11:45 – 12:05	Dr CONSTANTINOS VORGIAS, Professor of Biochemistry <i>Department of Biochemistry & Molecular Biology, Faculty of Biology, School of Science, National & Kapodistrian University of Athens, Greece</i>
12:05 – 12:25	Multiangl e dynamic light scattering: the method and some applications to protein solutions Dr STERGIOS PISPAS <i>Institute of Theoretical and Physical Chemistry, NHRF, Athens, Greece</i>
12:25 – 13:25	BioSAXS - Practical session
13:25 – 14:15	Lunch break
14:15 – 16:00	BioSAXS - Practical session
16:00 – 16:20	Coffee break
16:20 – 18:00	BioSAXS - Practical session

WEDNESDAY 09.04.2014

Bio Small Angle X-ray Scattering (SAXS) (part II)

- 09:00 – 10:30 **The use of high-resolution structures for SAXS-based modelling.**
Dr MAXIM PETOUKHOV
EMBL-Hamburg Unit, Germany
- 10:30 – 11:00 **Coffee break**
- 11:00 – 11:45 **Characterization of mixtures & intermolecular interactions by SAXS**
Dr PETR KONAREV
EMBL-Hamburg Unit, Germany
- 11:45 – 13:45 BioSAXS - Practical session
- 13:45 – 14:30 **Lunch break**
- 14:30 – 16:30 BioSAXS - Practical session
- 17:00 – 17:20 **Coffee break**
- 16:00 – 17:00 Questions and answers session
- 17:20 – 18:00 Presentations by the participants on their work and motivation to attend the workshop

THURSDAY 10.04.2014

Sample preparation (part II)

What is needed for a “successful” sample preparation

- Key objectives and how to benchmark the success of a strategy
- Bottlenecks and troubleshooting
- Decision making on method selection for 3D structure determination

- 09:00 – 09:45 **Highways, biways and detours in crystallization**
Prof. TERESE BERGFORS,
Department of Cell and Molecular Biology, Biomedical Centre, Uppsala University, Sweden
- 09:45 – 10:30 **Seeding Strategies for "Random" Crystal Screening and Crystal Optimization**
Dr PATRICK SHAW STEWART
Douglas Instruments Ltd, UK
- 10:30 – 12:00 Protein crystallization practical session
- 12:00 – 12:30 **Coffee break**
- 12:30 – 13:00 **New approaches to the search for crystallisation conditions and to crystal optimisation**
Dr EMMANUEL SARIDAKIS
Institute of Physical Chemistry NCSR- "Demokritos", Athens, Greece
- Users' perspective**
- 13:00 – 13:30 **From molten globules to bio-inspired materials**
Prof. MIKE KOKKINIDIS
University of Crete/Institute of Molecular Biology & Biotechnology, FORTH-Heraklion, Greece
- 13:30 – 14:00 **The Thermodynamic Stability of Proteins: Structure-based Issues and Interactions**
Dr GEORGE NOUNESIS, Deputy Director
Institute of Radioisotopes & Radiodiagnostic Products, NCSR- "Demokritos", Athens, Greece

14:00 – 14:30	Assessing protein integrity, stability and interactions by spectroscopy-based methods Dr Prof. ANASTASIA S. POLITOU , Assist. Prof. of Biological Chemistry <i>Medical School, University of Ioannina/Biomedical Research Division-FORTH-Ioannina, Greece</i>
14:30 – 15:15	Lunch break
14:00 – 15:45	Case studies – participants will form groups and discuss Brain storming and problem solving along with the tutors
15:15 – 15:45	Coffee break
15:45 – 18:00	Questions and answers session
18:00 – 18:15	Closing remarks