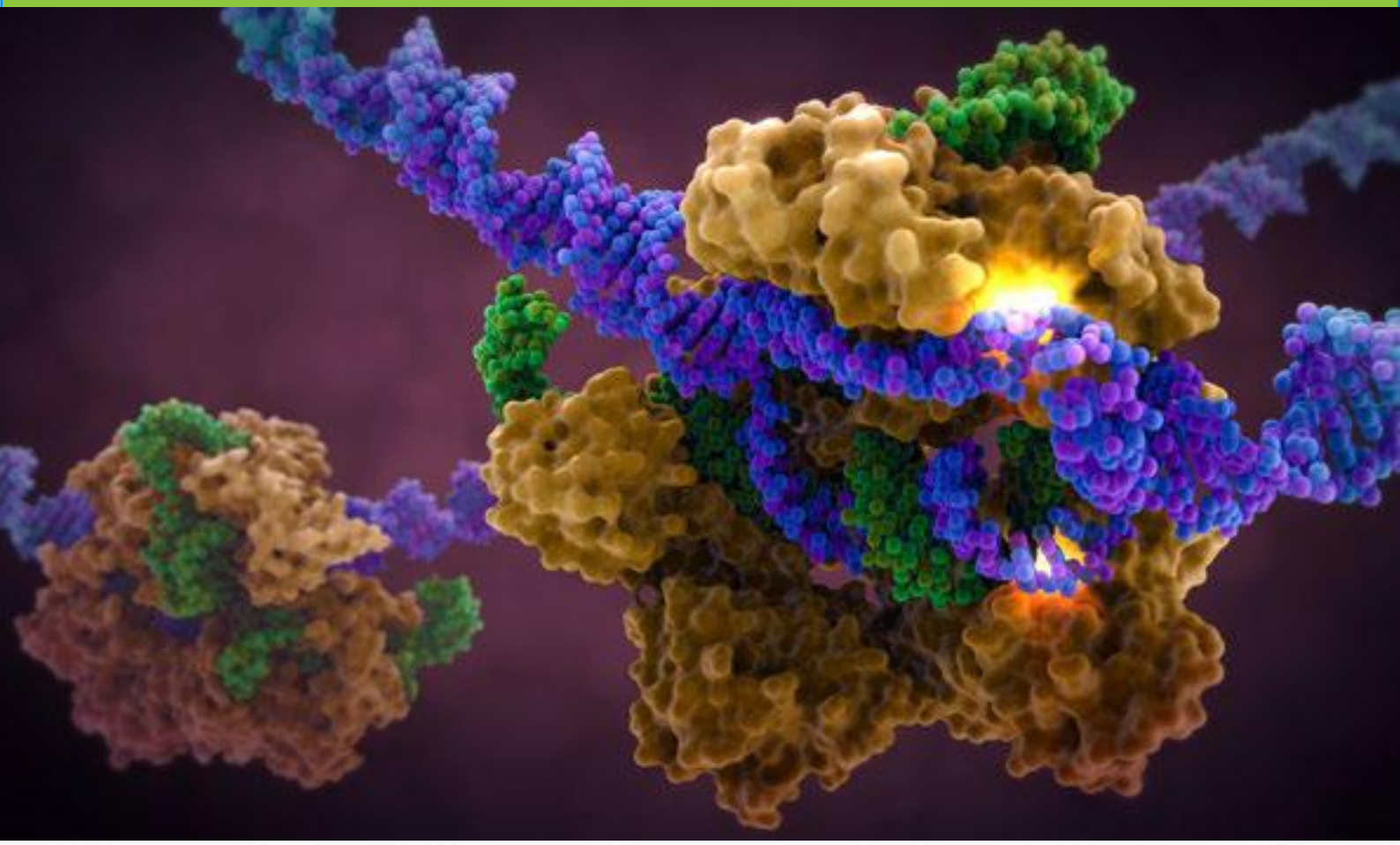
Tentative Program

9th International Conference on

Structural Biology

September 18-19, 2017 Zurich, Switzerland



Interactive Sessions

Keynote Lectures

Plenary Lectures

Workshops

Exhibitors

B2B Meetings

For available speaker slots

structuralbiology@biochemconferences.org

Conference Highlights

- 3D Structure Determination
- Computational Approaches in Structural Biology
- Hybrid Approaches for Structure Prediction
- Structural Biology and Single Molecules
- Sequencing
- Structural Biology Databases
- Cell Signalling
- Molecular Modelling and Simulation
- Drug Designing
- Structural Biology in Cancer Research
- Frontiers in Structural Biology
- Structural Biology Complexity Arenas
- Recent Advances in Structural Biology

Tentative Program

conferenceseries.com

9th International Conference on

Structural Biology

Structural Biology 2017

September 18-19, 2017 Zurich, Switzerland

Day 1

Opening Ceremony

Keynote Session

Title: The centers of premeltons signal the beginning and ends of genes
Henry M. Sobell, University of Rochester, USA

Title: Membrane protein structure, dynamics & function: Oriented sample and magic angle sample spinning solid state NMR
Timothy A. Cross, Florida State University, USA

2 Keynote slots available

Networking & Refreshments

Speaker Sessions

Title: Mechanical stability of talin rod controls traction force generation and cell migration
Vesa P. Hytönen, University of Tampere, Finland

Title: Structural insights into cholesterol regulation of inwardly-rectifying K+ channels
Irena Levitan, University of Illinois, USA

Title: Allosteric control of transcription regulation by nuclear receptors: an integrative structural biology approach
Dino Moras, IGBMC, France

Title: Transport protein superfamilies: Structure, function and evolution
Milton H. Saier, Jr., University of California at San Diego, USA

Title: Microrobotics enables non-contact, fully automated protein crystal harvesting
David F. Sargent, ETH Zurich, Switzerland

Title: Strategies to target protein-protein interactions
Marta Westwood, UCB Celltech, UK

Title: Cryo-Electron Microscopy Grid Preparation from Nanoliter-Sized Protein Samples and Single-Cell Extracts
Thomas Braun, University of Basel, Switzerland

Lunch

Title: On two ways to predict the protein folding process over a chaotic model
Christophe Guyeux, Université de Bourgogne Franche-Comté, France

Title: Influence of water environment on protein structure
Irena Roterman, Jagiellonian University, Poland

Title: How Does Domain Motion Contribute to Transition-State Stabilization? Combinatorial Thermodynamic Cycle Analysis of Conformational Coupling During Tryptophan Activation
Charles W. Carter, University of North Carolina at Chapel Hill, USA

Title: Calcium, calmodulin and the plasma membrane calcium pump
Joachim Krebs, Max Planck Institute for Biophysical Chemistry, Germany

Title: An unbiased view of functional shifts and co-evolution across large protein families: Integrating computational and experimental approaches
Bryan Kolaczkowski, University of Florida, USA

Title: Achievements and challenges in computational protein design: from membrane protein parameterization to new functional proteins
Ilan Samish, Weizmann Institute of Science and Amai Proteins Ltd., Israel

Title: Structure of DdrB-DNA complex reveals mechanism for accurate DNA strand annealing
Murray Junop, Western University, Canada

Title: RNA structural biology: from global to detail
Yun-Xing Wang, National Institutes of Health, USA

Special Session on 'The Structural Biology of Biomembranes' by **Qiu-Xing Jiang**, University of Florida, USA

Networking & Refreshments

5 speaker slots available

Day 2

Keynote Session

Tilman Schirmer, University of Basel, Switzerland

Shigeyuki Yokoyama, RIKEN, Japan

3 Keynote slots available

Networking & Refreshments

Speaker Sessions

Title: Nanoscale structure and dynamics of centromere nucleosomes
Yuri L. Lyubchenko, University of Nebraska Medical Center, USA

Title: Monitoring protein structural changes on a proteome-wide scale
Paola Picotti, ETH Zurich, Switzerland

Title: Extract the thermodynamic and kinetic information from protein simulations using dimensionality reduction
Shuanghong Huo, Clark University, USA

Title: LD Motif Interacting networks in cell-matrix adhesion
Igor L. Barsukov, University of Liverpool, UK

Title: Solution NMR relaxation and µs Molecular Dynamics Simulations of Dynamic Protein-Protein and Protein-Membrane Complexes
Matthias Buck, Case Western Reserve University, USA

Title: Structural basis for PTPN3-p38gamma complex involved in colon cancer progression
Tzu-Ching Meng, Academia Sinica, Taiwan

Title: Structural Conformational changes report biased agonism: The case of Galanin receptors
Arfaxad Reyes Alcaraz, Korea University College of Medicine, South Korea

Lunch

Title: Does the dynamics of their transmembrane domain qualify bitopic membrane proteins as substrates for intramembrane proteolysis?
Christina Scharnagl, Technical University of Munich, Germany

Title: Role of metavinculin in actin reorganization and force transmission
Sharon L. Campbell, University of North Carolina, USA

Title: Type I BIR domain inhibitors in cancer therapy: Designing drugs to modulate the NF-κB pathway
Federica Cossu, Institute of Biophysics at the National Research Council (IBF-CNR), Italy

Title: Structural biology of RAGE
Alex Shekhtman, State University of New York, USA

Title: Structural Plasticity allows for diversity in GTP hydrolysis mechanisms: Implications for GAP-dependent hydrolysis mechanisms for Rab proteins
Balaji Prakash, CSIR- Central Food Technological Research Institute, India

Title: Role of conformational equilibrium in molecular recognition and capsid assembly: The case of flavivirus capsid proteins
Fabio C. L. Almeida, Federal University of Rio de Janeiro, Brazil

Title: Investigation of allosteric coupling in human Beta2-adrenergic receptor (Beta2-AR) in the presence of intracellular loop 3
E. Demet Akten, Kadir Has University, Turkey

Networking & Refreshments

6 speaker slots available

Young Researcher's Forum
Poster Presentations

Title: Evolution of Immunoglobulins through studies of structure and function of IgE and IgM
Rosemary Nyamboya, King's College London, UK

Title: Searching drugs for NF-κB pathway regulation: BIR1 domains of IAPs as promising new targets
Luca Sorrentino, Institute of Biophysics at the National Research Council (IBF-CNR), Italy

Title: Dissecting the molecular basis of Nrf2-p21 interaction
Nadun Karunatilleke, The University of Western Ontario, Canada

Title: The challenge of improve disease treatment with protein engineering. The contributions of X-ray crystallography
Stephanie Bath de Morais, FIOCRUZ, Brazil

Title: Ameliorating effect of coconut water on the epithelium and gastric goblet cells of albino wistar rats induced with castor oil
Gabriel Donatus Edem, University of Uyo, Nigeria

Title: Genome variation and drug resistance: Effects of non-synonymous polymorphisms on Topoisomerase II α (Top2-α) drug interaction
Sadeq Vallian, University of Isfahan, Iran

Title: Development of Cytokine inhibitors ; Application of MD simulation
Ruqaiya Khalil, University of Karachi, Pakistan

Award Ceremony

Website: http://structuralbiology.conferenceseries.com/

Last updated January 31, 2017

Glimpses of Structural Biology 2016





TOURIST DESTINATIONS IN ZURICH, SWITZERLAND



Baden



Bahnhofstrasse



Bellerive Museum



Paradeplatz



China Garden



Confiserie Sprüngli



Fraumünster



Ganymed



Giacometti Hall



Laufen Castle

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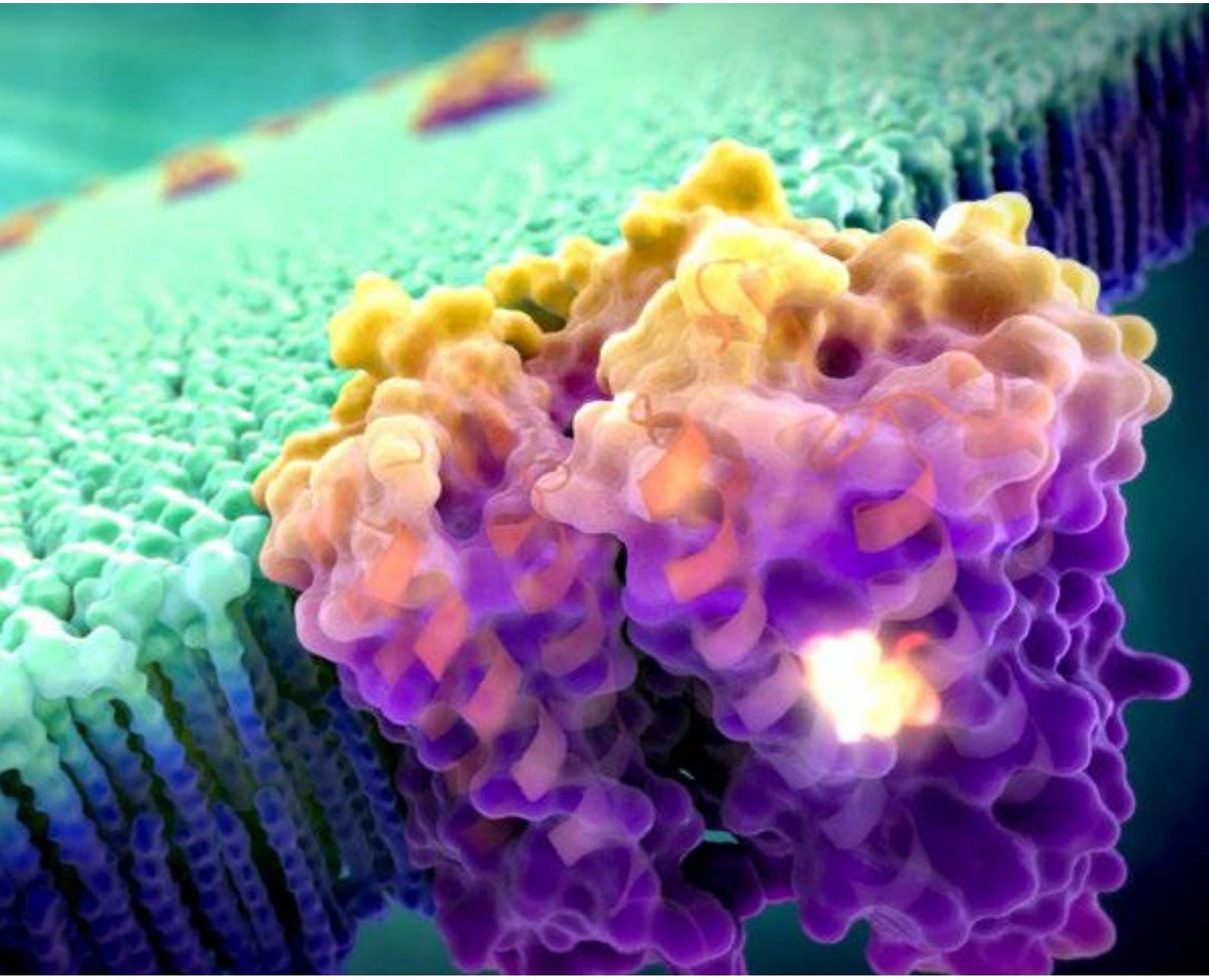
630th Conference

Scientific Program

6th International Conference on

STRUCTURAL BIOLOGY

August 22-23, 2016 New Orleans, USA



Hosting Organization: Conference Series LLC

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Scientific Program

Day 1 August 22, 2016

Registrations

Hall C

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Opening Ceremony

Keynote Forum

Introduction

Title: Comprehensive prediction of drug-protein interactions and side effects for the human proteome
Jeffrey Skolnick, Georgia Institute of Technology, USA

Symposium on "New technologies in structural studies by cryoEM"

Session Introduction

Session Chair: Qiu-Xing Jiang, University of Florida, USA
Session Co-Chair: Kenneth A Taylor, Florida State University, USA

Title: The structure of Zika virus
Michael G Rossmann, Purdue University, USA
Title: Random spherically constrained (RSC) single particle cryo-EM for structures of membrane proteins in a membrane environment
Liguo Wang, University of Washington, USA
Title: ChemiC: Chemically functionalized nanometer-thick carbon films for cryo-EM studies of biological complexes
Qiu-Xing Jiang, University of Florida, USA

Group Photo

Networking & Refreshments

Title: Deciphering intermolecular interactions in complex heterogeneous systems
Kenneth A Taylor, Florida State University, USA
Title: Breathing life into static images: Combining electron cryotomography and coarse-grained simulations to explore mechanisms of eukaryotic cell division machinery *in vivo*
Matthew T Swulius, California Institute of Technology, USA
Title: Advanced machine-learning algorithms for improving high-performance cryo-EM data analysis
Youdong Jack Mao, Harvard Medical School, USA

Lunch

Keynote Forum

Title: A personal history of structural virology
Michael G Rossmann, Purdue University, USA

Sessions: Structural Biology | 3 D Structure Prediction | Computational Approaches

Session Introduction

Session Chair: Fumio Hirata, Toyota Physical & Chemical Research Institute, Japan
Session Co-Chair: John H Miller, Victoria University of Wellington, New Zealand

Title: Tubulin structural interactions with small molecule microtubule-stabilizing agents
John H Miller, Victoria University of Wellington, New Zealand
Title: Molecular behavior of a higher-order complex of multiple transcription factors on enhancer site upon phosphorylation
Kazuhiro Ogata, Yokohama City University Graduate School of Medicine, Japan
Title: Structure, fluctuation, and function of biomolecules in solution explored by the 3D-RISM/RISM theory
Fumio Hirata, Toyota Physical & Chemical Research Institute, Japan
Title: Discovery of a GTP sensor using a structural reverse genetic approach
Toshiya Senda, High Energy Accelerator Research Organization (KEK), Japan
Title: Structural and synthetic biology of aminoacyl-tRNA synthesis and translation
Shigeyuki Yokoyama, RIKEN Structural Biology Laboratory, Japan

Networking & Refreshments

Sessions: Signalling Biology | Drug Designing | Sequencing

Session Introduction

Session Chair: Takeshi Kikuchi, Ritsumeikan University, Japan
Session Co-Chair: Elizabeth J Goldsmith, University of Texas Southwestern Medical Center at Dallas, USA

Title: Development of anti-HIV double variable domain antibodies that bind both gp120 and gp41 on the envelope protein
Seth Pincus, Children's Hospital and Louisiana State University School of Medicine, USA
Title: The excursion model for MAP kinase signaling in health and disease
Elizabeth J Goldsmith, University of Texas Southwestern Medical Center at Dallas, USA
Title: Sequence analysis on the information of folding nuclei in proteins with interesting 3D properties
Takeshi Kikuchi, Ritsumeikan University, Japan
Title: Ligand binding preferences of pathogenesis-related class 10 (pr-10) allergens
Barry K Hurlburt, United States Department of Agriculture, USA
Title: Genomes repetitive sequences in progressive evolution
Olga I Podgornaya, Far Eastern Federal University, Russia

Special Session on "Prediction of protein-protein interaction in the Twilight Zone"

Session Introduction

Session Chair: Thomas Scior, Benemérita Universidad Autónoma de Puebla, Mexico

Title: Prediction of protein-protein interaction in the twilight zone of low homology for the modulation of an ion channel: An analogy approach
Thomas Scior & Ángel Islas, Benemérita Universidad Autónoma de Puebla, Mexico

Panel Discussions

Day 2 August 23, 2016

Hall C

Keynote Forum

Title: The use of soft X-ray diffraction for native-SAD and for chromatic exploration of uncharted aspects of metals in macromolecules: A new pilot program at the APS
Bi-Cheng Wang, University of Georgia and Advanced Photon Source, USA
Title: How do voltage sensors in trans-membrane channels work?
Robert Stroud, University of California, San Francisco, USA

Special Session on "Modularity and metastability: Structural aspects of nanomedicine"

Session Chair: R Holland Cheng, University of California, Davis, USA
Session Co-Chair: Toshiya Senda, High Energy Accelerator Research Organization (KEK), Japan

Title: Modularity and metastability: Structural aspects of nanomedicine
R Holland Cheng, University of California, Davis, USA
Title: Structure and function of the C-terminal ID region of onco-protein CagA
Toshiya Senda, High Energy Accelerator Research Organization (KEK), Japan

Networking & Refreshments

Title: Talin - Insights into mechanosignalling and integrin activation
Vesa P Hytonen, University of Tampere, Finland
Title: Individual-Particle Electron Tomography (IPET): An approach to study flexible protein structure and dynamics
Gang Ren, Lawrence Berkeley National Laboratory, USA
Title: Integrated methods enable re-engineering of transcription factors allowing cell fate modification
Prasanna R Kolatkar, Hamad bin Khalifa University, Qatar
Title: A comprehensive strategy to obtain high quality crystals
Miki Senda, High Energy Accelerator Research Organization (KEK), Japan
Title: Symmetry ambiguity and its implications to the molecular functions of the MAVS prion-like filaments
Qiu-Xing Jiang, University of Florida, USA

Lunch

Title: Crystal structure of a human Fab in complex with a dominant antigen from Neisseria meningitidis
Jacinto López-Sagaseta, GlaxoSmithKline Vaccines S.r.l., Italy
Title: The atomic structures of Dragon grouper nervous necrosis virus from cryo-em in comparing with x-ray diffractions
Chan-Shing Lin, National Sun Yat-sen University, Taiwan

Sessions: Molecular Modelling | Complexity Arenas in Structural Biology | Structural Biology in Cancer

Session Introduction

Session Chair: Irena Roterman-Konieczna, Jagiellonian University, Poland
Session Co-Chair: Jyh-Yeuan Lee, University of Texas Southwestern Medical Center at Dallas, USA

Title: Impact of the water environment on protien folding
Irena Roterman-Konieczna, Jagiellonian University, Poland
Title: ABCG5/ABCG8: Structural role in ABC transporter-mediated sterol transport
Jyh-Yeuan Lee, University of Texas Southwestern Medical Center at Dallas, USA
Title: The molecular effects of processing on the peanut allergens
Soheila J Maleki, United States Department of Agriculture, USA

Title: Structural analysis of PTPN3-p38γ interactions by chemical cross-linking coupled with mass spectrometry
Tzu-Ching Meng, Academia Sinica, Taiwan

Networking & Refreshments

Title: Predicting the binding free energy of the inclusion process of Cyclodextrin derivatives and small molecules by means of MM/3D-RISM method
Masatake Sugita, Ritsumeikan University, Japan
Title: Is molecular docking the holy grail of computational modelling?
Ihosvany Camps, Federal University of Alfenas, Brazil
Title: Proton tunneling accelerates ATP hydrolysis in Eg5 kinesin
Sunyoung Kim, Louisiana State University School of Medicine & Health Sciences Center, USA
Title: Molecular docking of potent inhibitors of oncoproteins using Swiss dock and visualization by UCSF chimera
Mohammed Ahmed Mohammed Mansour, Tanta University, Egypt

Poster Presentations from

P01	**Title: In silico development of new acetylcholinesterase inhibitors** **Ihosvany Camps**, Federal University of Alfenas, Brazil
P02	**Title: Cation roles in the structure of a fish nodavirus observed in Electron Cryo-Microscopy and X-Ray diffraction** **Chan-Shing Lin**, National Sun Yat-sen University, Taiwan
P03	**Title: Estimation of binding free energy based on the MM/3D-RISM method for the Pim1-ligand system** **Takeshi Haesegawa**, Ritsumeikan University, Japan
P04	**Title: Motor domain β-sheet motifs contribute to diversification of kinesin-microtubule interactions** **Jessica Richard**, Louisiana State University School of Medicine & Health Sciences Center, USA
P05	**Title: SATB2 suppresses invadopodia formation in colorectal cancer cells via palladin inhibition** **Mohammed Ahmed Mohammed Mansour**, Tanta University faculty of Science, Egypt
P06	**Title: Folding region predictions by amino acid analysis of lysozyme superfamily proteins** **Takuto Nakashima**, Ritsumeikan University, Japan
P07	**Title: Small spatial motifs within a surface-exposed loop in kinesin motor proteins codes for mechanochemical coupling** **Rebecca Buckley**, Louisiana State University School of Medicine & Health Sciences Center, USA

Panel Discussion and Certificate Distribution