

	29.05 - SUNDAY	30.05 - MONDAY	31.05 - TUESDAY	01.06 - WEDNESDAY
8:00 – 9:15	BREAKFAST (8:00 – 9:30)			
9:15 – 9:30	L101 (9:15 – 10:00) Weontae Lee Non-Thermal Plasma and Protein Folding.	L201 (9:15 – 10:00) Victor Muñoz Experimental Studies of Fast-Folding Proteins: Lessons about Free Energy Barriers and Pre-Exponential Factors in Protein Folding.	L301 (9:30 – 10:15) Keehyoung Joo Protein Structure Determination by Global Optimization Using Sparse and Ambiguous NMR restraints.	L401 (9:30 – 10:15) Haiguang Liu Revealing structures and dynamics of proteins using the X-ray lasers: from a computational perspective.
9:30 – 9:45				
9:45 – 10:00				
10:00 – 10:15	L102 (10:00 – 10:45) Koliński Andrzej Efficient Modeling of Protein Structure and Dynamics.	L202 (10:00 – 10:45) Maciej Kozak The order-disorder processes in solution SAXS studies of the protein folding.	L302 (10:15 – 11:00) Igor Konieczny Homology Modeling, Molecular Dynamics, Biochemistry and Mass Spectrometry for Description of Nucleoprotein Complex of DNA Replication Initiator.	L402 (10:15 – 11:00) Martin Zacharias Exploring biomolecular dynamics and interactions using enhanced sampling approaches.
10:15 – 10:30				
10:30 – 10:45	L102 (10:45 – 11:00) Agnieszka Lipska Molecular dynamics of protein A and a WW domain with a united-residue model including hydrodynamic interaction.	L203 (10:45 – 11:00) Qianyi Cheng Folding Mechanisms of Small Proteins GB1 and LB1.		
10:45 – 11:00				
11:00 – 11:30	COFFEE (11:00 – 11:30)			
11:30 – 11:45	L103 (11:30 – 12:15) Jooyoung Lee Protein Structure Prediction/Determination by Global Optimization.	L204 (11:30 – 12:15) Masaki Sasai Dynamical Energy Landscape Perspective of Protein Functioning.	L303 (11:30 – 12:00) Sylwia Rodziewicz – Motowidło Structural studies of the HVEM protein and its fragments interacting with BTLA and CD160 proteins.	L403 (11:30 – 12:15) Changbong Hyeon Contact statistics highlight distinct organizing principles of RNA and proteins.
11:45 – 12:00				
12:00 – 12:15				
12:15 – 12:30	L104 (12:15 – 12:45) Yi He Correlations of the structure - dynamics – function of structurally homologous CheY-like proteins.	L205 (12:15 – 12:45) Stanislaw Oldziej Foldons, Protein Folding Nucleation Sites.	L304 (12:00 – 12:45) Joanna Skórko-Głonek Regulation of the Proteolytic Activity of the HtrA (DegP) Protease from <i>Escherichia Coli</i> ; Role of the Regulatory Loops.	L404 (12:15 – 13:00) Antti Niemi Solution X-ray scattering (S/WAXS) and the formation of large scale structure in protein dynamics.
12:30 – 12:45				
12:45 – 13:00	L105 (12:45 – 13:00) Balachandran Manavalan Support –vector-machine-based protein single-model quality assessment.	L206 (12:45 – 13:00) Marta Wiśniewska Buried Water Molecules in Protein Kinases.	LUNCH 12:50 – 13:25 (FOR MALBORK PARTICIPANTS)	CLOSING REMARKS
13:00 – 14:00	LUNCH (13:00 – 14:00)			
14:00 – 14:15	L106 (14:00 – 14:45) Philippe Derreumaux All-atom and coarse-grained simulations for protein structure prediction and protein aggregation dynamics and thermodynamics.	L207 (14:00 – 14:45) Joanna Sulkowska Energy landscape of Supercoiled Protein Folding.	MALBORK (13:30 – 19:00)	
14:15 – 14:30				
14:30 – 14:45				
14:45 – 15:00	L107 (14:45 – 15:00) InSuk Jung Study of conformational diversity of GC-repeating ds-DNA.	L208 (14:45 – 15:15) Adam Sierdzan Unique Features of Telomers.		
15:00 – 15:15	L108 (15:00 – 15:15) Pawel Krupa Role of disulfide bonds on folding and structures of proteins.			
15:15 – 15:30	L109 (15:15 – 15:30) Seung Hwan Hong A fragment search method for Dynamic Fragment Assembly.	L209 (15:15 – 15:45) Jaroslaw Marszalek Evolutionary Dynamics of Protein - Protein interactions, a Case Study of Proteins Involved in Iron - Sulfur Cluster Biogenesis.		
15:30 – 15:45	COFFEE (15:30 – 16:00)			
15:45 – 16:00	COFFEE (15:45 – 16:15)			
16:00 – 16:15	L110 (16:00 – 16:45) Kunihiro Kuwajima The problem of protein folding.	GDAŃSK		
16:15 – 16:30				
16:30 – 16:45				
16:45 – 17:00	L111 (16:45 – 17:30) In-Ho Lee Pathway Model of Mini-Protein BBA5.			
17:00 – 17:15				
17:15 – 17:30				
17:30 – 17:45	L112 (17:30 – 17:45) Magdalena Mozolewska Mechanism of Fe/S cluster transfer.			
17:45 – 18:30				
18:30 – 19:30	POSTER SESION (18:30 – 19:30)			
19:30 – 20:00	CONFERENCE BANQUET		19:00 – 20:00 SUPPER in the Hotel	GDAŃSK
20:00 – 22:00			20:00 – 21:00 SUPPER in the Hotel	