

PROGRAMME

Thursday,	March	16
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13:00 - 15:00 Registration

15:10 - 15:15 Introductory remarks

15:15 - 16:30 **Lecture session I**, chairman: *V. Sychrovský* L1 p. 3

Tomáš Obšil

Regulation by molecular interference: How 14-3-3 protein controls activity of forkhead transcription factor FOXO4 L2

Karel Bezouška

Optical spectroscopy, mass spectrometry and NMR are essential tools in the production of soluble receptors of natural killer cells

L3 p. 5

Jiří Vlach

A single-point mutation of M-PMV matrix protein causes reorientation of protein domains and changes the phenotype of the virus

L4 p. 5

Zuzana Vokáčová

NMR parameters in RNA molecules and their correlation with molecular structure

16:30 - 17:00 Coffee break

17:00 - 18:00 **Plenary lecture I**, chairman: *V. Sychrovský* PL1 p. 6

Vladimír Sklenář

BioNMR: Contemporary Trends, Challenges, and Future Prospects

18:30 - 22:30 Banquet

Friday, March 17

7:00 - 8:45 Breakfast

9:00 - 10:00 **Plenary lecture II**, chairman: *D. Štys* PL2 p. 6

Vladimír Baumruk

Methods of optical spectroscopy for structural biology

10:00 - 10:20 Coffee break

10:20 - 12:00 Lecture session II, chairman: D. Štys L5 p. 6

Jan Florian

DNA replication fidelity: Theory and experiment L6

Pavel Macek

Molecular dynamics study of major urinary proteinpheromone interactions: A structural model for ligand induced flexibility increase

L7 p. 7

Jiří Šebek

MD and Ab Initio Modeling of Electronic Spectra of

N-methylacetamide and Peptides in Water Solutions

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Peter Flecker

Molecular Modelling studies with Bowman Birk inhibitors

Michal Hušák

The use of GPU based calculations in structural biology L10 p. 9

Patrik Florek

Bacterial programmed cell death - genetical, biochemical, and structural insight

12:15 - 14:00 Lunch

14:15 - 15:00 **Plenary lecture III**, chairman: *B. Schneider* PL3 p. 10

Rudiger Ettrich

Molecular systems biology: Molecular simulations of proteins and their complexes

15:00 - 15:15 Coffee break

15:15 - 16:55 **Lecture session III**, chairman: *B. Schneider* L11 p. 10

Matěj Lexa

The accuracy of domain boundary detection in proteins based on segment co-occurence compared to other methods

L12 p. 11

Peter Palenčár

Crystal structure of photosystem II from *Thermosyne-chococcus elongatus* refined and studied by molecular dynamics

L13 p. 12

Jiří Brynda

From non-peptide towards non-carbon protease inhibitors: metallacarboranes as specific and potent inhibitors of HIV protease

L14 p. 13

Veronika Krejčiříková

Structural studies of galectin-4 N-terminal domain in complex with lactose

L15 p. 13

Daniel Svozil

DNA Conformational Families

16:55 - 17:15 Coffee break

17:15 - 18:00 **Plenary lecture IV**, chairman: *B. Schneider* PL4 p. 14

Jiří Šponer

Structure and dynamics of RNA and DNA. Advanced computational studies

18:30 - 19:30 Dinner

19:45 - 22:00 Poster session

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Saturday, March 18

7:00 - 9:10 Breakfast

9:15 - 10:15 Plenary lecture V, chairman: J. Brynda p. 14

Jan Dohnálek

Structure elucidation by diffraction methods - contribution to structural biology

10:15 - 10:30 Coffee break

10:30 - 11:50 Lecture session IV, chairman: J. Damborský L16 p. 15

Zbyněk Prokop

Discovery of Stereoselective Haloalkane Dehalogenase: New Tool for Assymetric Synthesis

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Tomáš Mozga

Structure-enantioselectivity relationships of haloalkane dehalogenases

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Radka Chaloupková

Physical variables as a control of enzyme stereoselectivity L19 p. 17

Táňa Chrobáková

Engineering of enantioselective haloalkane dehalogenase by cumulative mutagenesis

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Martin Klvana

Modelling of Product Release and Identification of Export Routes in Haloalkane Dehalogenase DhaA

11:55 - 12:00 Concluding remarks

12:15 - 14:00 Lunch

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K. Hofbauerová

Human protein CD69 studied by combination of vibrational spectroscopy and molecular

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E. Chovancová

Evolution of haloalkane dehalogenase protein family

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P. Jeřábek:

Computer-Assisted Design of Fluorescent Probes for Solvent Relaxation Experiments

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NMR study of protein CD69



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M. Navrátil

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PAIN: Program for Analysis of Internal motion. An Application to MD Simulated Motions of Major Urinary Protein-I

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14-3-3 protein changes conformation of nuclear localization sequence of forkhead transcription factor FOXO4

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P. Pavelčíková

SpoIIS components of Bacillus subtilis programmed cell death

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A. Samad

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Refined structural model of photosystem II from *Thermosynechococcus elongatus*, structural changes in reaction centre upon light induced oxidation of active accessory chlorophyll and reduction of active pheophytin

Castech

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