



PROGRAMME

Thursday, March 29

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| 12:00 - 14:00 | Registration |
| 14:15 - 14:20 | Introductory remarks |
| 14:20 - 15:20 | Lecture session I , chairperson: <i>P. Řezáčová</i> |
| L1 | <i>Pavel Plevka</i> The Structure of Ryegrass Mottle Virus |
| L2 | <i>Petr Kolenko</i> Crystallographic studies of intermolecular interactions of Fc-fragment of immunoglobulin |
| L3 | <i>Michal Kolář</i> Protein as a Flexible Structure: Case Study of the HIV Protease |
| 15:20 - 15:30 | Coffee break |
| 15:30 - 16:20 | Lecture session II , chairperson: <i>P. Řezáčová</i> |
| L4 | <i>Jiří Brynda</i> TLS analysis of antibody structure shows stabilization by epitope binding |
| L5 | <i>Tomáš Obšil</i> Interaction of DNA-binding domain of transcription factor FoxO4 with DNA |
| 16:20 - 16:30 | Coffee break |
| 16:30 - 17:05 | Plenary lecture I , chairperson: <i>B. Schneider</i> |
| L6 | <i>Filip Lankáš</i> Structural mechanics of DNA: from atomic scale to bioinformatics |
| 17:05 - 17:15 | Coffee break |
| 17:45 - 18:15 | Plenary lecture II , chairperson: <i>J. Hašek</i> |
| L7 | <i>Tomáš Polívka</i> Time-resolved structural dynamics - A challenge for future |
| 18:00 - 19:00 | Concert of violin duo Lukáš Kuta & Václav Polívka J.M Leclair, J.S. Bach, S. Prokofjev and others <i>Detailed program will be available on registration desk</i> |
| 19:00 - 22:30 | Banquet |

Friday, March 30

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| 7:00 - 8:45 | Breakfast |
| 9:00 - 10:50 | Lecture session III , chairperson: <i>R. Hrabal</i> |
| L8 | <i>Lukáš Žídek</i> Orientalional restraints in NMR structure determination |
| L9 | <i>Pavel Srb</i> Is local flexibility of M-PMV matrix protein related to its binding affinity with a molecular motor? |
| L10 | <i>Richard Štefl</i> RNA recognition by the ADAR2 dsRBMs: a 50 kDa protein-RNA complex by NMRL8 |
| L11 | <i>Vladimír Sychrovský</i> Calculation of Scalar Couplings in the Backbone of Nucleic Acid |
| 10:50 - 11:00 | Coffee break |
| 11:00 - 12:20 | Lecture session IV , chairperson: <i>L. Žídek</i> |
| L12 | <i>Zuzana Vokáčová</i> Theoretical NMR Study of Watson-Crick/Sugar Edge RNA Base Pair Family |
| L13 | <i>Jaroslav Burda</i> Computational study on spectral properties of the selected pigments from various photosystems; structure--transition energy relationship |
| L14 | <i>Jiří Mrázek</i> Electronic structure of the non-heme iron center in photosynthetic apparatus of plants and bacteria. Computational DFT study |
| L15 | <i>Tomáš Zimmerman</i> Computational study of cysteine and methionine interaction with platinum(II) complexes in water |
| 12:30 - 14:00 | Lunch |
| 14:00 - 15:20 | Lecture session V , chairperson: <i>J. Pačes</i> |
| L16 | <i>Jan Adam</i> Computational chemistry as a helpful tool for structure-functional study and correlation, en route to protein engineering |
| L17 | <i>Rudiger Ettrich</i> Structure and dynamics of the oxygen evolving complex of photosystem II: Role of the N-terminal loop |
| L18 | <i>Ondřej Šulák</i> New approaches to structure and function studies of RS20L lectin from <i>Ralstonia solanacearum</i> |



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| L19 | p. 24 | Saturday, March 31 |
| Navnit Mishra | | |
| Computational Studies on PA-IIL Lectin-Carbohydrate Interactions | | 7:00 - 9:10 Breakfast |
| 15:20 - 15.30 Coffee break | | |
| 15:30 - 17:25 Lecture session VI | | |
| chairperson: <i>I. Kratochvílová</i> | | |
| L20 | p. 25 | 9:15 - 10:15 Lecture session VII chairperson: <i>R. Zika</i> |
| <i>Bohdan Schneider</i> | | L23 p. 27 |
| Conformational space of nucleic acids | | <i>Irena Kratochvílová</i> |
| L21 | p. 26 | New principle of the molecular photo-FET |
| <i>Dana Nachtigallová</i> | | L24 p. 27 |
| Electronic coupling in the excited electronic state of stacked DNA bases | | <i>Jan Pačes</i> |
| L22 | p. 12 | CHIPAS, Tool for ChIPonChip analysis |
| <i>Dalibor Štys</i> | | L25 p. 27 |
| Mathematically-based studies of complex systems in physics, mathematics, chemistry, biology, economy and politics in Nove Hrady in the first half of 19th century and at the beginning of 21st century | | <i>Eva Brumovská</i> |
| <i>Radomír Kužel</i> : Forthcoming meetings of the Czech and Slovak Crystallographic Association - short information. Correct typesetting of papers - general rules on correct typing and writing references. | | Structural and functional characterization of human protein kinase CKI epsilon |
| Future evaluation of publications. | | |
| <i>Peter Šebo</i> | | 10:15 - 10:30 Coffee break |
| Progress in building if the Biotechnology center in Vestec | | |
| 17:30 - 18:30 Dinner | | 10:30 - 11:40 Lecture session VIII , chairperson: <i>T. Obšil</i> |
| 19:00 - 22:00 Poster session | | L2 p. 28 |
| | | <i>Jan Dohnálek</i> |
| | | Subtilases and metal binding - the weak binding site of subtilisins revisited |
| | | L27 p. 28 |
| | | <i>Jindřich Hašek</i> |
| | | Ligand docking to biochemical targets - crystallography and modeling |
| | | L28 p. 30 |
| | | <i>Pavlína Řezáčová</i> |
| | | Structural studies of the repressor CggR from <i>Bacillus subtilis</i> |
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| | | 11:40 - 12:00 Concluding remarks |
| | | 12:00 - 13:00 Lunch |

POSTERS

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| P1 | p. 31 | P5 | p. 33 |
| <i>L. Benda</i> | | <i>J. Chmelík</i> | |
| Calculation of the Effective Chemical Shielding Anisotropy in L-Alanyl-L-alanine, conformational and charge dependence study | | Structure of <i>Bombyx mori</i> Chemosensory Protein 1 in Solution | |
| P2 | p. 31 | P6 | p. 33 |
| <i>I. Beššeová</i> | | <i>E. Fadrná</i> | |
| Computational simulations study of Helixes 90-92 | | Computational study of restriction endonuclease hincii | |
| P3 | p. 32 | P7 | p. 34 |
| <i>E. Bouřa</i> | | <i>Z. Futera</i> | |
| Crystal structure of RGS3 RGS domain | | Computational Study on Ruthenium(II) Complexes and their Interactions with NA bases | |
| P4 | p. 32 | P8 | p. 34 |
| <i>L. Bumba</i> | | <i>L. Gorčík</i> | |
| Cryo-scanning electron microscopy of <i>Bordetella pertussis</i> adenylate cyclase toxin embedded in lipid vesicles | | NMR study of protein CD6 | |



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| P9 <i>L. Gryčová</i> Lys735 participates on the ATP binding to the isolated C-terminus TRPV1 | p. 34 | P21 <i>J. Matulová</i> A cost-effective approach for amino-acid-type selective isotope labeling of proteins expressed in <i>Leishmania tarentolae</i> | p. 41 |
| P10 <i>D. Ham</i> Structure – function relationships of protein-DNA interactions | p. 35 | P22 <i>L. Michera</i> Theoretical Study of Cisplatin Interactions with Glycine in Gas-phase and Implicit Water Solution – COSMO | p. 41 |
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**Conference is co-financed by project MATEO,
sub-project E-clue of the INTERREG III C program of
European Union**

