

# XVIII Discussions in Structural Molecular Biology and 5th User Meeting of CIISB

24-26 March 2022

## Programme

### Thursday, March 24

11:30 - 13:30 Registration and accommodation, coffee

All talks are 20 minutes long unless indicated otherwise. All times are INCLUDING discussion.

14:00 **Opening of XVIII Discussions and 5th User Meeting of CIISB** (*Jan Dohnálek, Vladimír Sklenář*)

14:15 - 16:05 **Session I** *Chair person: Jan Dohnálek*

*Lecture no. – page of abstract*

<b>Zdeněk Lánský</b> Institute of Biotechnology, Czech Academy of Sciences	Tau generates protective envelopes on microtubules <span style="float: right;">L1 – 3</span>
<b>Miroslav Kloz</b> Institute of Physics, Czech Academy of Sciences	Structural changes of carotenoid echinenone in Orange Carotenoid Protein studied by femtosecond Raman spectroscopy <span style="float: right;">L2 – 4</span>
<b>Rohit Joshi</b> Institute of Physiology, Czech Academy of Sciences	Nedd4-2 binding to 14-3-3 modulates the accessibility of its catalytic site and WW domains (Student talk) <span style="float: right;">L3 – 4</span>
<b>Aneta Kozeleková</b> CEITEC, Masaryk University	Phosphomimicking Mutations ≠ Phosphorylation – a case study of 14-3-3 protein (Student talk, CIISB user meeting) <span style="float: right;">L4 – 5</span>
<b>Mark Johnson / Lukáš Gajdoš</b> Institut Laue-Langevin, Grenoble	Neutrons for Structural Biology at the Institut Laue Langevin / Neutron diffraction for deciphering lectin-carbohydrate interactions in bacterial infection (30 min) <span style="float: right;">L5 – 6</span>

16:05 - 16:35 *Coffee break*

16:35 - 18:25 **Session II** *Chair person: Pavlína Maloy Řezáčová*

<b>Ján Bíňovský</b> CEITEC, Masaryk University	Baseplate structure of bacteriophage phi812 and mechanism of cell wall binding and penetration (Student talk, CIISB user meeting) <span style="float: right;">L6 – 7</span>
<b>Zuzana Trebichalská</b> CEITEC, Masaryk University	<i>In situ</i> cryo-electron tomography of enterovirus cell entry (Student talk, CIISB user meeting) <span style="float: right;">L7 – 7</span>
<b>Miroslav Homola</b> CEITEC, Masaryk University	Mechanism of virion formation of the <i>Emiliana huxleyi</i> virus 201 enveloped by two membranes (Student talk, CIISB user meeting) <span style="float: right;">L8 – 8</span>
<b>Tomáš Koval'</b> Institute of Biotechnology, Czech Academy of Sciences	Bacterial helicase-like transcription-associated factor HelD (CIISB user meeting) <span style="float: right;">L9 – 8</span>
<b>Sebastian Zoll</b> Institute of Organic Chemistry and Biochemistry, Czech Academy of Sciences	The structural basis of complement inhibition by the human parasite <i>Trypanosoma brucei gambiense</i> <span style="float: right;">L10 – 9</span>
<b>Piotr Wardęga</b> NanoTemper Technologies	Characterize your most challenging interactions. The New Monolith (10 min) <span style="float: right;">L11 – 9</span>

18:30 Dinner

20:00 **CIISB Executive committee**

Friday, March 25

7:30 - 9:00 *Breakfast*

9:00 - 10:50 **Session III** *Chair person: Lukáš Židek*

<b>Edward Curtis</b> Institute of Organic Chemistry and Biochemistry, Czech Academy of Sciences	Supernova: a deoxyribozyme that catalyzes a chemiluminescent reaction <i>L12 – 11</i>
<b>Maroš Huličiak</b> Institute of Biotechnology, Czech Academy of Sciences	New Class of Structurally Robust Non-Antibody Protein Scaffolds for Directed Evolution (Student talk) <i>L13 – 11</i>
<b>Jan Příbyl</b> CEITEC, Masaryk University	Atomic force microscopy in structural biology (CIISB user meeting) <i>L14 – 12</i>
<b>Petr Pompach</b> Institute of Biotechnology, Czech Academy of Sciences	Structural Mass Spectrometry – An Advanced Tool in Protein Structure Analysis (CIISB user meeting) <i>L15 – 13</i>
<b>Vilhelmiina Kontkanen</b> Faculty of Science, University of South Bohemia	Structural Changes and Their Consequences for Azurin Oxidation in Vacuum and on Gold Interfaces <i>L16 – 13</i>
<b>Patrick King</b> SPECION, s.r.o.	Microfluidic Modulation Spectroscopy (MMS): A Novel IR-Based Technique Providing Automated, Highly-Sensitive Protein Secondary Structure Characterisation in <i>in situ</i> Conditions (10 min) <i>L17 – 14</i>

10:50 - 11:10 *Coffee break*

11:10 - 12:50 **Session IV** *Chair person: Pavel Plevka*

<b>Kristýna Adámková</b> Institute of Biotechnology, Czech Academy of Sciences	Zinc-dependent nuclease from <i>Stenotrophomonas maltophilia</i> : Structural analysis of the active site (Student talk, CIISB user meeting) <i>L18 – 15</i>
<b>Ondrej Cehlár</b> Institute of Neuroimmunology, Slovak Academy of Sciences	Biophysical characterization of novel monoclonal antibodies targeting epitopes on the SARS-CoV-2 Spike protein (CIISB user meeting) <i>L19 – 16</i>
<b>Martin Malý</b> Faculty of Nuclear Sciences and Physical Engineering, Czech Technical University	Integrative structural analysis of antibiotic-inactivating enzyme from <i>Stenotrophomonas maltophilia</i> (Student talk, CIISB user meeting) <i>L20 – 17</i>
<b>Michal Koblížek</b> Institute of Microbiology, Centre Algatech, Czech Academy of Sciences	2.4 Å structure of the double concentric ringed light harvesting complex from phototrophic bacterium <i>Gemmatimonas phototrophica</i> <i>L21 – 18</i>
<b>Zuzana Hlavenková</b> Thermo Fisher Scientific	Thermo Scientific™ Tundra Cryo-TEM: 100kV Cryo-TEM dedicated for Single Particle Analysis (10 min) <i>L22 – 18</i>
<b>Poster flash talks, 2 minutes each</b>	
<b>Alexandra Náplavová</b> (Faculty of Science, Masaryk University) Multiple approaches for protein phosphorylation: a story of 14-3-3 (Student poster)	
<b>Jan Šimek</b> (Faculty of Science, Masaryk University) To homodimerize or to heterodimerize: story of 14-3-3 protein dimer formation (Student poster)	
<b>Pavel Pohl</b> (Institute of Physiology, Czech Academy of Sciences) 14-3-3-protein regulates Nedd4-2 by modulating interactions between HECT and WW domains (Student poster)	
<b>Olivia Petrvalská</b> (Institute of Physiology, Czech Academy of Sciences) 14-3-3 directly interacts with the kinase domain of CaMKK1 and inhibits calmodulin binding	

12:50 - 14:00 *Lunch*

14:00 - 15:50 **Session V** Chair person: **Vojtěch Spiwok**

<b>Martin Lepšík</b> Institute of Organic Chemistry and Biochemistry, Czech Academy of Sciences	On the Importance of Physically Correct Models for Protein-Ligand Binding <i>L23 – 19</i>
<b>Michal Malý</b> Institute of Biotechnology, Czech Academy of Sciences	"ReDNATCO" - making DNATCO more useful for crystallographers <i>L24 – 20</i>
<b>Ladislav Bartoš</b> Faculty of Science, Masaryk University	Designing transmembrane proteins to enhance transport of peptides across cell membranes (Student talk) <i>L25 – 21</i>
<b>Zdenek Futera</b> Faculty of Science, University of South Bohemia	Mechanisms of Charge Transfer through Multiheme Protein Junctions, Their Distance and Band-Alignment Dependencies <i>L26 – 22</i>
<b>Lubomír Rulišek</b> Institute of Organic Chemistry and Biochemistry, Czech Academy of Sciences	Exploring Reaction Mechanisms of Metalloproteins by Correlating Theory and Experiment <i>L27 – 23</i>
<b>Martin Máša</b> Beckman Coulter, Life Sciences Sales Dpt	EMnetik 24 System - DNA Cleanup for Genetic Engineering (10 min) <i>L28 – 24</i>

15:50 - 16:10 *Coffee break*

16:10 - 18:00 **Session VI** Chair person: **Tomáš Koval'**

<b>Martin Marek</b> Faculty of Science, Masaryk University	Can misfolded enzymes be beneficial? Yes, they can <i>L29 – 25</i>
<b>Jitka Plucarová</b> Faculty of Science, Masaryk University	Interactions of adaptor protein Grb2 with microtubule associated protein 2c (Student talk, CIISB user meeting) <i>L30 – 25</i>
<b>Radim Novotný</b> Faculty of food and biochemical technology, University of Chemistry and Technology	Structure and biological functions of TBEV Capsid protein (Student talk) <i>L31 – 26</i>
<b>Raju Mandal</b> Faculty of Science, Charles University	Biophysical characterization of the FOXO4:p53 complex (Student talk) <i>L32 – 27</i>
<b>Dávid Tužinčín</b> CEITEC, Masaryk University	Seeing the invisible – Study of transiently formed protein conformation found in domain 1.1 of bacterial transcription factor (Student talk, CIISB user meeting) <i>L33 – 28</i>
<b>Poster flash talks, 2 minutes each</b>	
<p><b>Andreas Santamaria</b> (Institut Laue-Langevin, Grenoble) Strikingly different roles of SARS-CoV-2 fusion peptides uncovered by neutron scattering (Student poster)</p> <p><b>Lucie Valentová</b> (CEITEC, Masaryk University) A hunting strategy and virion structure of <i>P. aeruginosa</i> bacteriophage JBD30 revealed by cryo-electron microscopy (Student poster)</p> <p><b>Peter Pajtinka</b> (CEITEC, Masaryk University) Synergistic antimicrobial activity of magainin 2 and PGLa revisited (Student poster)</p> <p><b>Patricia Hrašnová</b> (Institute of Neuroimmunology of Slovak Academy of Sciences) Structural and Biophysical Aspects of Lactoferrin and Its Interaction with Plasminogen (Student poster)</p> <p><b>Andrej Bitala</b> (Institute of Virology, Biomedical Research Center Slovak Academy of Sciences) Immunoprecipitation pull-down assay revealed binding between human CD160 and viral UL144 (Student poster)</p>	

18:00 - 19:00 **General Assembly of the Czech Society for Structural Biology**

The meeting will be held in the Czech and/or Slovak languages

1. Volba zapisovatele
2. Schválení programu
3. Výroční zpráva Rady ČSSB o činnosti a hospodaření
4. Zpráva revizní komise
5. Schválení výroční zprávy Rady ČSSB o činnosti a hospodaření
6. Infrastruktury se vztahem ke strukturální biologii, přístupy, financování, vývoj Instructu a jiných
7. Různé

19:00 - 20:00 *Dinner*

20:00 - 22:00 **Poster session**

## Saturday, March 26

7:30 - 9:00 *Breakfast*

9:00 - 10:20 **Session VII** *Chair person: Jiří Pavlíček*

<b>Václav Veverka</b> Institute of Organic Chemistry and Biochemistry, Czech Academy of Sciences	A ubiquitous disordered protein interaction module orchestrates transcription elongation  <i>L34 – 28</i>
<b>Zuzana Cieniková</b> CEITEC, Masaryk University	Structures of phage 812 neck suggest a mechanism for genome retention and release in <i>Herelleviridae</i> (CIISB user meeting)  <i>L35 – 29</i>
<b>Markéta Šoltysová</b> Institute of Organic Chemistry and Biochemistry, Czech Academy of Sciences	How do SorC proteins recognize their DNA operators? (Student talk)  <i>L36 – 29</i>
<b>Vladena Bauerová</b> Institute of Molecular Biology, Slovak Academy of Sciences	Structure-functional insights into the dantrolene binding site of the human cardiac ryanodine receptor: towards a deeper understanding of heart arrhythmias  <i>L37 – 30</i>

10:20 - 10:40 *Coffee break*

10:40 - 12:20 **Session VIII** *Chair person: Veronika Obšilová*

<b>Dalibor Košek</b> Institute of Physiology, Czech Academy of Sciences	Mechanism of replicative "rolling-circle" DNA transposition in eukaryotes  <i>L38 – 31</i>
<b>Robert Vácha</b> CEITEC, Masaryk University	How Viruses and Virus-like Nanoparticles Can Release Their Cargo/Genome (CIISB user meeting)  <i>L39 – 32</i>
<b>Tibor Füzik</b> CEITEC, Masaryk University	Structure of tick-borne encephalitis virus immature particle solved by cryo-electron microscopy and sub-tomogram averaging (CIISB user meeting)  <i>L40 – 33</i>
<b>Richard Štefl</b> CEITEC, Masaryk University	Cryo-EM of mouse RNase III–RNA complexes (CIISB user meeting)  <i>L41 – 34</i>
<b>Gabriel Demo</b> CEITEC, Masaryk University	Cryo-EM ensemble reveals the mechanism of +1 ribosomal frame shifting  <i>L42 – 35</i>

12:20 - 12:40 **Evaluation of student presentations, prizes, concluding remarks**

12:40 *Lunch & departure*