

Scientific program

Sunday

14:00–17:00	Registration	
17:00–17:15	Conference opening	
17:20–18:05	P. Hobza	Noncovalent Interactions: Theory and Applications
19:00–21:30	Welcome party	

Monday

chairman	<i>W. A. Sokalski</i>	
9:00–9:30	L. Gorb	DNA bases in Dickerson dodecamer: results of recent quantum-chemical calculations
9:30–10:00	P. Jurečka	Using QM Data in Force Field Development for RNA and DNA Simulations
10:00–10:30	J. Šponer	Computational studies of nucleic acids – the current trends.
	Coffee break	
chairman	<i>X. Assfeld</i>	
11:00–11:30	C. Lim	Principles Governing Biological Processes: Applications to drug design and drug target identification
11:30–12:00	T. Dudev	Determinants of Fe ²⁺ over M ²⁺ (M = Mg, Mn, Zn) Selectivity in Non-Heme Iron Proteins
12:00–12:30	N. Gresh	Polarizable molecular mechanics/dynamics potentials. Validations by QC computations on two sensitive issues: stacked guanine quartets and dizinc metalloenzyme binding sites.
	Lunch	
chairman	<i>P. Hobza</i>	
14:20–14:50	G. Mazzone	Computational Contribution to the Design of Effective Drugs in Photodynamic Therapy
14:50–15:10	S. Mutter	Metallobiology of Amyloid Beta Peptides by Ligand Field Molecular Mechanics
15:10–15:30	D. Bonhenry	Investigation of Ca ²⁺ permeation in the Orail channel.
15:30–15:50	K. Mazmanian	Preferred Hydrogen-Bonding Partners of Cysteine: Implications for Regulating Cys Functions
	Coffee break	
chairlady	<i>C. Lim</i>	
16:20–16:50	W. A. Sokalski	Use of catalytic fields in biocatalyst design
16:50–17:20	J. Poater	How non-terran bio-solvents affect the structure and stability of B-DNA
17:20–17:40	R. Matute	Revisiting the Mechanism of DNA Polymerases with an Empirical Valence Bond Model
17:40–18:00	M. Růžička	Study of Structural and Elastic Properties of DNA Mutation Motifs Using Computational Methods
19:00–21:00	Poster session	

Tuesday**chairman***R. Ettrich*

9:00–9:30	T. Clark	Metadynamics Simulations of G-Protein Coupled Receptors
9:30–10:00	M. Hall	Modelling Hydrogenase Biomimetics
10:00–10:30	N. Russo	Enzyme promiscuity: A Computational Approach

Coffee break**chairman***L. Gorb*

11:00–11:30	P. Paneth	Binding isotope effects in design of HIV-1 reverse transcriptase inhibitors
11:30–12:00	S. Zaric	Modeling and elucidating role of aromatic and aliphatic side chain interactions in stability of amyloids
12:00–12:30	V. Moliner	Revealing the Origin of Enzyme Catalysis from Computational Studies

Lunch**chairman***T. Clark*

14:20–14:50	J. Kozelka	Anion- π interactions in flavoproteins involve a substantial charge-transfer component
14:50–15:20	P. Cysewski	Water solubility advantage of drugs cocrystallized with pharmaceutically acceptable cofomers.
15:20–15:50	D. Řeha	Application of Charge Transfer Calculations on Computational Study of Proteins

Coffee break**chairman***P. Paneth*

16:20–16:50	R. Ettrich	Computational Modeling of 3'-Phosphoadenosine 5'-Phosphosulfate Synthase PAPSS
16:50–17:20	T. Wesolowski	
17:20–17:50	J. Urban	MP1- membrane interactions, MD study

19:30 **Concert in the Cathedral of St. Bartholomew**Performed by **Pavel Šmolík**.**Compositions:**

- Georg Muffat: Toccata prima (5')
- Johann Pachelbel: Ciacona in f (9')
- Johann Sebastian Bach: Toccata d moll (9')
- Johann Sebastian Bach: O mensch, bewein (5')
- Felix Mendelssohn-Bartholdy: Sonata c moll (Grave, Andante, Allegro Moderato, Fuga) (11')
- Pavel Šmolík: Improvizace (8')

Wednesday**chairman***T. Wesolowski*

- 9:00–9:30 K. Ruud Modeling one- and multiphoton absorption in biomolecular systems
- 9:30–10:00 P. Bouř Computations of Magnetic Circular Dichroism in Molecules
- 10:00–10:30 M. J. Ramos Advances and Pitfalls in Computational Enzymatic Catalysis

Coffee break**chairman***S. Roszak*

- 11:00–11:30 A. Michalak ETS-NOCV decomposition of the reaction force
- 11:30–12:00 X. Assfeld Modeling Excited States Interactions in Biomolecules.
- 12:00–12:30 H. Dong Exploring the Gating Mechanism of the Calcium Release-Activated Calcium Channel with Molecular Dynamics Simulations

Lunch

- 14:30–16:00 **Sightseeing tour in Pilsen**
- 16:00–18:00 **Excursion to the Pilsner brewery**
- 18:00 **Dinner in the brewery restaurant**

Thursday**chairman** *N. Gresh*

- 9:00–9:30 B. Szefczyk AIMD analysis of model protic ionic liquids
- 9:30–10:00 R. Friedman Computer simulations to improve the understanding of drug resistance in cancers
- 10:00–10:30 S. Roszak The breathing MOF's – chemistry of MIL-53 from first principles

Coffee break**chairman** *J. Kozelka*

- 11:00–11:30 S. Zálíš Modelling of electron transfer in metal labelled proteins and systems containing multiple redox centers
- 11:30–12:00 H. Dos Santos Predicting standard reduction potential for anticancer Au(III)-complexes: A DFT study
- 12:00–12:30 M. Srnec Computational Electrochemistry of Mononuclear Non-Heme Iron Complexes: Redox Properties and their Contributions to Reactivity

Lunch**chairman** *P. Bourř*

- 14:20–14:50 A. Wierzbicki A modeling study of the binding of novel indenenes to the catalytic site of mutant KRAS
- 14:50–15:10 T. Jeliński Application of COSMO-RS methodology for screening of ionic liquids and deep eutectic solvents as effective media for enhancement of extraction and solubility
- 15:10–15:30 Z. Chval Bonding and non-bonding interactions of Pt(II)-complexes
- 15:30–15:50 F. Šebesta Reduction of Pt(IV) complexes by small biomolecules

Coffee break**chairman** *M. Hall*

- 16:20–16:50 T. Ishida Computational modeling of carbohydrate recognition in protein complex
- 16:50–17:20 W. Rocha Release of NO from Ruthenium Nitrosyl Complexes in Aqueous Solution.
- 17:20–17:40 J. Novotný Revealing structure of i-motif with purine stretches using NMR and molecular modelling
- 17:40–18:00 J. Jungwirth Quantitative Determination of Ala-Ala Conformer Ratios by Decomposition of Raman Optical Activity Spectra

Friday

chairman	<i>S. Grabowski</i>
9:00–9:30	M. Biczysko Understanding structure and function of bio-molecular systems from spectroscopic 'fingerprints'
9:30–10:00	Z. Futera Heterogeneous Electron Transfer Between Gold Surface and Adsorbed Deca-Heme Protein MtrF
10:00–10:30	R. Marek Supramolecular Carriers for Anticancer Metallo drugs
Coffee break	
chairman	<i>B. Szeferczyk</i>
11:00–11:30	T. Mančal Ten years of coherent quantum biology: Paradigm change that almost happened
11:30–12:00	S. Grabowski Dual character of boron as a Lewis acid center – ab initio calculations and QTAIM analysis
12:00–12:30	Closing ceremony

Poster presentations

Poster no.	Author name	Poster title
1	E. Ahlstrand	Force Field Based Calculations of Zn Interaction Energies in Proteins
2	P. L. Bora	Supramolecular Covalence in Bifurcated Chalcogen Bonding
3	T. Bouchal	Influence of non-Watson-Crick base pairing on structure of DNA
4	R. Cajzl	Excited state dynamics of conjugated polyenes Ab-initio and semiempirical calculation of electronic absorption spectra
5	D. Cheshmedzhieva	Competition between Abiogenic Al^{3+} and Native Mg^{2+} , Fe^{2+} and Zn^{2+} Ions in Protein Binding Sites: Implications for Aluminium Toxicity
6	B. J. Cuyacot	Calculation of Pt-195 NMR Chemical Shifts for Pt(IV)-Based Metallo-drugs
7	P. Czeleń	Docking and molecular dynamics study of CDK2 inhibition by new indoline derivatives
8	O. Dvořáčková	The effect of substituents on the kinetics of the hydration reactions of trans-Platinum complexes
9	H. Georg	Elucidating the structure of merocyanine dyes with the ASEC-FEG method
10	W. Jedwabny	Testing a simple non-empirical scoring model on inhibitors of EphA2-ephrinA1 protein-protein interaction
11	N. Kircheva	Gallium as a Therapeutic Agent: A Thermodynamic Evaluation of the Competition between Ga^{3+} and Fe^{3+} Ions in Metalloproteins
12	P. Krawczyk	The reactive group effects on the photophysical and biological properties of the fluorescent probe based on chalcone derivatives
13	M. Krupová	Circularly Polarized Luminescence in Study of Protein Folding – Experimental and Theoretical Study
14	E. Kutálková	Molecular-dynamics simulations of hyaluronan-ions interactions
15	S. Malali	Paramagnetic NMR study of Ru(III) complex for calculating NMR chemical shift
16	M. Melicherčík	The influence of site Mutations on Activity of XPB Protein
17	M. Misini Ignjatovic	Structure and thermodynamics of water at the binding site of galectin-3 with two diastereomers of different affinities
18	V. Nikolova	Determinants of the host-guest interactions Q1 between α -, β - and γ -cyclodextrins and group IA, IIA and IIIA metal cations: a DFT/PCM study
19	I. Niksic-Franjic	Reactions of free radicals with haloorganics in buffered aqueous solutions: modeling proton-coupled electron transfer and competing mechanisms
20	J. Petrova	Assessment of the Effect of Oligomer-Water Interaction on Polyaniline Properties
21	M. Pizl	Vibrational spectroscopy of metal complexes derived from vitamin B6: Experimental and DFT study
22	L. Plačková	Theoretical study of PAPSS2 kinase domain
23	T. Smetanová	Estimation of Ir(III) empirical parameters for molecular mechanical force fields
24	Z. Sochorová Vokáčová	Theoretical study of a lipase-catalyzed reaction in organic solvent
25	B. Szeffler	Theoretical Study on Cisplatin Binding to B-Vitamins: Could the therapeutic effect of cisplatin be decreased if a patient with lung cancer drinks carrot or beet juice?
26	M. Turner	Modelling $Pt^{II}(\text{Ligand}) - \text{Amyloid-}\beta$ Interactions: Prediction of Ligand Effects
27	M. Zgarbová	Recent Force Fields for RNA Simulations – Performance, Comparison and Challenges