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**Sunday, June 19<sup>th</sup>**

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9:50 City Walk (only for registered participants)  
13:30 – 15:30 Registration  
15:00 Welcome snack  
15:20 Meeting opening

Session: Nucleic Acids I - Chair: Tom Cheatham

15:30 **Modesto Orozco**  
*Simulating DNA from the electron to the chromosome*

16:10 **Montgomery Pettitt**  
*DNA confined: Advantages of non-linear behavior*

16:50 Coffee break

17:10 **Mahdi Bagherpoor Helabad**  
*The role of Protein-DNA interactions in the DNA binding specificity of hormone receptors*

17:30 **Lennart Nilsson**  
*Triplex forming oligonucleotides and gene therapy*

18:10 **Vlad Cojocaru**  
*From decoding dynamics to tailoring cooperativity in protein-DNA interactions*

19:00 Dinner at Terminus hotel (only for registered participants)

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**Monday, June 20<sup>th</sup>**

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Session: Nucleic Acids II - Chair: Lennart Nilsson

8:30 **Thomas E. Cheatham III**  
*Challenges and successes in modeling RNA structure, dynamics and interactions with ions.*

9:10 **Joanna Trylska**  
*Interactions of aminoglycoside antibiotics with RNA*

9:30 **Filip Lankas**  
*Modelling allosteric effects in DNA*

9:50 Coffee break

Session: protein-lipid interactions - Chair: Michele Cascella

10:10 **Carmen Domene**  
*Computational approaches to the molecular thermometers of the human body*

10:50 **Syma Khalid**  
*Molecular simulations that put the chemical complexity into model bacterial membranes*

11:30 **Stefano Vanni**  
*Taking advantage of membrane diversity in intracellular trafficking pathways*

11:50 **Michael Schauperl**  
*Explaining the recognition process of ice binding proteins*

12:10-13:30 Lunch Break

Session: ISQBP awards talks - Chair: Alex MacKerell

13:30 **Rebecca Wade** - ISQBP Award in Computational Biology

*In silico prediction of biomolecular recognition*

14:30 **Marta Filizola** - ISQBP Loew Lectureship

*New Challenges and Opportunities in G Protein-Coupled Receptor Drug Discovery*

15:30 Coffee Break and **Poster session**

17:40 Recommended departure to Welcome ceremony (by foot)

18:00 Welcome ceremony by the City of Bergen (at the University Aula, Cf. map)

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**Tuesday, June 21<sup>st</sup>**

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Session: Drug design I - Chair: Annick Dejaegere

8:30 **William L.Jorgensen**

*Design and Discovery of Potent Enzyme Inhibitors*

9:10 **Daniel Cappel**

*Rigorous Free Energy Calculations Applied to Protein Homology Models*

9:30 **Elsa Sánchez-García**

*Supramolecular Ligands as Regulators of Biomolecular Interactions*

9:50 Coffee Break

Session: Drug design II - Chair: Bjørn Olav Brandsdal

10:10 **Carmay Lim**

*Using an old drug to target a new drug site*

10:50 **Sanja Zivanovic**

*Multi-level strategy for analysis of bioactive drug conformations*

11:10 **Chandra Verma**

*Stapled diets: tackling resistance*

11:30 ISQBP General Assembly

12:00-13:30 Lunch Break

Session: Methodological Advances I - Chair: William L. Jorgensen

13:30 **Alex MacKerell**

*Overview of the Classical Drude Oscillator Polarizable Force Field for Biomolecules*

14:10 **Charles L. Brooks III**

*High-throughput, free energy based ligand discovery and optimization using multi-site  $\lambda$ -dynamics*

14:50 **Hima Bindu Kolli**

*Hybrid Particle-Field Approach with Electrostatics for Mesoscale Biomolecular Simulations*

15:10 **Iain Bethune**

*Molecular Integration Simulation Toolkit - interfacing novel integrators with Molecular Dynamics codes.*

15:30 Coffee Break

18:30 Recommended departure from Grand Hotel Terminus (by foot, Cf. map)

19:00 Boat departure to *Cornelius på Holmen*, Conference Dinner, sponsored by the Journal of Chemical Theory and Computation (JCTC)

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**Wednesday, June 22<sup>nd</sup>**

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Session: Methodological Advances II – chair: Vlad Cojocaru

8:50 **Thomas Gaillard**

*Computational Protein Design with an MMGBSA Energy Function*

9:10 **Kwangho Nam**

*Development and acceleration of multiscale QM/MM methods for simulations of complex biomolecular systems*

9:30 **Xabier Lopez**

*Aluminum and bioligand interactions with phosphate containing groups.*

9:50 Coffee Break

Session: Enzyme and protein dynamics – chair: Nathalie Reuter

10:10 **Annick Dejaegere**

*Structural dynamics and signalling mechanisms in nuclear retinoid receptors*

10:50 **Carol Post**

*An unusual allosteric mechanism for regulating protein interactions of Syk tyrosine kinase*

11:30 **Natacha Gillet**

*Understanding Charge Transfer in Cryptochromes and Photolyases via QM/MM Simulations: Application to PhrB Protein*

11:50 **Thibault Tubiana**

*Norovirus Capsid Assembly*

12:10 Closing remarks

12:30 Lunch