



MGMS and RSC MMG Young Modellers' Forum 2011

Presentations

9.00 – 9.30	Coffee and Registration
9.30 – 9.40	Welcome and Introduction Steve Maginn and Louise Birch
9.40 – 10.00	Flexibility Controls Specificity of Snake Venom Metalloproteases Hannes Wallnoefer, <i>University of Innsbruck, Austria</i>
10.00 – 10.20	Flipping Inositol Phosphates: A Molecular Dynamics Approach to Understanding the Selectivity of Protein Kinase B Sarah Rosen, <i>Imperial College London</i>
10.20 – 10.40	Protein-ligand binding free energies from large-scale DFT calculations with the ONETEP program Stephen Fox, <i>University of Southampton</i>
10.40 – 11.00	Combination of structure- and ligand-based virtual screening for the discovery of novel non-LBP antiandrogens as chemical tools to treat prostate cancer Laura Caboni, <i>Trinity College Dublin, Ireland</i>
11.00 – 11.20	Atomic Detail Studies of P-Glycoprotein and Drug Permeation in Model Membranes Jerome Ma, <i>University of Oxford</i>
11.20 – 11.40	From Development to Application: Molecular Mechanical Study of Halogen Bonding in Drug Discovery Mahmoud Ibrahim, <i>University of Manchester</i>
11.40 – 13.30	Lunch and Poster Session
13.30 – 13.50	Druggability and structural analysis of bromodomain acetylated-lysine binding sites Lewis Vidler, <i>Institute of Cancer Research</i>
13.50 – 14.10	Advances in Free Energy Calculation: The Enveloping Distribution Sampling Method Sereina Riniker, <i>ETH Zurich, Switzerland</i>
14.10 – 14.30	Metabolite Space and Metabolite-Likeness Julio Peironcely, <i>Leiden University/TNO, The Netherlands</i>
14.30 – 15.00	Tea
15.00 – 15.20	A Ligand-Assisted Proton Shuttle (LAPS) Mechanism: Using Coordinated Acetate Ligands to Affect Chemical Change David Johnson, <i>University of York</i>
15.20 – 15.40	Interpreting Experimental Analysis of the Human Ether-a-go-go Related Gene Product (hERG) with Molecular Dynamics

	and <i>in-silico</i> Docking Charlotte Colenso, <i>University of Bristol</i>
15.40 – 16.00	A Novel Computational Approach to Fragment-Based Drug Discovery Michael Bodnarchuk, <i>University of Southampton</i>
16.00	Fun event – “Name That Molecule”
16.30	Judges Deliberations
16.45	Prize Presentations
17.00	End

Posters

Poster 1	Revisiting the General Solubility Equation: <i>In Silico</i> Prediction of Aqueous Solubility incorporating the Effect of Topographical Polar Surface Area Jogoth Ali, <i>University of Hertfordshire</i>
Poster 2	The Development and Application of a simplified QM/MM method for Free Energy Simulations Michael Carter, <i>University of Southampton</i>
Poster 3	Visualization and Analysis of Helix Flexibility Using Bendix Caroline Dahl, <i>University of Oxford</i>
Poster 4	Ligand- and Structure-Based Computational Approaches for the Design of Potent and Selective Compounds towards Adenosine Receptors. David Rodríguez, <i>Unilever Centre: University of Cambridge</i>
Poster 5	Rapid Prediction and Scoring of Water Molecules in Protein Binding Sites Gregory Ross, <i>University of Oxford</i>
Poster 6	The Assessment of Computationally Derived Protein Ensembles in Protein-Ligand Docking Barbara Sander, <i>University of Southampton</i>
Poster 7	Actinyl reduction potentials in solution: An assessment of computational methods to achieve experimental accuracy Krishnamoorthy Arumugam, <i>University of Manchester</i>
Poster 8	The Relationship between Drug-Lipid Interaction Energy and Non-Specific Binding Callum Dickson, <i>Imperial College London</i>
Poster 9	Understanding the Competing Reactions of Polyurethane Foam Formation: A Computational Study Jack Gibb, <i>Unilever Centre: University of Cambridge</i>
Poster 10	Exploring the mycolic acid potential energy surface: the ups and downs Wilma Groenewald, <i>Bangor University</i>
Poster 11	In silico target predictions using machine-learning methods: Naïve Bayes & Parzen-Rosenblatt Window Alexios Koutsoukas, <i>Unilever Centre: University of Cambridge</i>
Poster 12	An <i>In Silico</i> Scaffold Hopping Protocol for Identifying Novel Kinase Inhibitors Sarah Langdon, <i>Institute of Cancer Research</i>