

# Young Modellers' Forum 2005

## Presentations

9.00 – 9.30	<b>Coffee and Registration</b>
9.30 - 9.40	<b>Welcome and Introduction</b> Steve Maginn and Stuart Firth-Clark
9.40 – 10.00	<b>Evolution of metal binding sites</b> James Torrance - <i>European Bioinformatics Institute</i>
10.00 – 10.20	<b>Using multiobjective optimization to study the strengths of different interaction energies in protein-ligand complexes</b> Sally Mardikian - <i>University of Sheffield</i>
10.20 – 10.40	<b>Simulation of conformational transitions and Free Energy Calculations of PcrA DNA Helicase by Targeted Molecular Dynamics and WHAM</b> Hao Wang - <i>University of Nottingham</i>
10.40 – 11.00	<b>Drug binding to the hERG K<sup>+</sup> channel: a molecular modelling study</b> Phillip J. Stansfeld - <i>University of Leicester</i>
11.00-11.20	<b>Modelling drug metabolism in cytochrome P450 enzymes</b> Jolanta Zurek - <i>University of Bristol</i>
11.20-11.40	<b>Online Drug Design</b> Alasdair Laurie - <i>University of Leeds</i>
11.40-13.30	<b>Lunch and Poster Session</b>
13.30 – 13.50	<b>Molecular Surface Property Graphs</b> Vishwesh Venkatraman - <i>University of Portsmouth</i>
13.50 – 14.10	<b>Where do the protons go: Protonation states in protein-ligand complexes</b> Paul Czodrowski - <i>Philipps University Marburg</i>
14.10 – 14.30	<b>The Role of Water in Protein-Ligand Interactions: Implications for Rational Drug Design</b> Caterina Barillari - <i>University of Southampton</i>
14.30 – 15.00	<b>Tea</b>
15.00 - 15.20	<b>Investigation of the charge transfer mechanism in 9-mesityl-10-methylacridinium : Rationalizing electron transfer in perpendicular systems.</b> Ben Allen - <i>University Of Newcastle Upon Tyne</i>

15.20 – 15.40	<b><math>\pi</math>-stacking in aromatic structures: a DFT and Atoms in Molecules study</b> Arturo Robertazzi - <i>Cardiff University</i>
15.40 – 16.00	<b>Modelling Aromatic Interactions via Molecular Mechanics Simulation</b> Christopher M. Baker – <i>University of Oxford</i>
16.00	<b>Compound Countdown</b>
16.30	<b>Judges Deliberations</b>
16.45	<b>Presentations</b>
17.00	<b>End</b>

## Posters

Poster 1	<b>Can Clefts Aid the Prediction of Protein-Protein Interfaces</b> Nick Burgoyne - <i>University of Leeds</i>
Poster 2	<b>A Graph Theoretic Survey of RNA Base Triple Interactions</b> Mohd Firdaus Raih - <i>University of Sheffield</i>
Poster 3	<b>Scoring Functions and Enrichment: A Case Study on Hsp 90</b> Chrysi Konstantinou Kirtay - <i>University of Cambridge</i>
Poster 4	<b>QSAR Analysis of NK3 Activity</b> Natalie Akoma-Mordi – <i>University College, London</i>
Poster 5	<b>The Application of RAPPER to and it's Implications for the Limitations in Comparative Modelling</b> Nicholas Furnham - <i>Cambridge University</i>
Poster 6	<b>QM/MM Study of the Intramolecular Proton Transfer in Glycine</b> Phaedra Williams – <i>University of Bath</i>
Poster 7	<b>Regioselectivity of Soluble Epoxide Hydrolase: A QM/MM study</b> Simon Hoyle – <i>University of Bristol</i>
Poster 8	<b>Molecular Dynamics Simulations of LDAO detergent micelles and of Mistic</b> Emi Psachoulia - <i>University of Oxford</i>
Poster 9	<b>QSPR investigation into solvation properties of platinum complexes</b> Steven Oldfield - <i>Cardiff University</i>
Poster 10	<b>Docking and direct design in the binding pocket -libraries for serine protease inhibitors</b>

Christof Gerlach - *Philipps-Universität Marburg*

Poster 11 **QM/MM modelling of reactions in citrate synthase**  
Marc van der Kamp - *University of Bristol*

Poster 12 **Molecular Modelling of DNA Compression Caused by Physical Probing Techniques**  
Verity Hudson - *University of Nottingham*