

# A Biochemical Society Training Day Hands-on Introduction to Protein Simulations



mage kindly supplied by Marcos Sotomayor (Ohio State University, USA)

#### **Organizers:**

Amir Khan Marcos Sotomayor

### **Overview:**

Protein and enzyme function is determined by both structure and dynamic flexibility in solution. Molecular simulations provide insight into conformational

transitions that play a critical role in biological activity. This training course will enable biochemists that are non-experts in computational biology to perform protein simulations and evaluate their results, using a free and widely used software package. Participants will leave with the practical knowledge of running and analysing simulations on a protein system.

## 23–24 October 2014 Charles Darwin House, London, UK

## APPLICATION DEADLINE: 25 AUGUST 2014





### **Topics:**

Sponsored by:

- \* Visualization and analysis of protein structure
- \* Theoretical aspects of molecular dynamics simulations
- \* Preparation of a protein system for molecular dynamics simulations

g innovation

- \* Practical aspects of running a simulation
- \* Output of simulations and analysis of trajectories

portlandpresslimited

\* Steered molecular dynamics

For a full programme please visit: **www.biochemistry.org** 





# A Biochemical Society Training Day Hands-on Introduction to **Protein Simulations**

23-24 October 2014 **Charles Darwin House**, London, UK

**APPLICATION DEADLINE:** 25 AUGUST 2014



Registration fees starting at £85

This event is **CPD** approved and attendees can claim 42 credits

For a full programme please visit: www.biochemistry.org

Sponsored by: portlandpresslimited publishing innovation



#### **Trainers:**

Amir Khan **Hannes Loeffler Rachelle Gaudet** Valeria Losasso

(Trinity College, Dublin, Ireland) (Science and Technology Faculties Council, UK) Marcos Sotomayor (Ohio State University, USA) (Harvard University, USA) (Science and Technology Faculties Council, UK)