

Bioinformatics and Functional Genomics Seminars, Autumn 2008-2009

Monday, 13 October 2008

- Modelling metabolism: from topology to dynamics of metabolic networks
Ralf Steuer, MIB & Humboldt University of

Monday, 20 October 2008

- Rapid proteome profiling using positional signature peptides
Jennifer Siepen, University of Manchester

Monday, 27 October 2008

- Conserved regulatory networks in vertebrate development
Greg Elgar, Queen Mary, University of London

Monday, 03 November 2008

- Studying human genetic variation by whole genome sequencing
Richard Durbin, Sanger Institute

Monday, 10 November 2008

- Novel optimization techniques for de novo protein structure prediction
Julia Handl, University of Manchester
At 12:00 PM Michael Smith Seminar Room

Monday, 17 November 2008

- Software tools for the analysis of cancer-related datasets
Crispin Miller, Paterson Institute for Cancer Research, University of Manchester

Monday, 24 November 2008

- Discrete event simulation modelling of biological control logic
David Stockton/ Riham Khalil, De Montfort University, Leicester

Monday, 01 December 2008

- An ontological support for the laboratory automation
Larisa Soldatova, University of Wales, Aberystwyth

Monday, 08 December 2008

- Better Molecular docking
Richard Sessions, University of Bristol