



GeneGo User Group Meeting Agenda 2010 - Europe

Wednesday,
May 12th

TIME	TOPIC	PRESENTER
8:30 am	Registration and Breakfast	
9 - 9:30am	GeneGo Vision and Strategy Science and Technology	Yuri Nikolsky
9:30 - 9:45 am	2009 Review	Julie Bryant
9:45 - 10:15 am	Hunting for correctors of Cystic Fibrosis	Joanna Betts GlaxoSmithKline
10:15 - 10:45 am	MetaDrug: from compounds structure to affected pathways	Eugene Myshkin
10:45 - 11 am	Break	
11 - 11:30 am	Pathway Analysis of Motor Neuronal Specification and Differentiation in developing vertebrate	Dr. David Chambers King's College London
11:30 - 12 pm	Upcoming enhancements to toxicity analysis in MetaCore and MetaDrug	Richard Brennan
12 - 1 pm	Training Group 1: Using MetaCore to answer questions such as: What are my gene lists telling me? What processes are they involved in? <ol style="list-style-type: none"> 1. What is biologically going on with my experimental data? 2. Potential target - what biological effect altered if hit upstream and downstream 3. Search for known target-around topics to find information 4. From my gene lists what is most important 5. I want to find out information about my datasets such as: <ol style="list-style-type: none"> a. Which genes are important in x disease b. Which genes are known targets c. Which genes can be easily assayed and used as biomarkers 6. What are the top pathways for my genes (having maps or network answers confuse people) 7. What are my top genes and what are they interacting with? 	Eugene Myshkin
12 - 1 pm	Training Group 2: Tox Workflows <ul style="list-style-type: none"> • What are drug risk factors? • What is the mechanism? 	Richard Brennan
1 - 2pm	Lunch	
2 - 2:30 pm	Lessons learned about ECM of fibroids using pathway analysis of gene expression data	Jörg Mueller Bayer Schering Pharma AG
2:30 - 3 pm	Integration of MetaCore in the Genomics Platform workflow	Celine Delucinge-Vivier Data Analyst University of Geneva
3 - 3:15 pm	Break	
3:15 - 3:45 pm	Working with experimental data sets	Matt Newman Vice President Business Development OmicSoft

3:45 - 4:15 pm	Identifying androgen-responsive genes in the human endometrium	Elaine Marshall <i>University of Edinburgh Centre for Reproductive Biology MRC Human Reproductive Sciences Unit</i>
4:15 - 5:30 pm	Training Group 1: OmicSoft Training Group 2: microRNA	Matt Newman Eugene Myshkin
6:00 - 8:00 pm	GeneGo get to know each other over wine, beer and appetizers. Location: De Luca Cucina & Bar 83 Regent Street Cambridge CB2 1AW, United Kingdom http://www.delucacucina.co.uk/cambridge-cocktail-bar.htm Wine, beer, appetizers and good company-come and join us! RSVP to erin@genego.com when you register for the UGM	