

EuBIC Winter School 2017



Programme



<https://www.fh-ooe.at/eubic-ws17>

Day 1 – 10. January 2017

Educational Day

Time	Title		
09:10 - 09:30	Opening Welcome & Introduction		
09:30 - 09:45	Coffee Break		
09:45 - 12:00	Developing Spectral Libraries with Progenesis QI for Proteomics, Part I	ELIXIR and de.NBI Hackathon	Introduction to Proteomics Data Analysis, Part I
12:00 - 13:00	Lunch Break		
13:00 - 14:45	Developing Spectral Libraries with Progenesis QI for Proteomics, Part II	ELIXIR and de.NBI Hackathon	Introduction to Proteomics Data Analysis, Part II
14:45 - 15:00	Coffee Break		
15:00 - 19:00	Highlights of the Latest Developments of the Proteome Discoverer Framework	ELIXIR and de.NBI Hackathon	Introduction to Proteomics Data Analysis, Part III
19:00	Dinner		

For further information on the workshops, please see the electronic version of the programme or the winter school's webpage.

Day 2 – 11. January 2017

Bioinformatics Challenges in Identification & Quantification

Time	Title	Speaker(s)
09:00 - 09:05	Morning Welcome and Announcements	
09:05 - 09:55	Constructing community knowledge for peptide identification and quantification	Nuno Bandeira
09:55 - 10:15	Coffee Break	
10:15 - 11:05	Broadening the mission: quality assessment for quantitative and label-dependent MS	David Tabb
11:05 - 12:00	Poster Flash Talks of Young Investigators Part I	
12:00 - 13:00	Lunch Break	
13:00 - 13:50	A discovery portal for bioinformatics resources	Elixir
13:50 - 14:20	Developing structural interactomics and its application in cell biology	Fan Liu
14:20 - 14:40	Coffee Break	
14:40 - 18:40	Workshop Session	see workshop descriptions
18:40 - 19:40	Dinner	
19:40	Poster Session & Come Together	

Parallel Workshops

Fan Liu	Structural Interactomics by Cross-linking Mass Spectrometry
Nuno Bandeira	Global Big Data Challenge
Qiagen	Ingenuity Pathway Analysis
David Tabb	Proteomics without a Genome: Leveraging RNA-Seq from Non-Model Organisms

Day 3 – 12. January 2017

Result Interpretation

Time	Title	Speaker(s)
09:00 - 09:05	Morning Welcome and Announcements	
09:05 - 09:55	Open Data	Lennart Martens
09:55 - 10:15	Coffee Break	
10:15 - 11:05	Issues of LC-MS Quantification	Oliver Kohlbacher
11:05 - 12:00	Poster Flash Talks of Young Investigators Part II	
12:00 - 13:00	Lunch Break	
13:00 - 13:50	Reactome: A Curated Knowledgebase of Biomolecular Pathways	Antonio Fabregat Mundo
13:50 - 14:10	Coffee Break	
14:10 - 18:10	Workshop Session	see workshop descriptions
18:10 - 19:10	Dinner	
from 19:10	Social Event	

Parallel Workshops

Jürgen Cox	Perseus & MaxQuant
Oliver Kohlbacher	Automated Processing of Quantitative Proteomics Data with OpenMS
Antonio Fabregat Mundo	Accessing Reactome Services and Integrating its Widgets

Day 4 – 13. January 2017

From Proteomics to Multi-Omics in Bioinformatics

Time	Title	Speaker(s)
09:00 - 09:05	Morning Welcome and Announcements	
09:05 - 09:55	The MaxQuant and Perseus Computational Platforms for Comprehensive Analysis of Large-scale (Prote)Omics Data	Jürgen Cox
09:55 - 10:30	Coffee Break & Poster Session	
10:30 - 11:25	Integrating Data and Software using Ontologies	Magnus Palmblad
11:25 - 12:15	Best Flash Talk and Best Poster Award	
12:15 - 14:00	Lunch, Outlook and Workshop Closing	

Sponsors

