

Friday, 17 June



Theoretical and data analysis Sessions



9-10

 Network analysis of post-synaptic proteomic data reveals molecular substructures associated with neurological disorders, <u>J Douglas</u> <u>Armstrong</u> (University of Edinburgh)

10-12.30

 Data analysis and interpretation (data from practical session),
 <u>Viviana Greco</u> (Catholic University of the "Sacred Heart") and <u>Oksana</u>
 Sorokina (University of Edinburgh)

11-11.30

Coffee break

12.30-13.30

 Functional Proteomics: proteinprotein interactions, <u>Maria Monti</u> (CEINGE and University of Naples Federico II)

Closing Remarks and lunch

Scientific Committee

ItPA Executive Board:

Paola Roncada, President

Viviana Greco, Vice President

Tiziana Alberio, Treasurer, EuPA Education Committee

Alessio Soggiu, Secretary

Maria Monti, Training Programs

Damiana Pieragostino, International relationships

Lorenza Putignani, Fund raising

Margherita Ruoppolo, CEINGE and University of Naples Federico II, CEINGE Executive Coordinator

Mauro Fasano, University of Insubria, Fondazione ItPA onlus

Local Organizing Committee

Maria Monti, School Coordinator, ItPA Training Programs

Flora Cozzolino, lab activities coordinator

Margherita Ruoppolo, CEINGE Executive Coordinator

CEINGE staff, lab tutors:

Luisa Canè, Marianna Caterino, Simona Celentano,

Michele Costanzo, Ilaria Iacobucci, Vittoria Monaco

Secretariat contact (Claudia Mignoli): 081 3737824

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Theoretical-Practical EuPA Summer School



Understanding the Proteome:
from study design
to data analysis
moving through the bench

Naples, 14-17 June 2022





Tuesday, 14 June Theoretical Sessions





Wednesday, 15 June Hands-on Sessions





Thursday, 16 June Theoretical Sessions and social event





Registration and welcome coffee
Opening Remarks, <u>Paola Roncada</u> (ItPA
President)
 Opening lecture: The potential of
quantitative proteomics, <u>Andrea Urba</u>

(Catholic University of the "Sacred Heart")
Experimental design in proteomics, Tiziana Alberio (University of Insubria)

13-14 Lunch

12-13

• Analytical methods in proteomics, <u>Flora</u>
<u>Cozzolino</u> (CEINGE and University of Naples
Federico II)

15-15.30 Coffee break

15.30-16.30 • Analytical methods in proteomics for PTMs, <u>Roman A Zubarev</u> (Karolinska Institute)

16.30-17.30 • Protein identification, <u>Alessio Soggiu</u> (University of Milan)

9-18	1) Mass mapping and PTMs identification of H4
	histone.
	Peptide mixture will be analysed by MALDI-TOF/TOF.
	2) Shotgun label free differential

proteomic analysis.

Peptide mixtures will be analysed by LTQOrbitrap XL

11.30 Coffee Break

13 Lunch

16.30 Coffee Break

9.30-10.30	 Methodological approaches in 	
	Quantitative Proteomics, <u>Peter</u>	
	Horvatovich (University of Groningen)

10.30-11 Coffee Break

• Applicative tools for quantitative proteomics, <u>Marianna Caterino</u> (CEINGE and University of Naples Federico II)

• Feature selection in quantitative proteomics, <u>Mauro Fasano</u> (University of Insubria)

13-14 Lunch

15.30 Cultural event: visit to Campi Flegrei with a guide and social dinner