

#### Friday, 17 June



#### Theoretical and data analysis Sessions



9-10

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

10-12.30

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

11-11.30

Coffee break

12.30-13.30

 Functional Proteomics: proteinprotein interactions, Maria Monti (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

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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





| 10    | Registration and welcome coffee             |  |
|-------|---|--|
| 10.30 | Opening Remarks, <u>Paola Roncada</u> (ItPA |  |
|       | President)                                  |  |

 Opening lecture: The potential of quantitative proteomics, <u>Andrea Urbani</u> (Catholic University of the "Sacred Heart")

• Experimental design in proteomics, <u>Tiziana Alberio</u> (University of Insubria)

**13-14** Lunch

11-12

• 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 Sogqiu</u> (University of Milan)

9-18
1) Mass mapping and PTMs identification of H4
histone.
Peptide mixture will be analysed by MALDI-

TOF/TOF.

 Shotgun label free differential proteomic analysis.
 Peptide mixtures will be analysed by LTQ-Orbitrap XL

**11.30** Coffee Break

13 Lunch

**16.30** Coffee Break

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

| <b>10.30-11</b> C | offee Break |
|-------------------|-------------|
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• 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