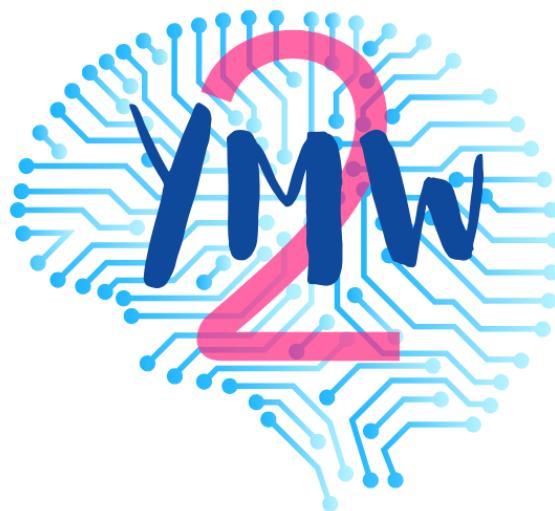


YOUNG MINDS AT WORK: BLENDING BIOCHEMISTRY AND BIOINFORMATICS

Program on-line workshop

10th -12th December 2024



December 10th, 2024

8:45-9:00	Join the workshop	
9:00-9:10	Welcome Prof. Anna Marabotti	
9:10-13:00	ELIXIR-IT training Prof. Luana Licata	Biological Network: topological analysis and visualization using Cytoscape software
13:00-14:30	Break	
14:30-14:45	Welcome Prof. Raffaele Calogero (Presidente BITS) Prof. Pier Luigi Martelli (Membro del direttivo SIB) Dr. Anna Villa (Direttore ITB-CNR)	

Opening Keynote lecture Chair: Mirko Treccani

14:45-15:15	Prof. Yinglong Miao	Accelerated Molecular Simulations and Drug Discovery
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Session 1 Chairs: Simone Pernice, Giacomo Baruzzo, Mirko Treccani

15:15-15:30	Ferrini Alessio	<i>Angiogenesis and vascular patterning: a calibrated mathematical model using the chicken chorioallantoic membrane</i>
15:30-15:45	Defilippo Annamaria	<i>Understanding Complex Systems through Differential Causal Networks</i>
15:45-16:00	Pietrafesa Davide	<i>Computational Benchmarking of Aptamers for Selective Inhibition of HuR</i>
16:00-16:15	Ghafouri Hamidreza	<i>Intrinsically Disordered Ensemble Tools (IDPET), A Python Package for Studying Conformational Ensembles of Disordered Proteins</i>

16:15-16:45 **Break**

16:45-17:00	Ansermino Chiara	<i>EPS-urine derived extracellular vesicles: multi-omics profiling for prostate cancer risk stratification</i>
17:00-17:15	Franchina Davide G.	<i>Integration of mass spectrometry and mass cytometry imaging for spatially resolved single cell profiling</i>
17:15-17:30	Lapi Francesco	<i>Multi-omics data analysis to characterize age-specific metabolic shifts during embryo development</i>

December 11th, 2024

9:00-9.15 **Join the workshop**

Opening Keynote Lecture Chair: Maria Cristina Aspromonte

9:15-9:45 **Prof. Sandra Macedo-Ribeiro** Pathogen-specific structural features in *C. albicans* Ras1 activation complex

Session 2 Chairs: Alfonso Trezza, Giulia Babbi, Lara Console

9:45-10:00 **Lopatriello Giulia** Molecular determinants of response to neoadjuvant chemotherapy in esophageal adenocarcinoma at single-cell spatial resolution

10:00-10:15 **Vono Niccolò** Exploring the molecular mechanisms that influence Retinoic Acid responsiveness or resistance in cancer stem cells within patient-derived cancer organoids.

10:15-10:45 **Break**

10:45-11:00 **Bellia Fabio** Exploring synthetic *SIL1* protein reintroduction to treat Marinesco-Sjogren Syndrome.

11:00-11:15 **Sgambati Domenico** Two new members of Ros/MucR family: *MucR2* and *MucR3* from *Sinorhizobium meliloti*

11:15-11:30 **Tisi Annamaria** Early biochemical alterations of the endocannabinoid system in the retina of Alzheimer's disease mice

11:30-11:45 **Barone Francesca** *OCTN1* (*SLC22A4*) displays two different transport pathways for organic cations or zwitterions

11:45-12:00 **Sabbioni Giuseppe** *Shwachman-Diamond Syndrome*: different signalling pathways driving chronic inflammation.

12:00-13:00 **Poster session**

13:00-14.00 **Break**

Session 3 Chairs: Giulia Cesaro, Laura Veschetti, Beatrice Casini

14:00-14:15 **Manfredi Matteo** *Alpha&ESMhFolds*: a web server for comparing 42,942 *AlphaFold2* and *ESMFold* models of the human reference proteome

14:15-14:30 **Semente Miriam Rainha** Bioinformatic neoantigens prediction from Next generation sequencing data

14:30-14:45	Baldan Matteo	<i>Simulation of bacteria interaction networks: from topology to species abundances</i>
14:45-15:00	Polito Maria Pia	<i>Biochemical characterization of the feedforward loop between CDK1 and FOXM1 in epidermal stem cells</i>
15:00-15:30 Break		
15:30-15:45	Giallongo Sebastiano	<i>Glutamate enrichment in myelofibrosis tumor microenvironment induces mesenchymal stromal cells senescence by promoting intracellular fumarate accumulation.</i>
15:45-16:00	Tussardi Gaia	<i>Addressing the Biological Interpretation of High-Dimensional Cell-Cell Communication Data from Single Cell Transcriptomics Through an Interactive Web Application</i>
16:00-16:15	Castro Naser Ximena Aixa	<i>Role of Integrins in Maternal-Embryonic Communication: Proteomic Analysis of Early Pregnancy in Alpacas (<i>Vicugna pacos</i>)</i>
16:15-16:30	Zarifa Osmanli	<i>Large-scale analysis of structured tandem repeat proteins across the tree of life</i>
Special Topic Chairs: Pasqualina D'Ursi, Paola Fossa		
16:30-17:00	Fabio Gasparini	<i>A Tool is a tool is a tool. The alignment of computational methods and clinical and laboratory practices in biomedical experimental arrangements</i>

December 12th, 2024

9:00-9.15 **Join the workshop**

Opening Keynote Lecture Chair: Lara Console

9:15-9:45 **Prof. Massimo Aureli** *Neuronal Differentiation and Degeneration: Role of the Plasma Membrane*

Session 4 Chairs: Giulia Babbi, Beatrice Casini, Maria Cristina Aspromonte

9:45-10:00 **Scarano Naomi** *Computational Strategies for the design of novel TAAR1 agonists*

10:00-10:15 **Mazza Tiziano** *3D structure of the ABC transporter MRP2 by Cryo-EM. Insights Post-translational modifications*

10:15-10:45 **Break**

10:45-11:00 **Buoli Comani Valeria** *Cryo-EM structures of hemoglobin-*lsdH* complexes: paving the way for novel protein-protein interaction inhibitors as antibiotics*

11:00-11:15 **Stelitano Giovanni** *Repurposing of approved drugs to fight *M. abscessus* infections.*

11:15-11:30 **Anteghini Marco** *X-MAP: Explainable Mutation Analysis Platform for Predicting Genetic Variant Impacts on Protein Structure, Stability, and Human Health*

11:30-11:45 **Buttarro Veronica** *Identifying disease biomarkers using a novel data analysis pipeline based on multi-view learning*

11:45-12:00 **Vestuto Vincenzo** *Biochemical Characterization of a New Class of Selective Haspin kinase Inhibitors as potential anticancer drugs*

12:00-12:15 **Conclusions**

12:15-12:30 **Best Oral and Best Poster awards**