Decemb	er 10th	. 2024
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8:45-9:00	Join the workshop	
9:00-9:10	Welcome Prof. Anna Marabotti	
9:10-13:00		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)	

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
Session 1	Chairs: Simone Pernice, Giacomo Baruzzo, Mirko Treccani	
15:15-15:30	Ferrini Alessio	Angiogenesis and vascular patterning: a calibrated mathematical model using the chicken chorioallantoic membrane
15:30-15:45	Defilippo Annamaria	Understanding Complex Systems through Differential Causal Networks
15:45-16:00	Pietrafesa Davide	Computational Benchmarking of Aptamers for Selective Inhibition of HuR
16:00-16:15	Ghafouri Hamidreza	Intrinsically Disorder Ensemble Tools (IDPET), A Python Package for Studying Conformational Ensembles of Disordered Proteins
16:15-16:45	Break	
16:45-17:00	Ansermino Chiara	EPS-urine derived extracellular vesicles: multi-omics profiling for prostate cancer risk stratification
17:00-17:15	Franchina Davide G.	Integration of mass spectrometry and mass cytometry imaging for spatially resolved single cell profiling
17:15-17:30	Lapi Francesco	Multi-omics data analysis to characterize

development

age-specific metabolic shifts during embryo

December 11th, 2024

12:00-13:00 Poster session

Break

13:00-14.00

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	a, 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.

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	Simulation of bacteria interaction networks: from topology to species abundances
14:45-15:00	Polito Maria Pia	Biochemical characterization of the feedforward loop between CDK1 and FOXM1 in epidermal stem cells
15:00-15:30	Break	
15:30-15:45	Giallongo Sebastiano	Glutamate enrichment in myelofibrosis tumor microenvironment induces mesenchymal stromal cells senescence by promoting intracellular fumarate accumulation.
15:45-16:00	Tussardi Gaia	Addressing the Biological Interpretation of High-Dimensional Cell-Cell Communication Data from Single Cell Transcriptomics Through an Interactive Web Application
16:00-16:15	Castro Naser Ximena Aixa	Role of Integrins in Maternal-Embryonic Communication: Proteomic Analysis of Early Pregnancy in Alpacas (Vicugna pacos)
16:15-16:30	Zarifa Osmanli	Large-scale analysis of structured tandem repeat proteins across the tree of life

Special Topic

Chairs: Pasqualina D'Ursi, Paola Fossa

16:30-17:00	Fabio Gasparini	A Tool is a tool is a tool. The alignment of
		computational methods and clinical and laboratory practices in biomedical experimental
		arrangements

December 12th, 2024

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

Opening Keyn	ote Lecture
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Chair: Lara Console

12:15-12:30

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-IsdH 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	Title: X-MAP: Explainable Mutation Analysis Platform for Predicting Genetic Variant Impacts on Protein Structure, Stability, and Human Health
11:30-11:45	Buttaro 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	

Best Oral and Best Poster awards