

BRIDGING THE GAP BETWEEN EXPERIMENTALISTS, BIOCURATORS AND ANALYSIS TOOL PROVIDERS

UCL-Rome meeting
funded by UCL Cities Partnerships Programme

ROME / MAY 2nd, 2023

Aula Seminari "Francesco Autuori", Department of Biology, Università degli Studi di Roma "Tor Vergata"

SCIENTIFIC COMMITTEE

Ruth Lovering - Functional Gene Annotation group, University College of London, UCL, UK.
Luana Licata - MINT database coordinator, University of Rome tor Vergata, Italy.

PROGRAMME

09:00-09:10	Welcome Antonella Canini, Director of the Dept. of Biology, Univ. "Tor Vergata" Luana Licata and Ruth Lovering, Scientific Organisers	12:15-12:45	Open discussion
09:10-09:30	Gene Ontology annotation of microRNAs Ruth Lovering, University College London, UCL	12:45-13:45	Lunch
09:30-09:55	PPI networks and system biology approaches for the study of complex neurodegenerative disorders Claudia Manzoni, University College London, UCL	13:45-14:00	Structural Bioinformatics: Exploitation of Proteins as Drugs, Drug Precursors, Drug Targets and Drug Carriers Veronica Morea, IBPM, CNR
09:55-10:20	Evolutionary and functional links between neuropilins and VEGF Snezana Djordjevic, University College London, UCL	14:00-14:15	Uncovering new signaling pathways modulated by Src kinase in cancer Daniela Barilà, University of Rome Tor Vergata
10:20-10:45	Seeing pathogenic "signals" in the microbiome "noise" – dynamic bacterial expression patterns in urinary tract infection Jennifer Rohn, University College London, UCL	14:15-14:30	A network-based approach to address complex diseases Francesca Sacco University of Rome Tor Vergata
10:45-11:15	Coffee/tea	14:30-14:45	SignalingProfiler: leveraging manually-curated data toward multi-omics integration Livia Perfetto, University of Rome Sapienza
11:15-11:30	Contextualizing molecular networks for human diseases Luana Licata, University of Rome Tor Vergata	14:45-15:00	Identification of conserved modules embedded into non-homologous lncRNA genes in human Manuela Helmer-Citterich, University of Rome Tor Vergata
11:30-11:45	Transcriptomics data analysis and functional interpretation of gene lists Teresa Colombo, IBPM, CNR	15:00-15:15	Deciphering Adipose Tissue Cell Heterogeneity: Development of scRNAseq-based Bioinformatics Tools Andrea Ninni and Daniele Lettieri Barbato, University of Rome Tor Vergata
11:45-12:00	Omics data integration for personalized medicine: challenges and opportunities Allegra Via, University of Rome Sapienza	15:15-15:45	Coffee/tea
12:00-12:15	The clinical potential of peripheral human tissues analysis in Parkinson's disease Tommaso Schirinzi, University of Rome Tor Vergata	15:45-17:00	Open discussion