16:15–17:45
BIOINF Main Track 5

Aula Magna

16:15–16:30	**PRIMA: a bidirectional state-space architecture and training approach for modelling protein-protein interactions** Arturo Fiorellini-Bernardis · Sebastien Boyer · Jean Quentin · Ghassene Jebali · Oliver Bent
16:30–16:45	**iPerturb: An interpretable framework for predicting gene expression changes under perturbations** Khang Ta Gia · Linh Huynh Viet
16:45–17:00	**Katz Kernel Based Stratification of Breast Cancer Patients** Francesco Cellitti · Lorenzo Di Rocco · Livia Perfetto · Veronica Lombardi · Umberto Ferraro Petrillo
17:00–17:15	**PANDORA: a multiparametric framework for sensitive detection of insertional mutagenesis** Riccardo Pizzichemi
17:15–17:30	**Towards a unified framework: a data structures benchmark for representing genomic information** Matteo Tommaso Ungaro · Mauricio Orantes Bonilla · Francesca Querques · Evelyne Tassone · Simona Giunta
17:30–17:45	**An XAI-Driven Transformer for Non-Canonical Splice-Site Prediction** Ahmed Negm