

**Finding regulatory elements in eucaryotes: a statistical approach using Gene Ontology - (session: Other)**

Davide Corà, Paolo Provero, Michele Caselle

Universita' degli Studi di Torino

The discovery of regulatory elements in eucaryotes is one of the most important challenges of bioinformatics. In this poster we present a method in which sets of genes, generated from statistical determination of overrepresented motifs, are correlated with the prevalence of Gene Ontology annotation terms.

POSTER ONLY