

Risultati

Computational analysis of biological structures and networks

Workshop on “Dissimilarity-based approaches in bioinformatics”

Results September 2017

A support vector machine based test for incongruence between sets of trees in tree space	15/15
Metabolic pathways variability and sequence/networks comparisons	14.5/15
The Repertoire Dissimilarity Index as a method to compare lymphocyte receptor repertoires	12/15