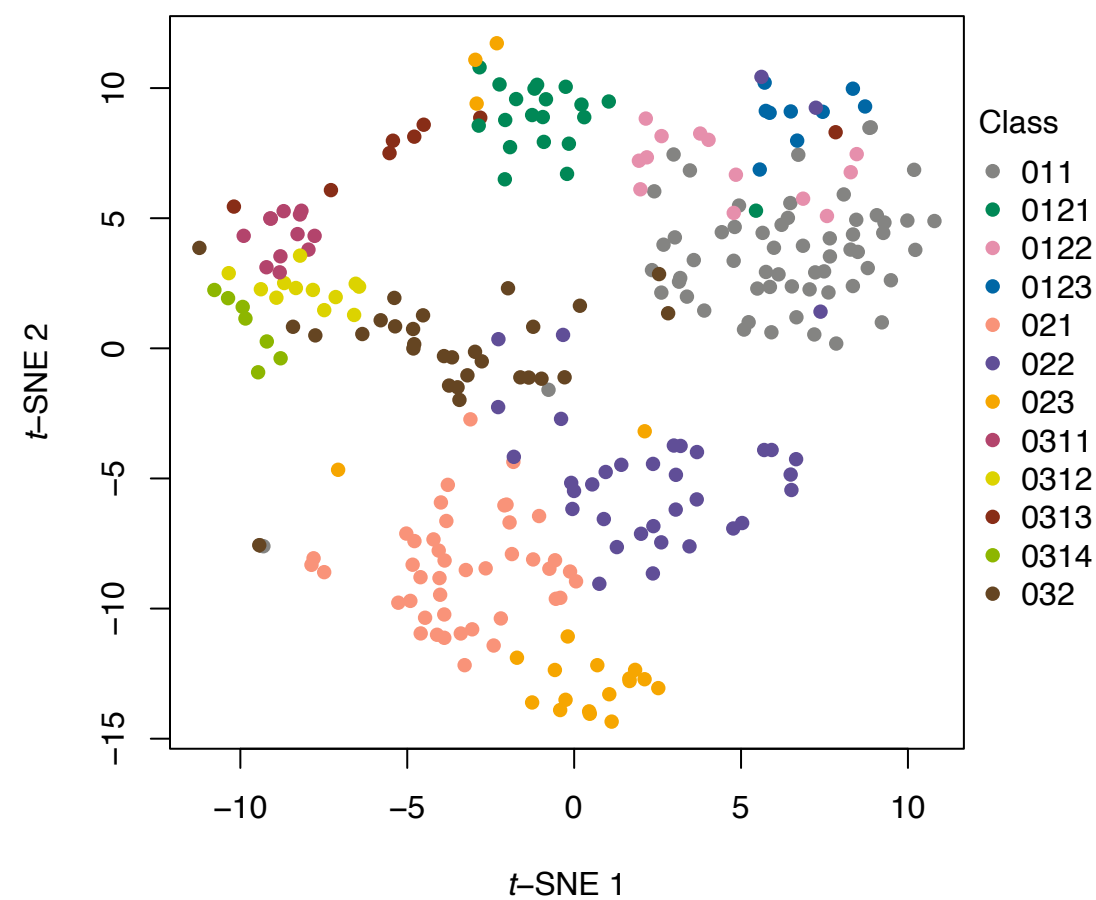
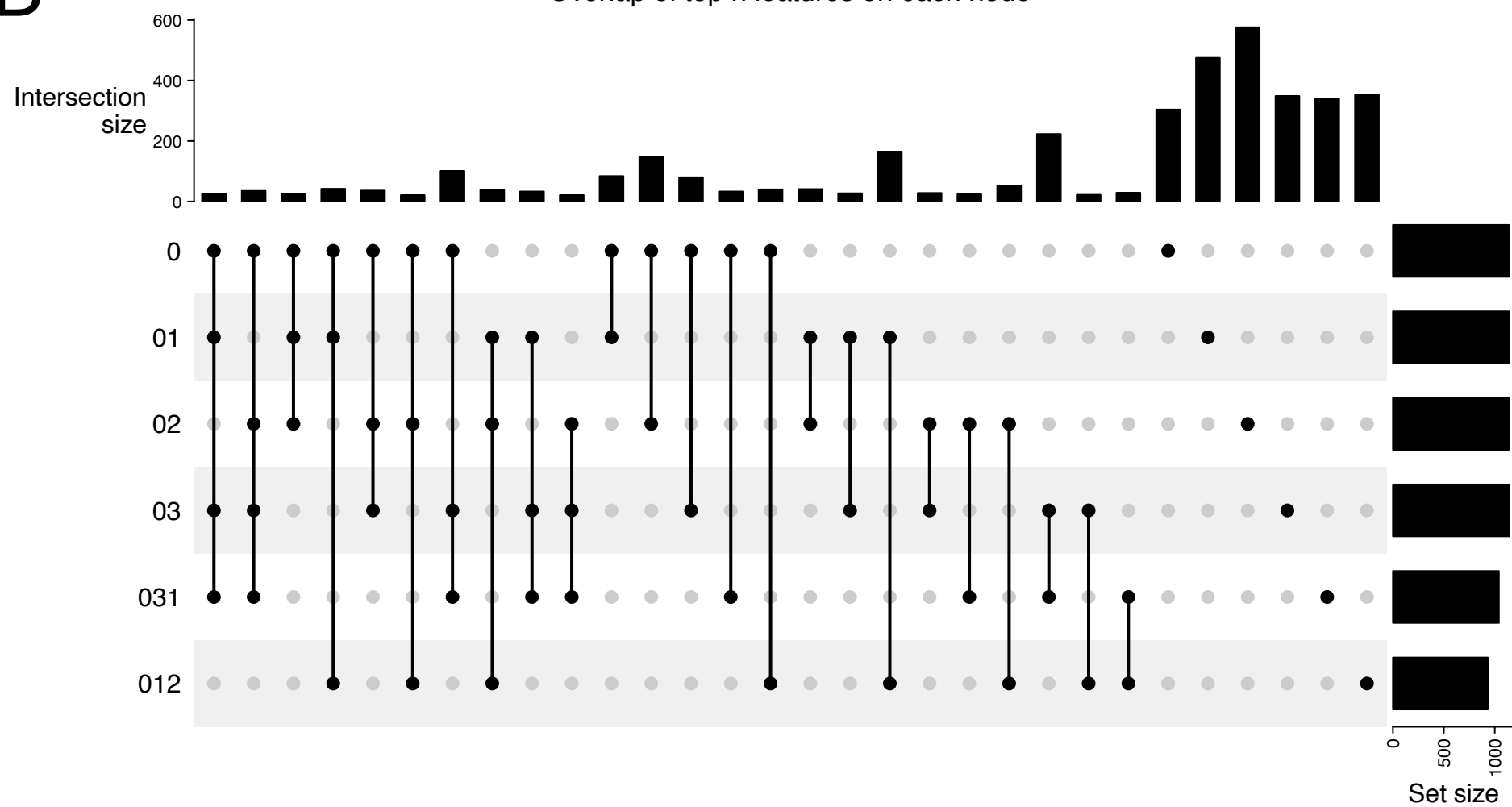


A

t-SNE on 1000 rows with the highest ATC scores

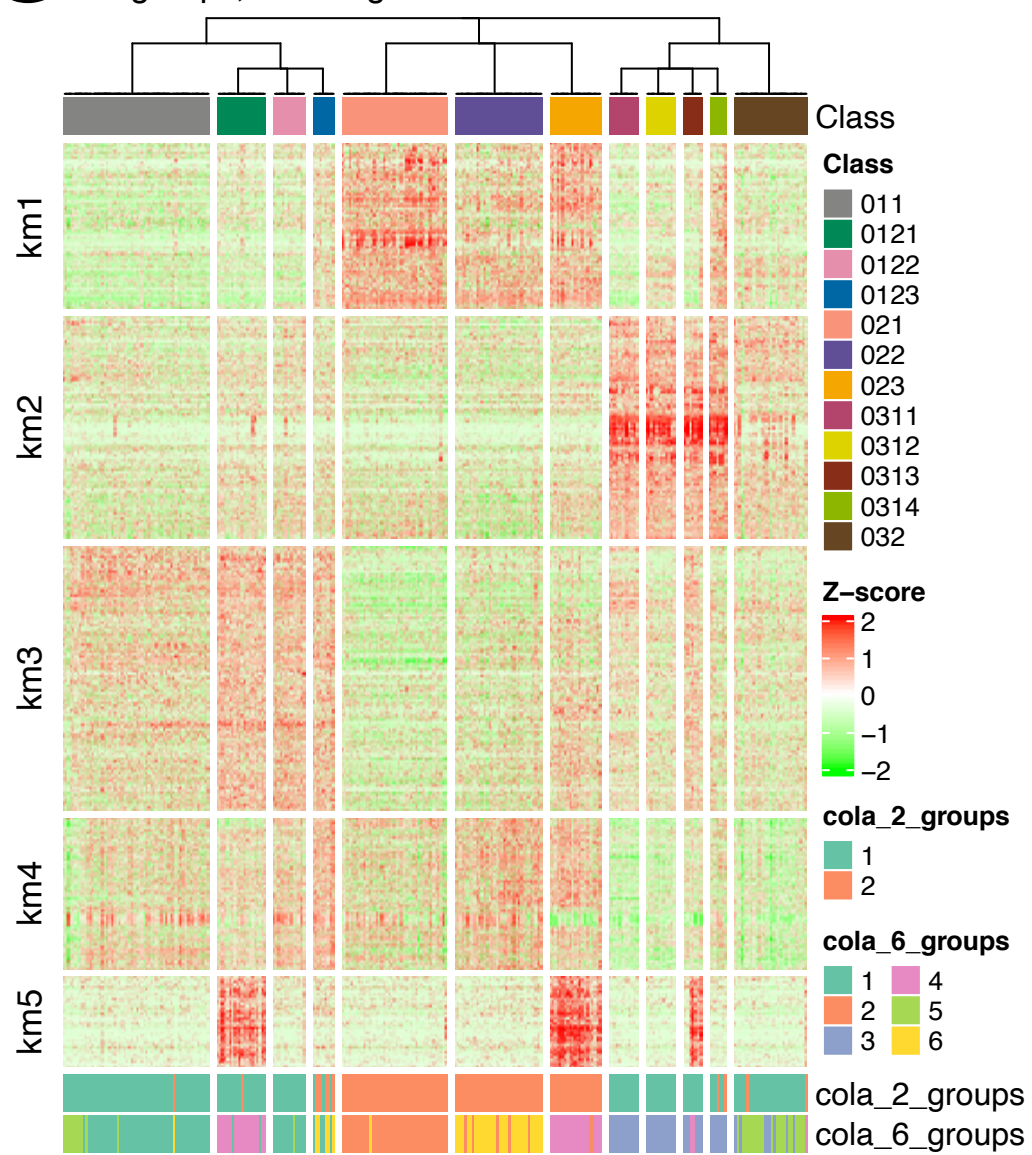


B

Overlap of top n features on each node

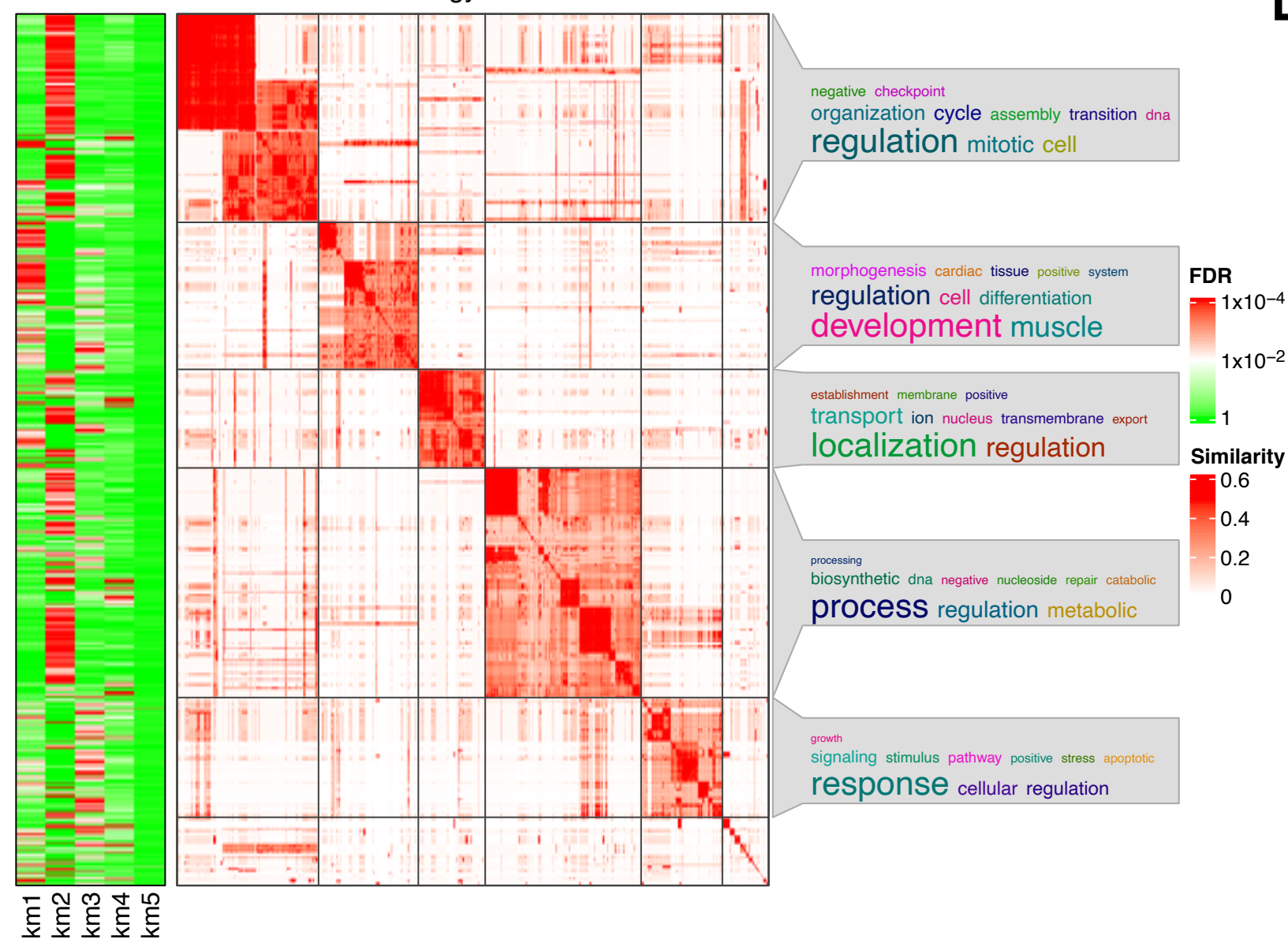
C

12 groups, 4683 signatures under FDR < 0.05



D

1150 Gene Ontology terms



E

Stability of classification from 20 runs

