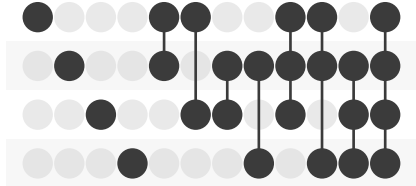
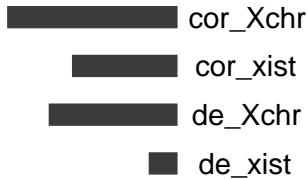
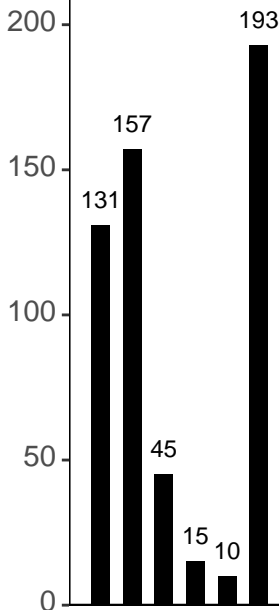


Number of significant genes at d1
across the four analyses



300 200 100 0
Set Size