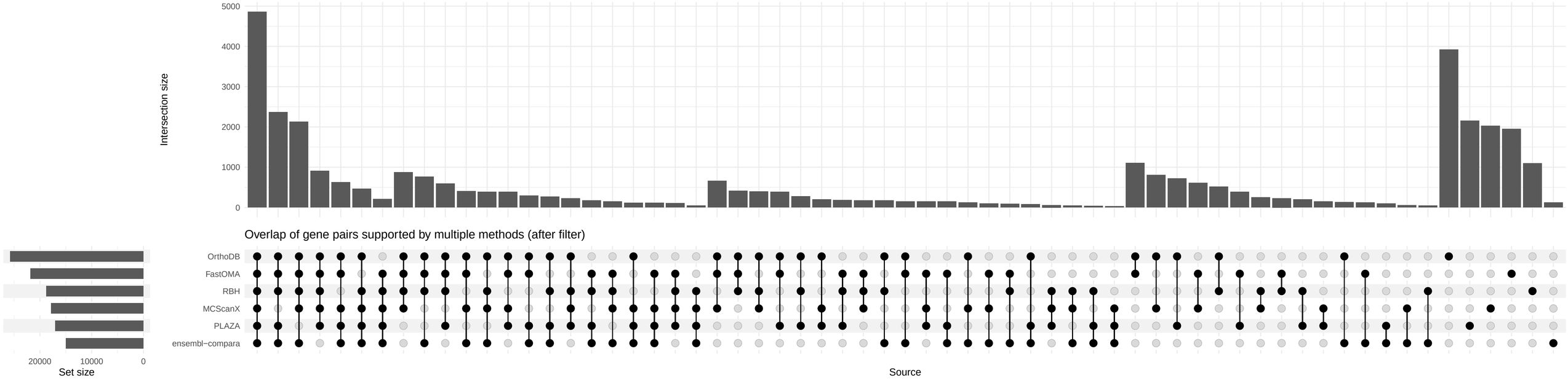
Intersection size
5000
4000
3000
2000
1000
0
Overlap of gene pairs supported by multiple methods (after filter)
OrthoDB
FastOMA
RBH
MCScanX
PLAZA
ensembl−compara
20000
10000
0
Set size
Source