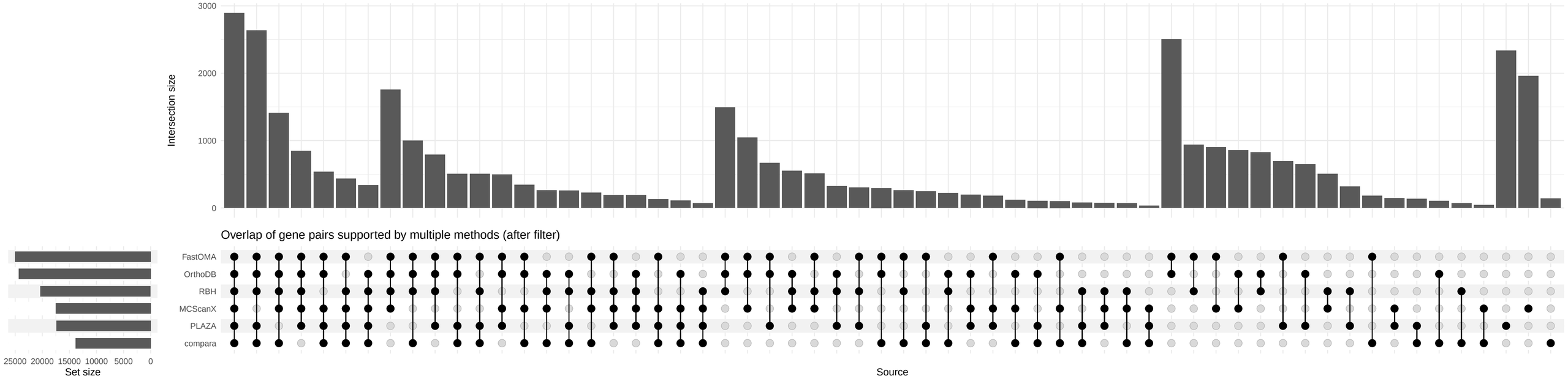
Intersection size
3000
2000
1000
0
Overlap of gene pairs supported by multiple methods (after filter)
FastOMA
OrthoDB
RBH
MCScanX
PLAZA
compara
25000 20000 15000 10000 5000 0
Set size
Source