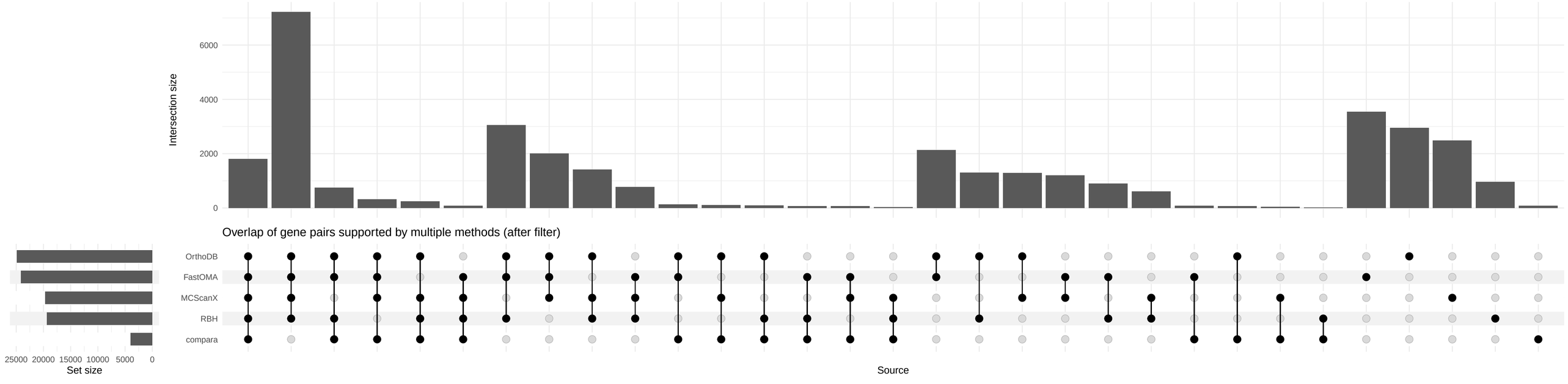
Intersection size
6000
4000
2000
0
Overlap of gene pairs supported by multiple methods (after filter)
OrthoDB
FastOMA
MCScanX
RBH
compara
25000 20000 15000 10000 5000 0
Set size
Source