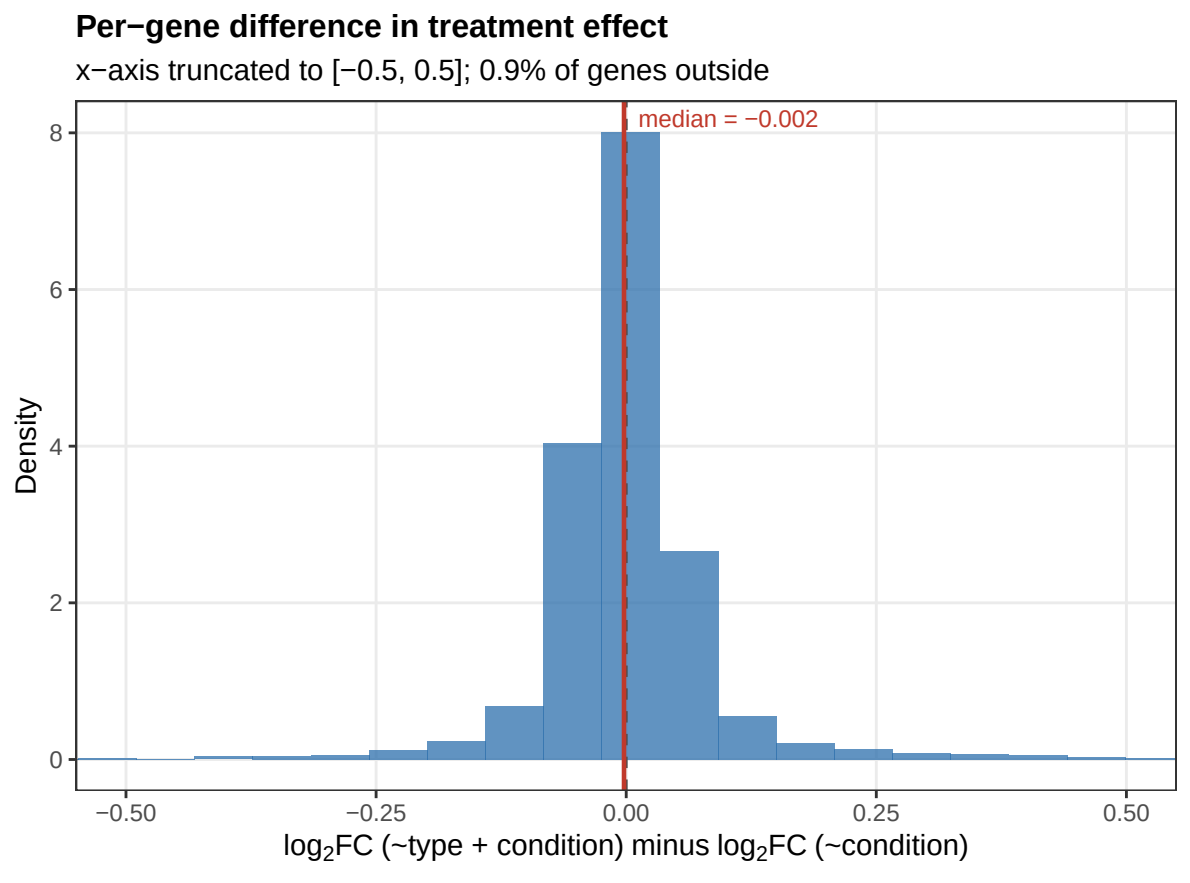
Per−gene difference in treatment effect
x−axis truncated to [−0.5, 0.5]; 0.9% of genes outside
median = −0.002
Density
8
6
4
2
0
−0.50
−0.25
0.00
0.25
0.50
log2FC (~type + condition) minus log2FC (~condition)