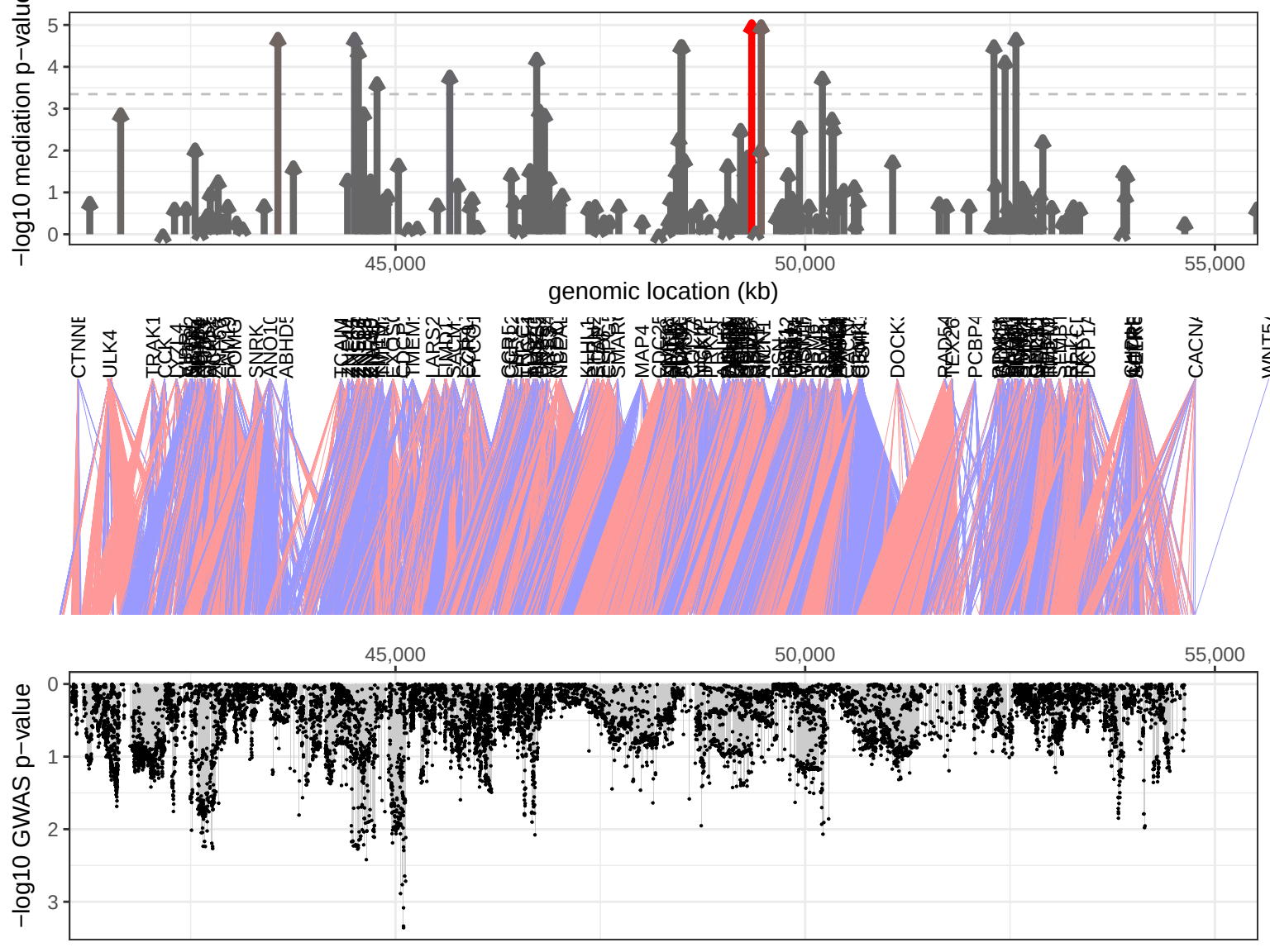
−log10 mediation p−value
0
1
2
3
4
5
45,000
50,000
55,000
genomic location (kb)
CTNNE
ULK4
TRAK1
SNRK
ANO10
ABHD
LARS2
MAP4
DOCK
RAD54
PCBP4
CACNA
45,000
50,000
55,000
−log10 GWAS p−value
0
1
2
3