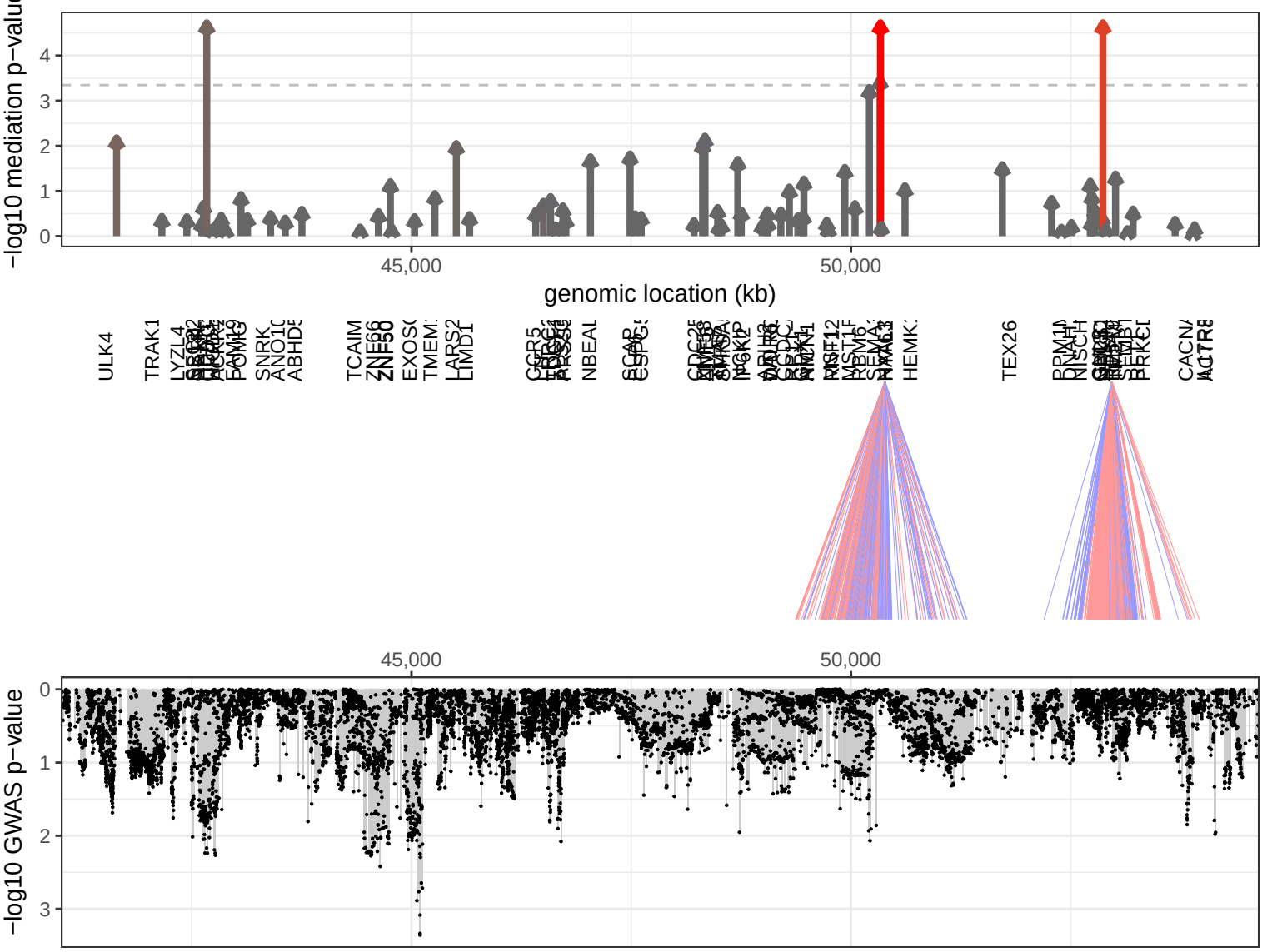
−log10 mediation p−value
0
1
2
3
4
45,000
50,000
genomic location (kb)
ULK4
TRAK1
SNRK
ANO10
ABHD
TCAIM
ZNF66
EXOSC
TMEM
LARS2
LIMD1
NBEAL
HEMK
TEX26
CACN
−log10 GWAS p−value
0
1
2
3