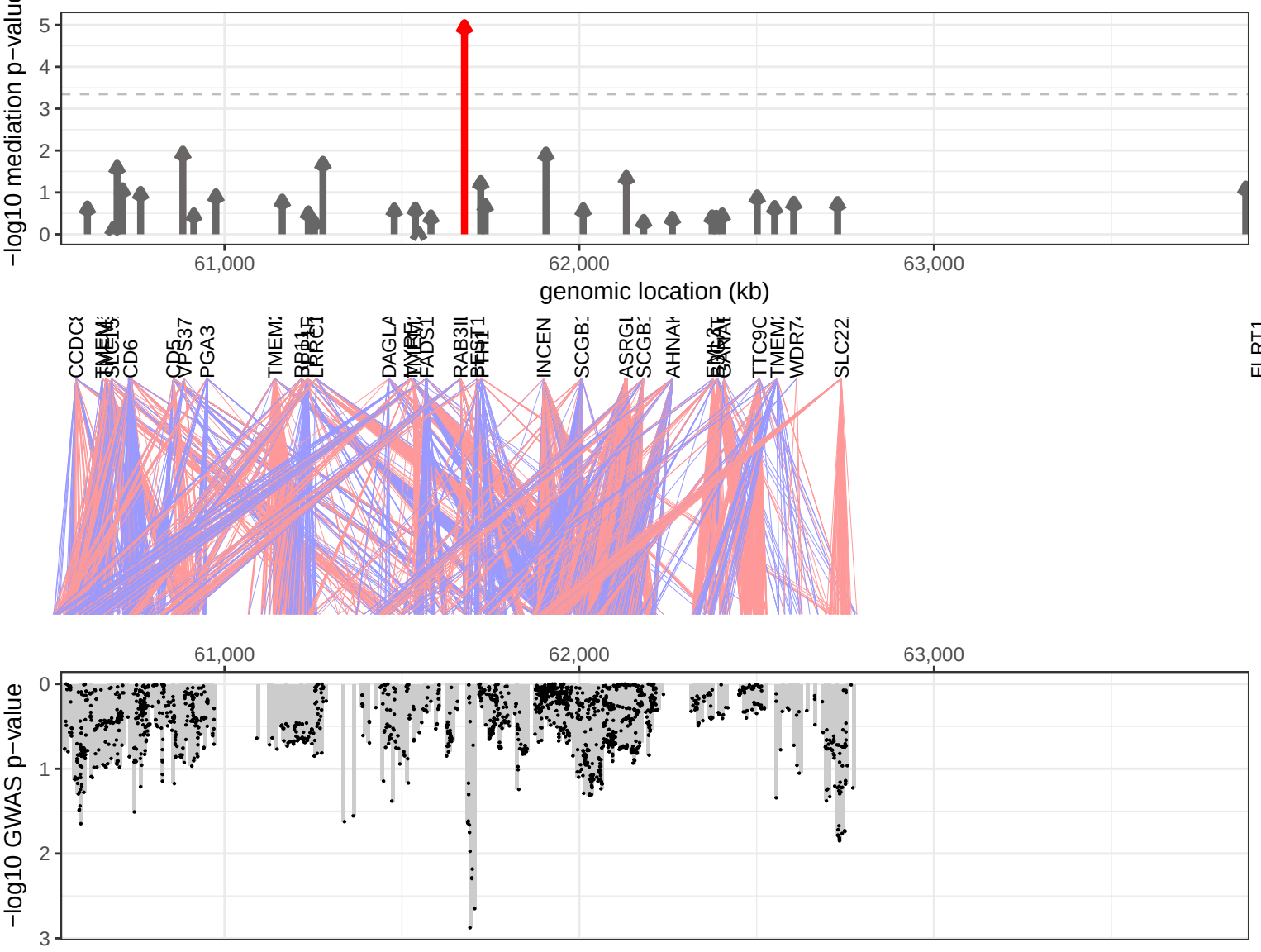
−log10 mediation p−value
5
4
3
2
1
0
61,000
62,000
63,000
genomic location (kb)
CCDC
CD6
CD5
PGA3
TMEM
LRRC1
DAGLA
FADS1
RAB3IL
INCEN
SCGB
ASRGL
AHNAK
TTC9C
TMEM
WDR74
SLC22
61,000
62,000
63,000
−log10 GWAS p−value
0
1
2
3