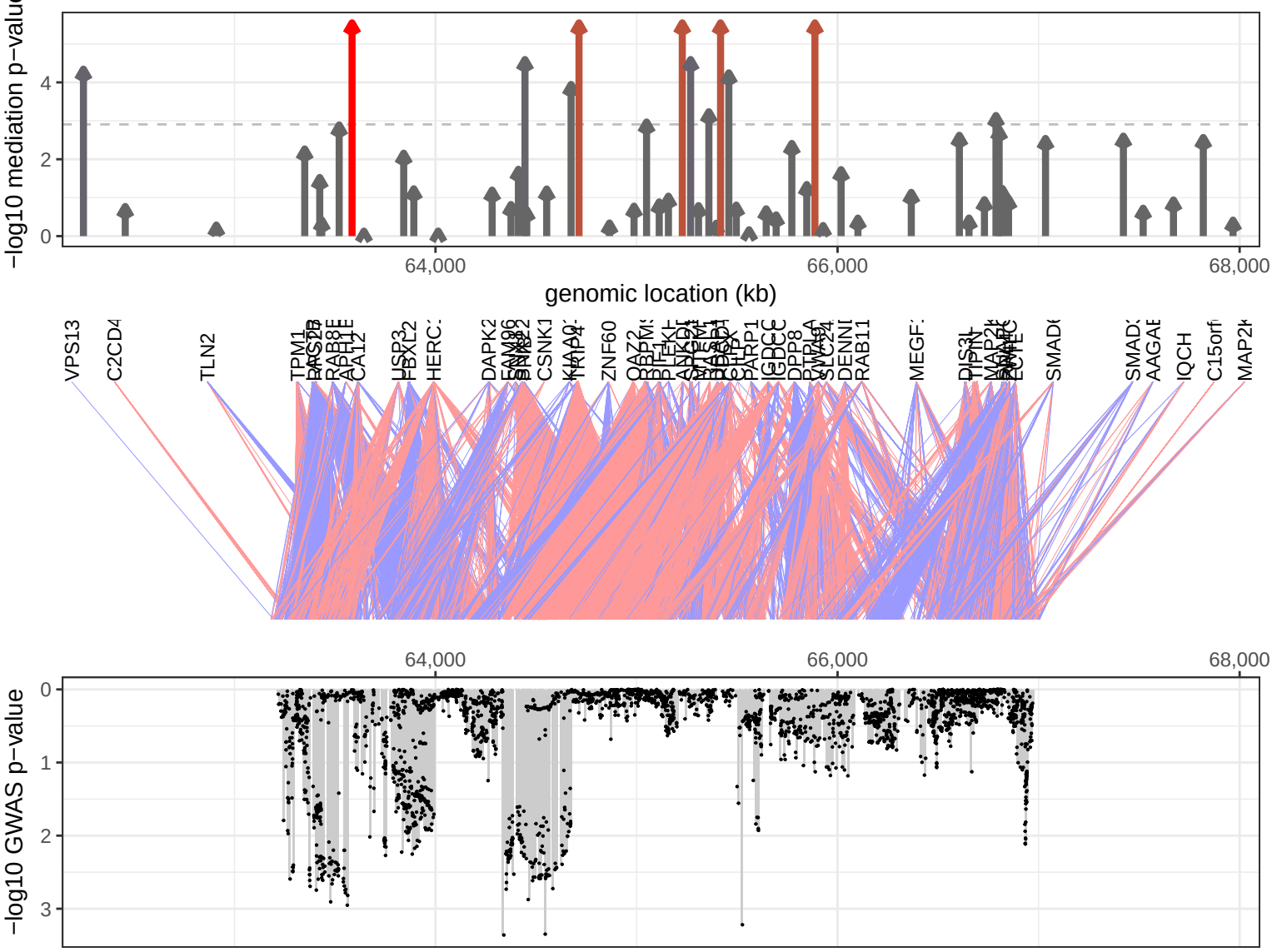
−log10 mediation p−value
0
2
4
64,000
66,000
68,000
genomic location (kb)
VPS13
C2CD4
TLN2
TPM1
CA12
USP3
HERC
DAPK2
CSNK1
ZNF60
PARP1
DPP8
RAB11
MEGF
SMAD
AAGAB
IQCH
C15orf
MAP2K
64,000
66,000
68,000
−log10 GWAS p−value
0
1
2
3