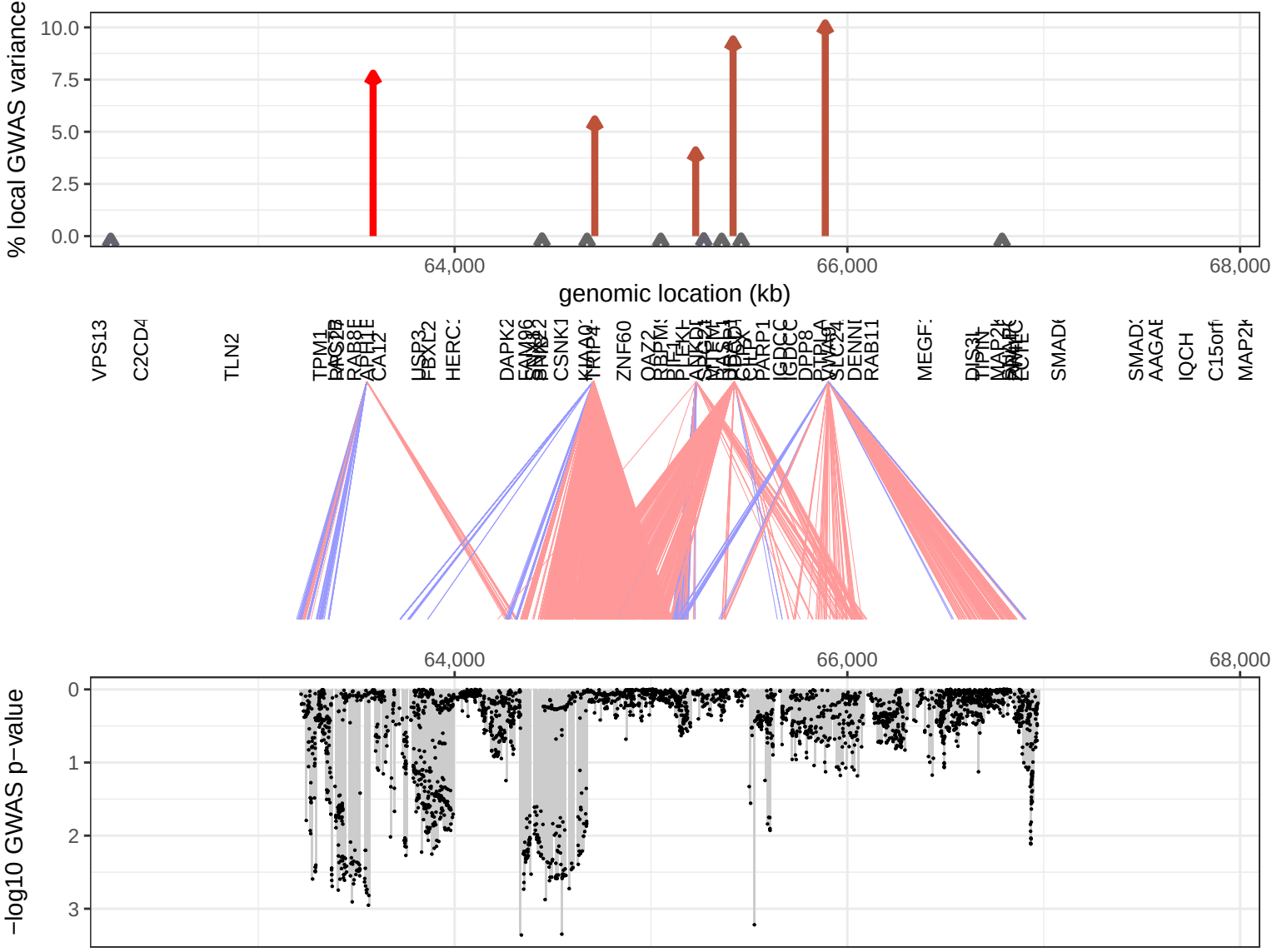
% local GWAS variance
10.0
7.5
5.0
2.5
0.0
64,000
66,000
68,000
genomic location (kb)
VPS13
C2CD4
TLN2
CA12
HERC
DAPK2
CSNK1
ZNF60
PARP1
DPP8
DENN
RAB11
MEGF
SMAD
SMAD
AAGAB
IQCH
C15orf
MAP2K
−log10 GWAS p−value
0
1
2
3