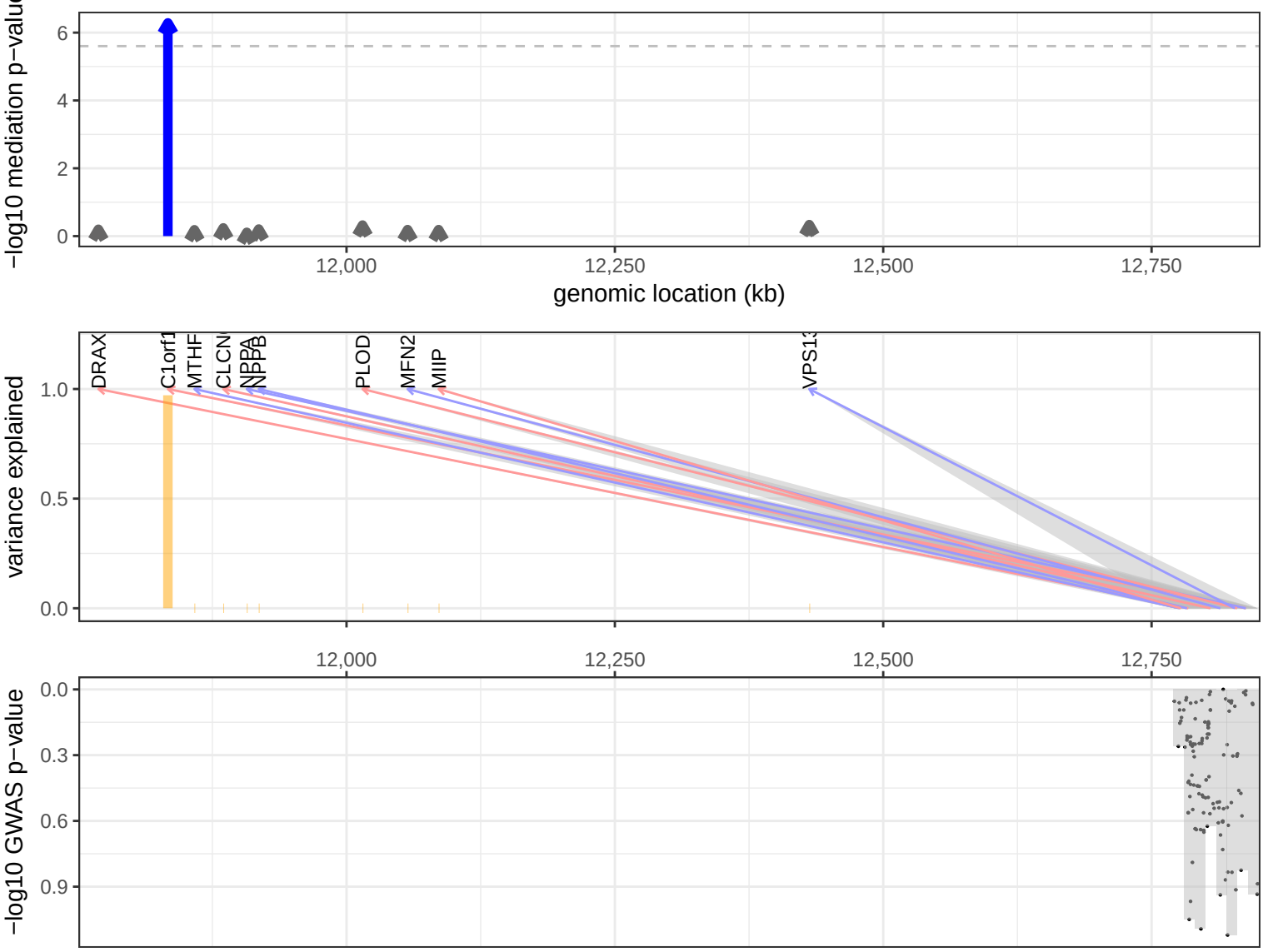
−log10 mediation p−value
6
4
2
0
12,000
12,250
12,500
12,750
genomic location (kb)
variance explained
1.0
0.5
0.0
DRAX
C1orf1
MTHF
CLCN
NPPA
NPPB
PLOD
MFN2
MIIP
VPS13
12,000
12,250
12,500
12,750
−log10 GWAS p−value
0.0
0.3
0.6
0.9