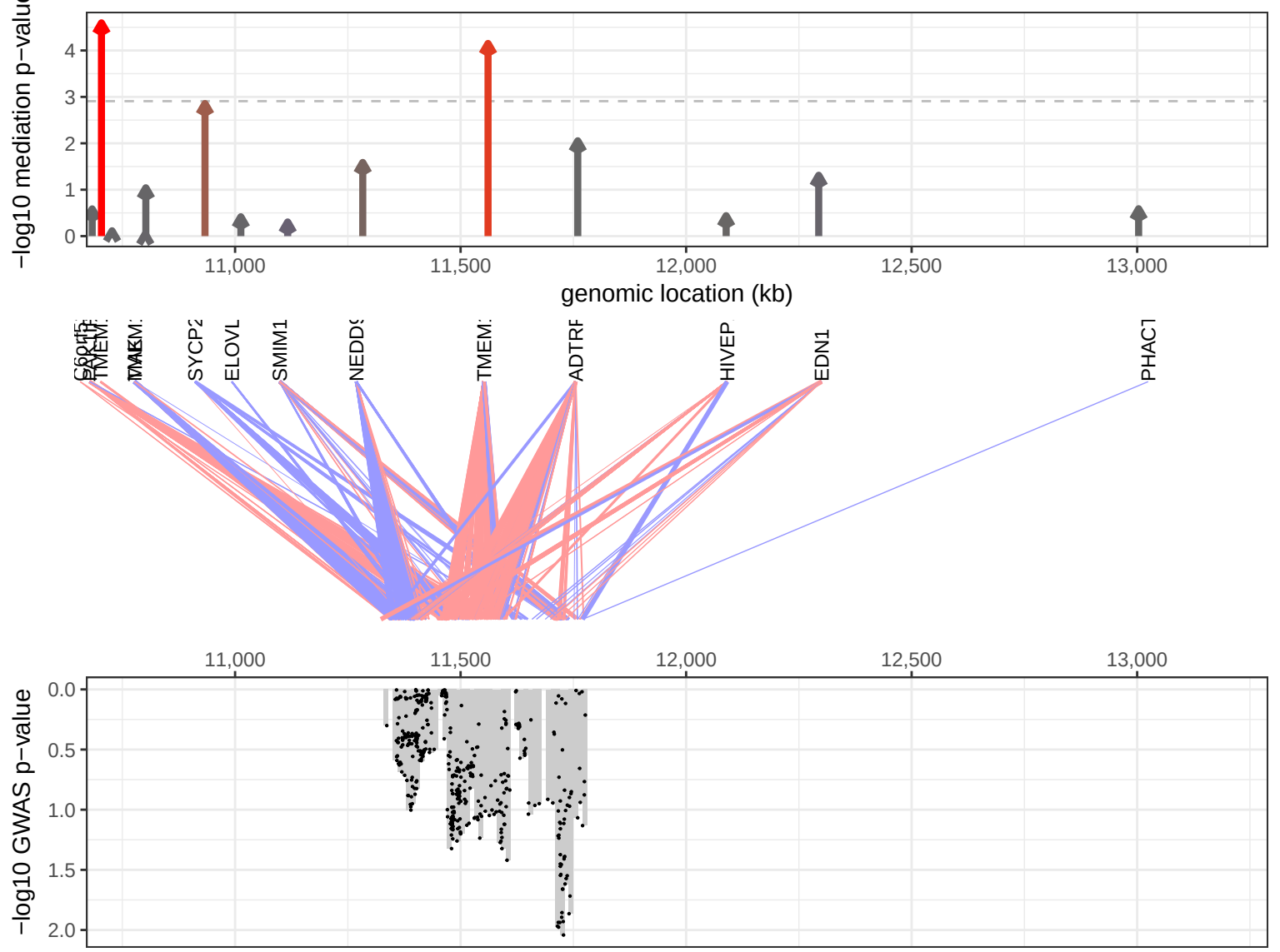
−log10 mediation p−value
4
3
2
1
0
11,000
11,500
12,000
12,500
13,000
genomic location (kb)
SYCP2
ELOVL
SMIM1
NEDD
TMEM
ADTRF
HIVEP
EDN1
PHACT
11,000
11,500
12,000
12,500
13,000
−log10 GWAS p−value
0.0
0.5
1.0
1.5
2.0