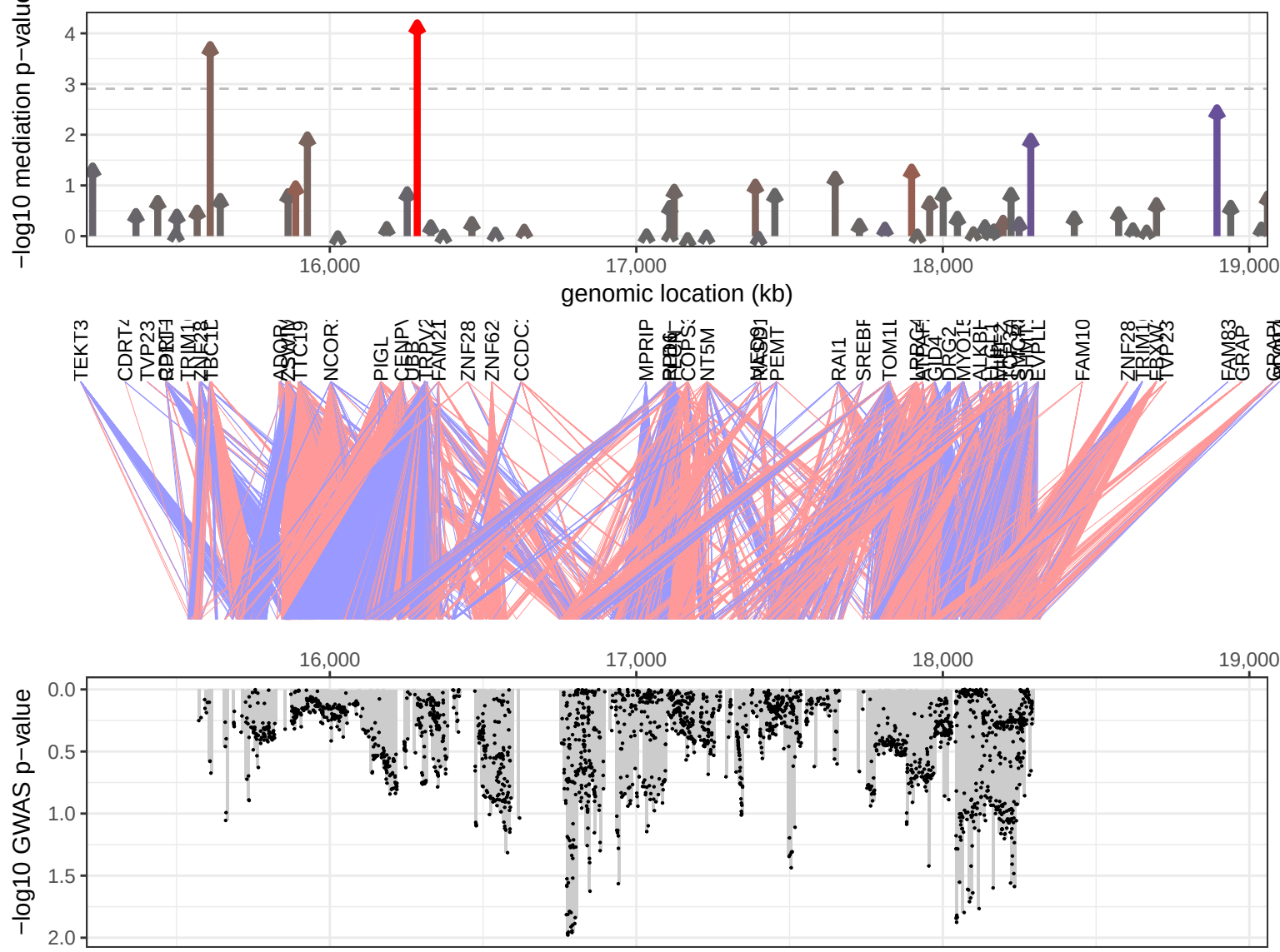
−log10 mediation p−value
4
3
2
1
0
16,000
17,000
18,000
19,000
genomic location (kb)
TEKT3
CDRT4
TVP23
TBC1D
NCOR2
PIGL
ZNF28
ZNF62
CCDC
MPRIP
COPS5
NT5M
PEMT
RAI1
SREBF
TOM1L
DRG2
MYO1E
FAM10
ZNF28
TVP23
FAM83
GRAP
16,000
17,000
18,000
19,000
−log10 GWAS p−value
0.0
0.5
1.0
1.5
2.0