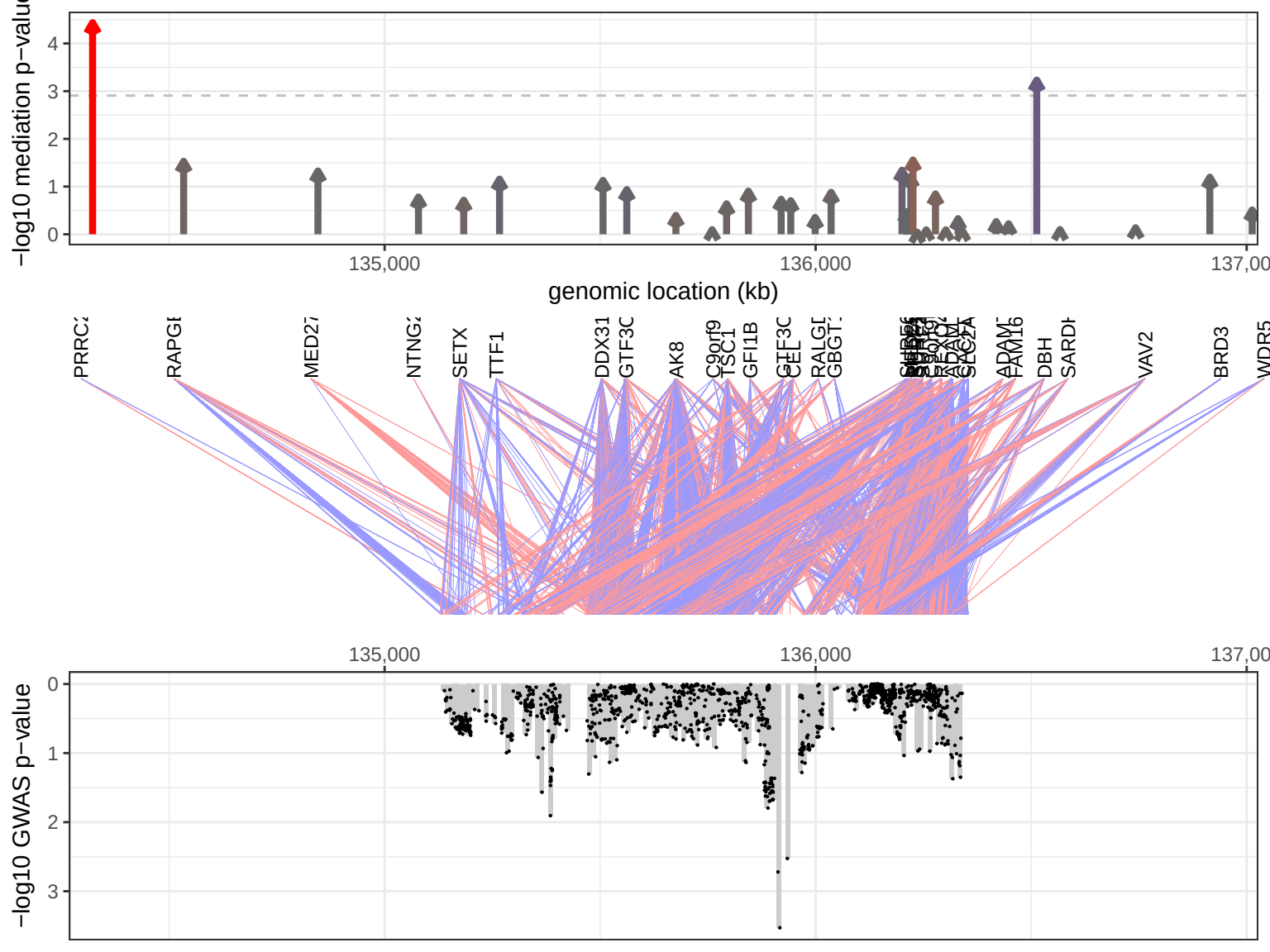
−log10 mediation p−value
4
3
2
1
0
135,000
136,000
genomic location (kb)
PRRC2
RAPGE
MED27
NTNG2
SETX
TTF1
DDX31
GTF3C
AK8
C9orf9
TSC1
GFI1B
GTF3C
CEL
RALGD
GBGT1
ADAM
FAM16
DBH
SARDH
VAV2
BRD3
WDR5
135,000
136,000
−log10 GWAS p−value
0
1
2
3