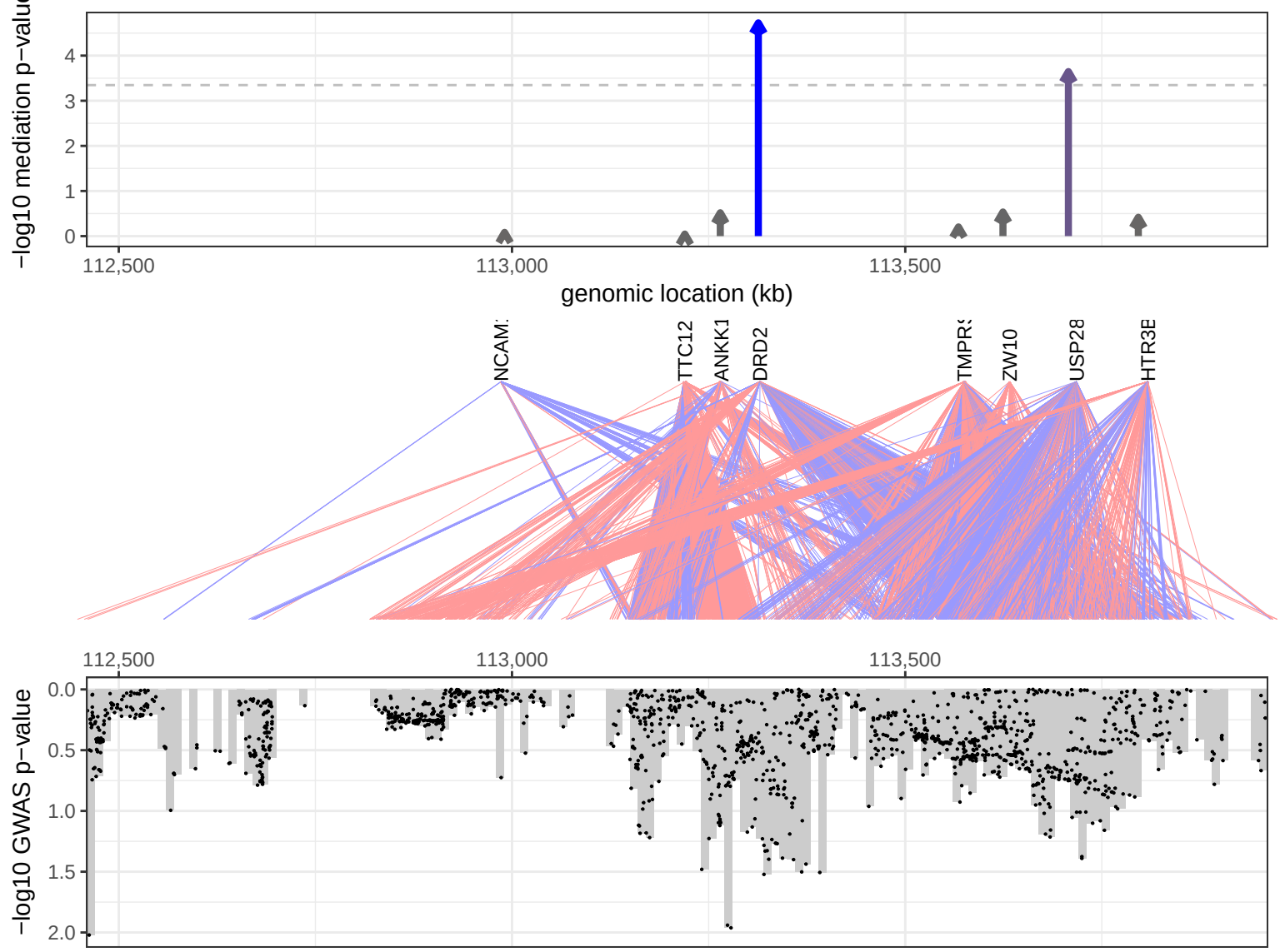
−log10 mediation p−value
4
3
2
1
0
112,500
113,000
113,500
genomic location (kb)
NCAM
TTC12
ANKK1
DRD2
TMPRS
ZW10
USP28
HTR3E
112,500
113,000
113,500
−log10 GWAS p−value
0.0
0.5
1.0
1.5
2.0