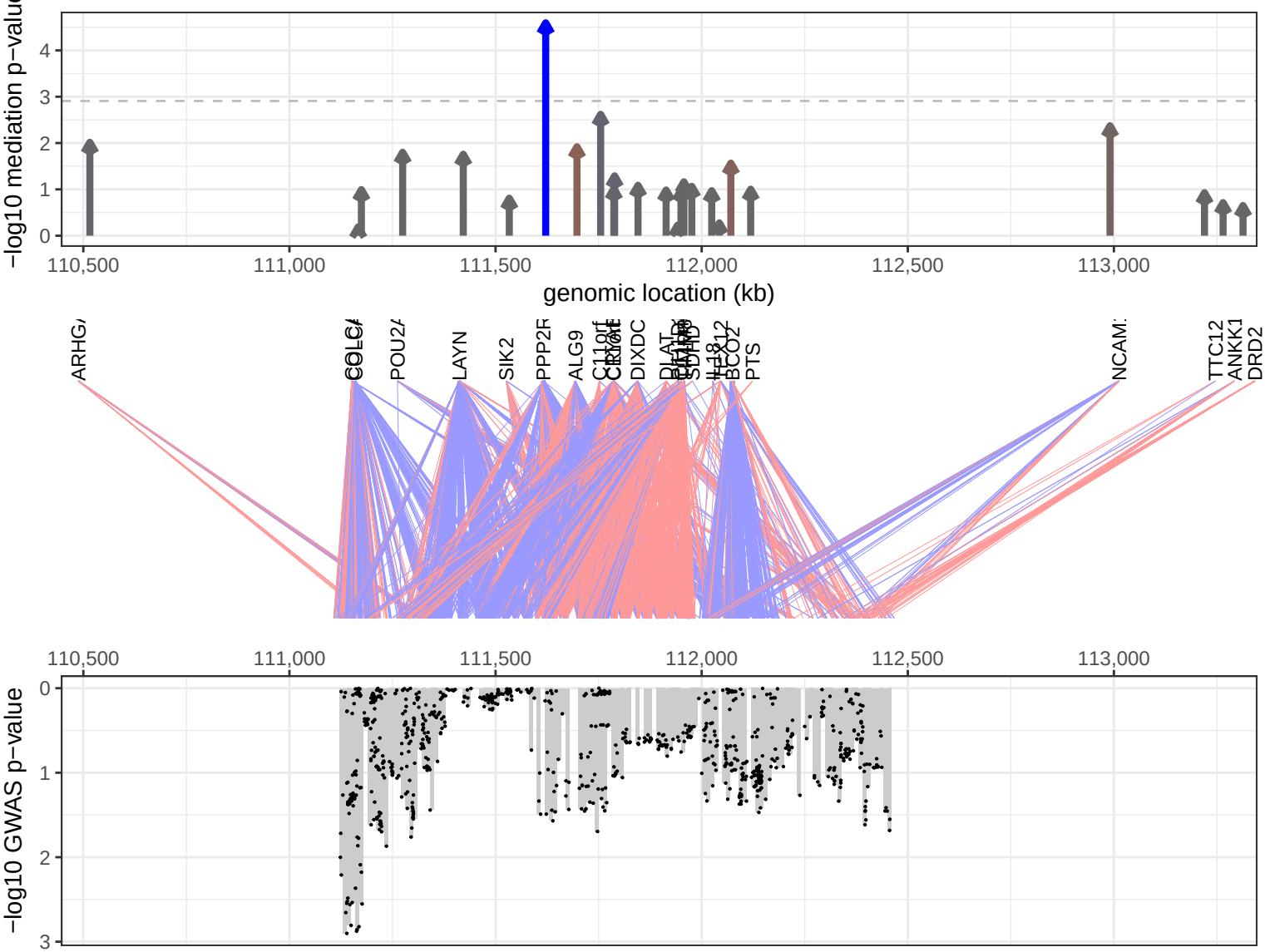
−log10 mediation p−value
0
1
2
3
4
110,500
111,000
111,500
112,000
112,500
113,000
genomic location (kb)
ARHG
POU2
LAYN
SIK2
PPP2R
ALG9
DIXDC
BCO2
PTS
NCAM
TTC12
ANKK1
DRD2
110,500
111,000
111,500
112,000
112,500
113,000
−log10 GWAS p−value
0
1
2
3