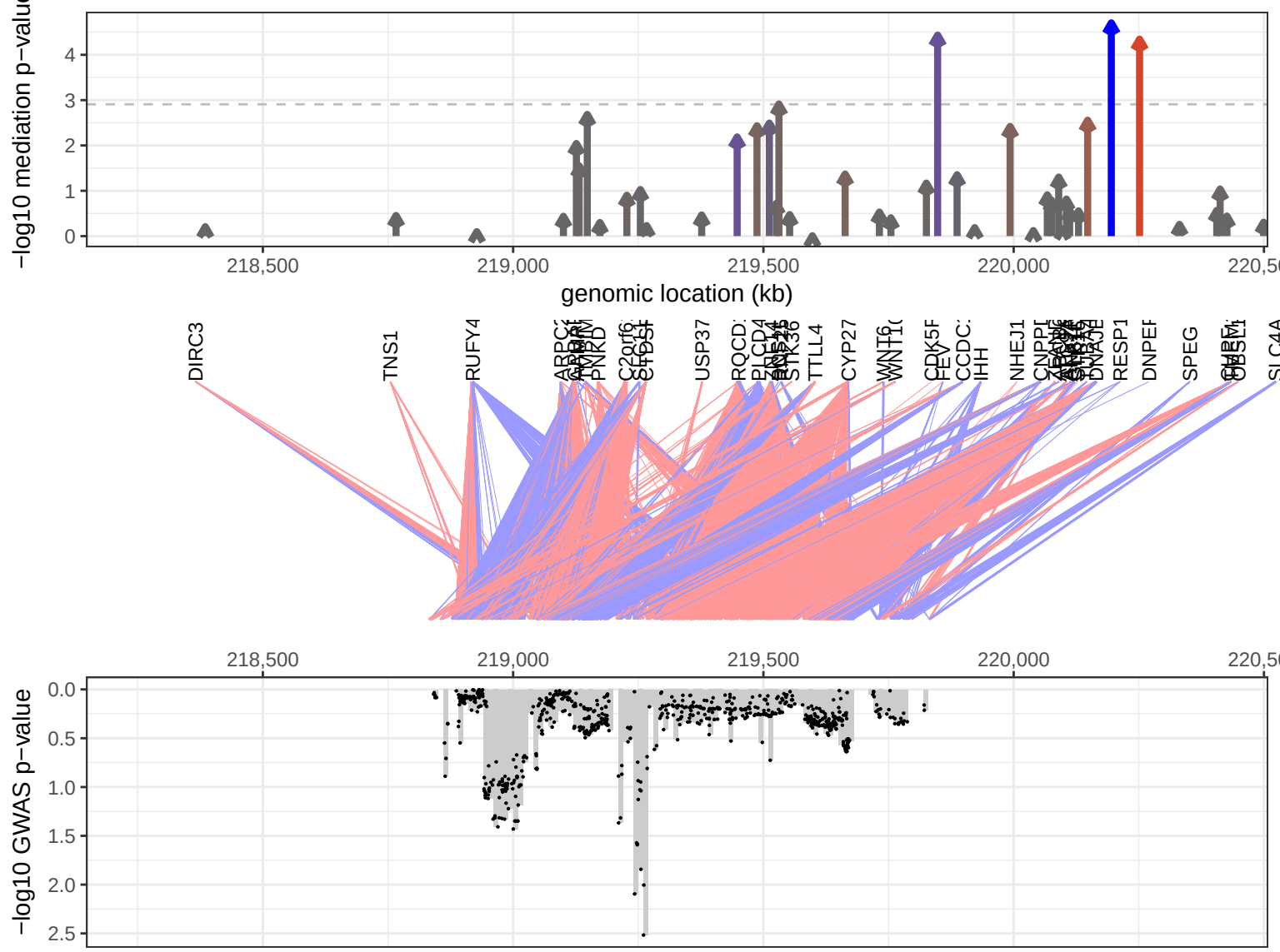
−log10 mediation p−value
0
1
2
3
4
218,500
219,000
219,500
220,000
genomic location (kb)
DIRC3
TNS1
RUFY4
USP37
TTLL4
CYP27
NHEJ1
RESP1
SPEG
218,500
219,000
219,500
220,000
−log10 GWAS p−value
0.0
0.5
1.0
1.5
2.0
2.5