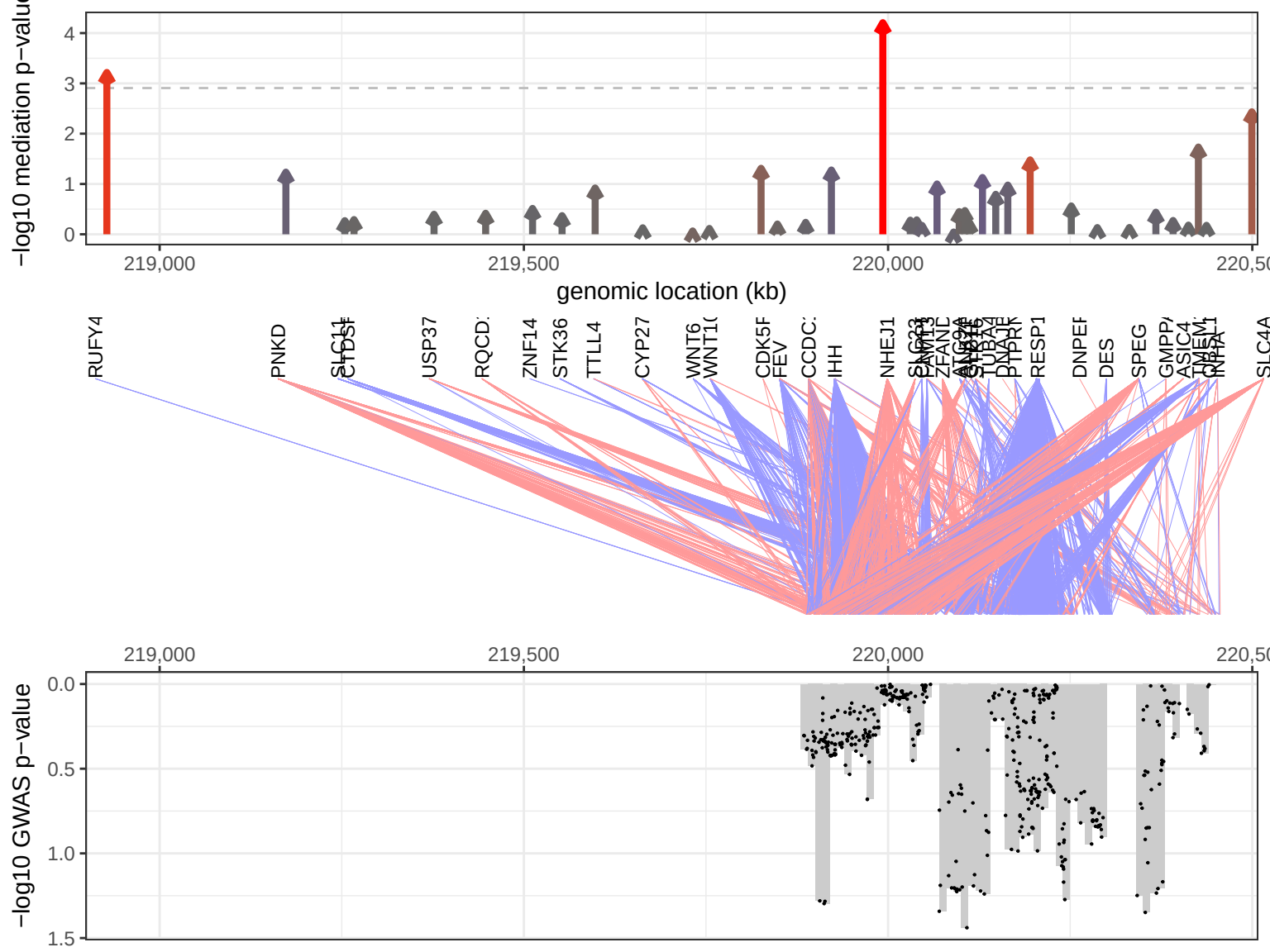
−log10 mediation p−value
4
3
2
1
0
219,000
219,500
220,000
genomic location (kb)
RUFY4
PNKD
USP37
ZNF14
STK36
TTLL4
CYP27
CDK5F
FEV
IHH
NHEJ1
RESP1
DNPEF
DES
SPEG
−log10 GWAS p−value
0.0
0.5
1.0
1.5