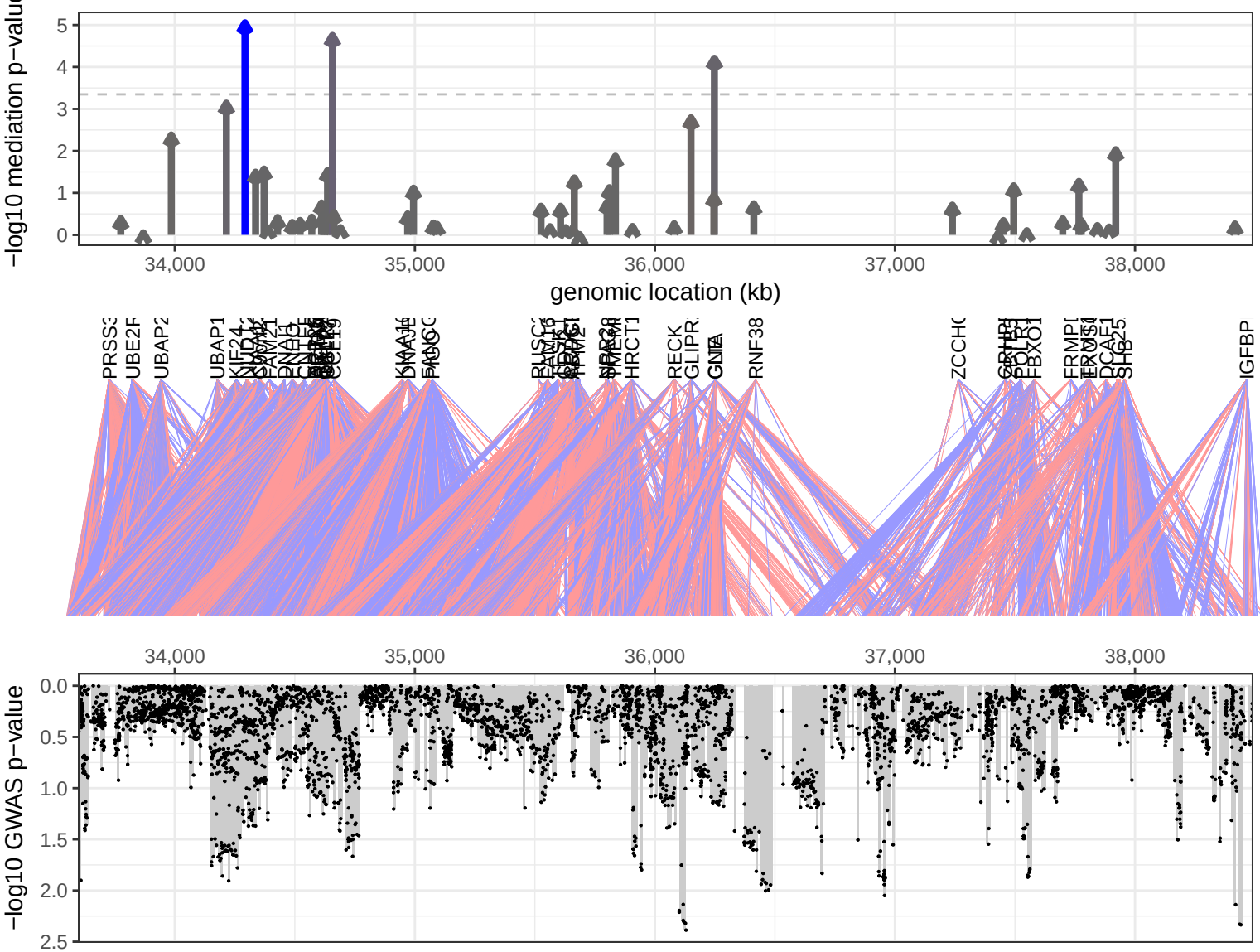
−log10 mediation p−value
5
4
3
2
1
0
34,000
35,000
36,000
37,000
38,000
genomic location (kb)
PRSS
UBE2F
UBAP1
RNF38
ZCCHC
IGFBP
34,000
35,000
36,000
37,000
38,000
−log10 GWAS p−value
0.0
0.5
1.0
1.5
2.0
2.5