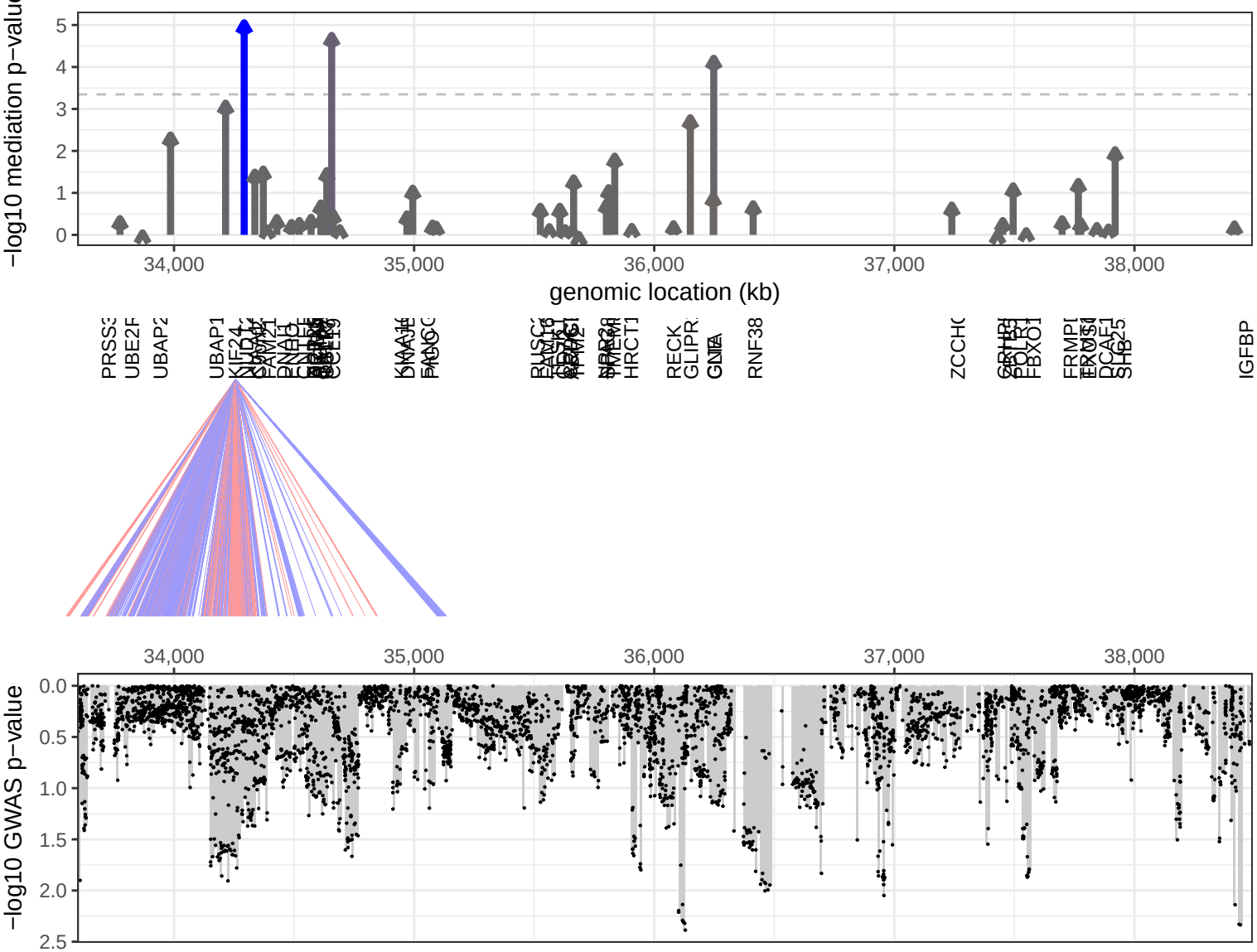
−log10 mediation p−value
genomic location (kb)
34,000
35,000
36,000
37,000
38,000
PRSS
UBE2F
UBAP
UBAP1
KIF24
KIAA
HRCT1
RECK
GLIPR
RNF38
ZCCHC
FBXO1
FRMPD
IGFBP
−log10 GWAS p−value