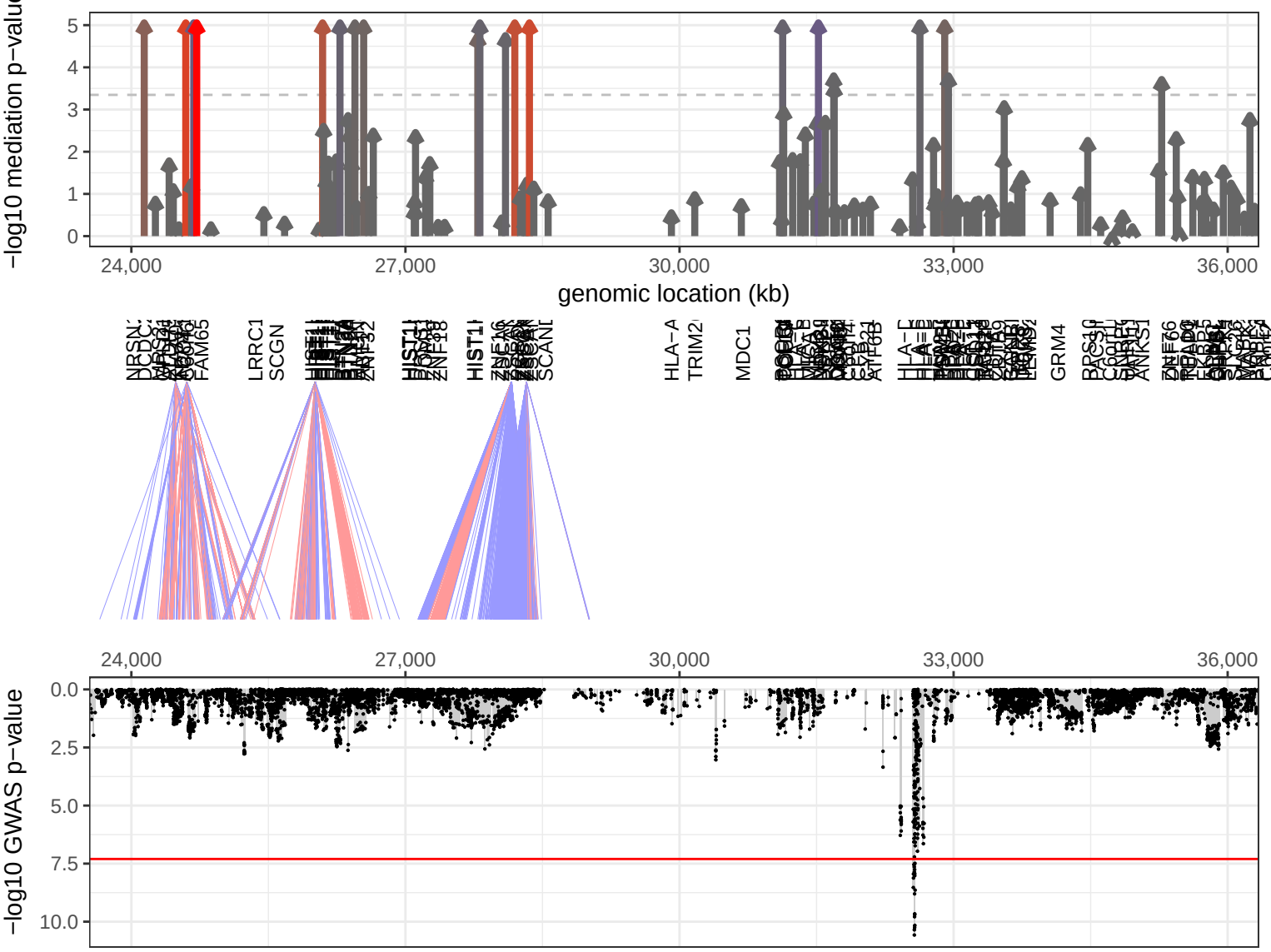
−log10 mediation p−value
genomic location (kb)
24,000
27,000
30,000
33,000
36,000
LRRC1
SCGN
HIST1
HLA−A
TRIM2
MDC1
GRM4
−log10 GWAS p−value