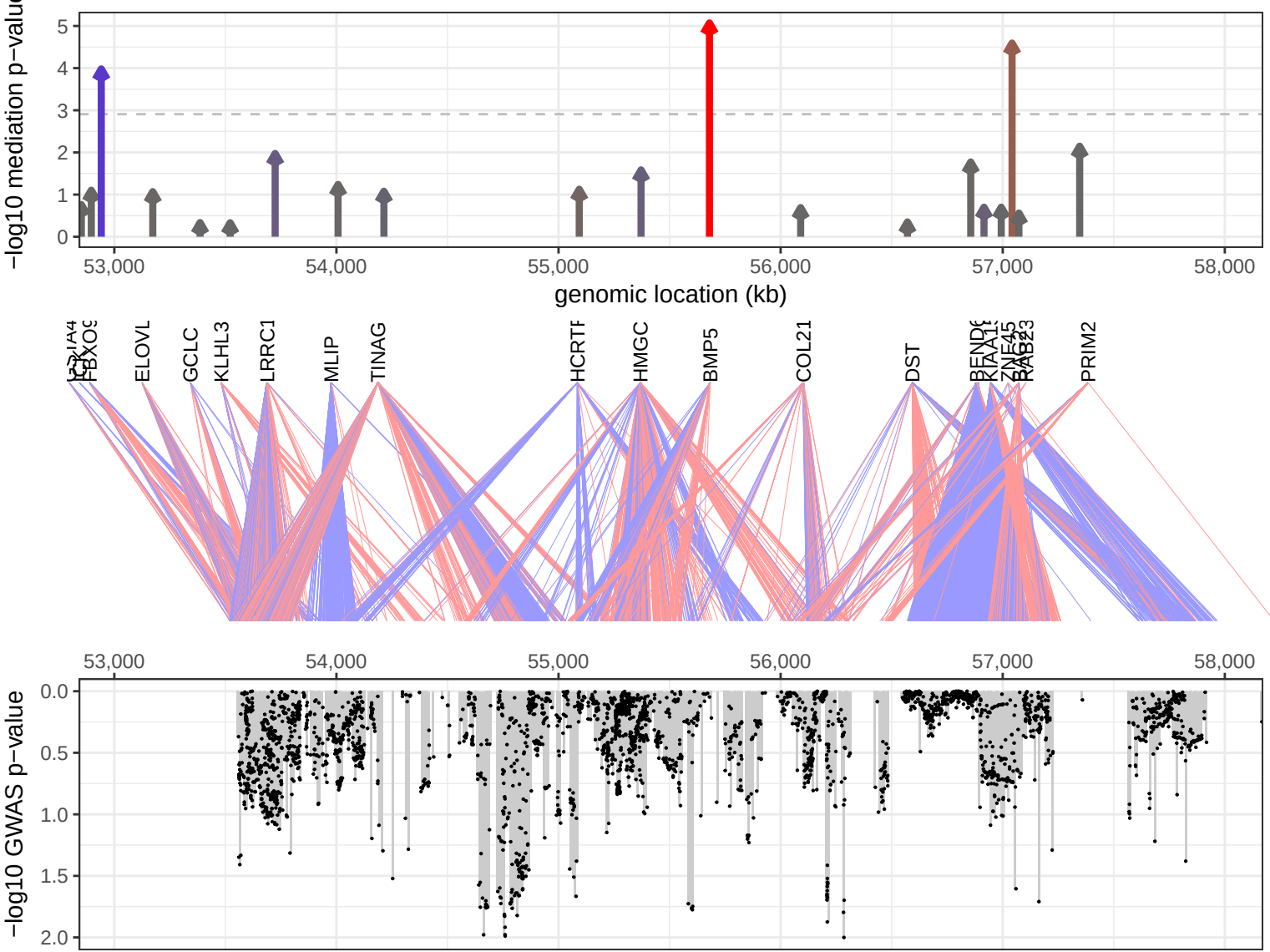
−log10 mediation p−value
genomic location (kb)
53,000
54,000
55,000
56,000
57,000
58,000
ELOVL
GCLC
KLHL3
LRRC1
MLIP
TINAG
HCRTF
HMGC
BMP5
COL21
DST
PRIM2
−log10 GWAS p−value