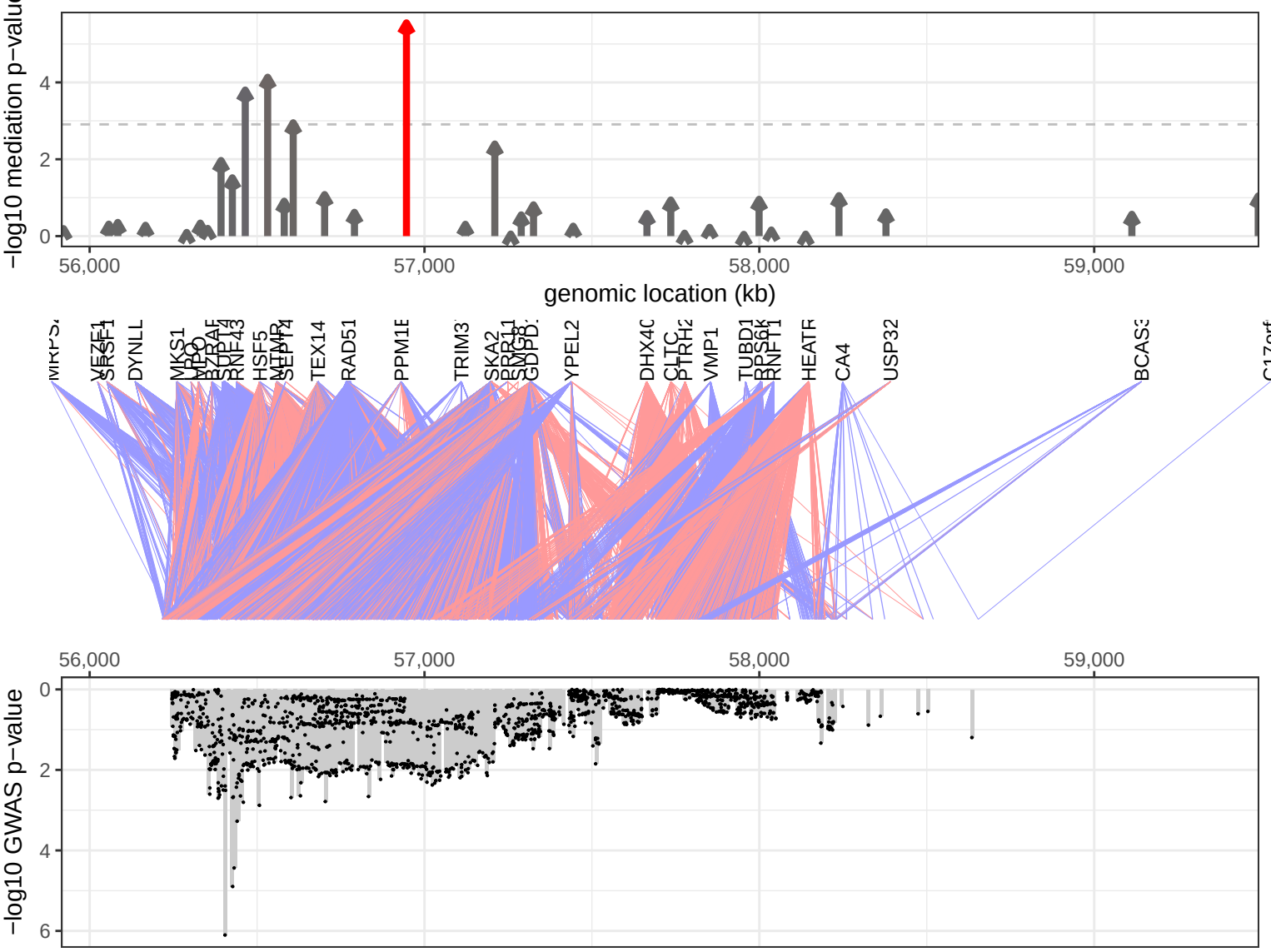
−log10 mediation p−value
0
2
4
56,000
57,000
58,000
59,000
genomic location (kb)
DYNLL
MKS1
RNF43
HSF5
TEX14
RAD51
PPM1E
TRIM3
SKA2
YPEL2
DHX40
VMP1
HEATR
CA4
USP32
BCAS3
−log10 GWAS p−value
0
2
4
6