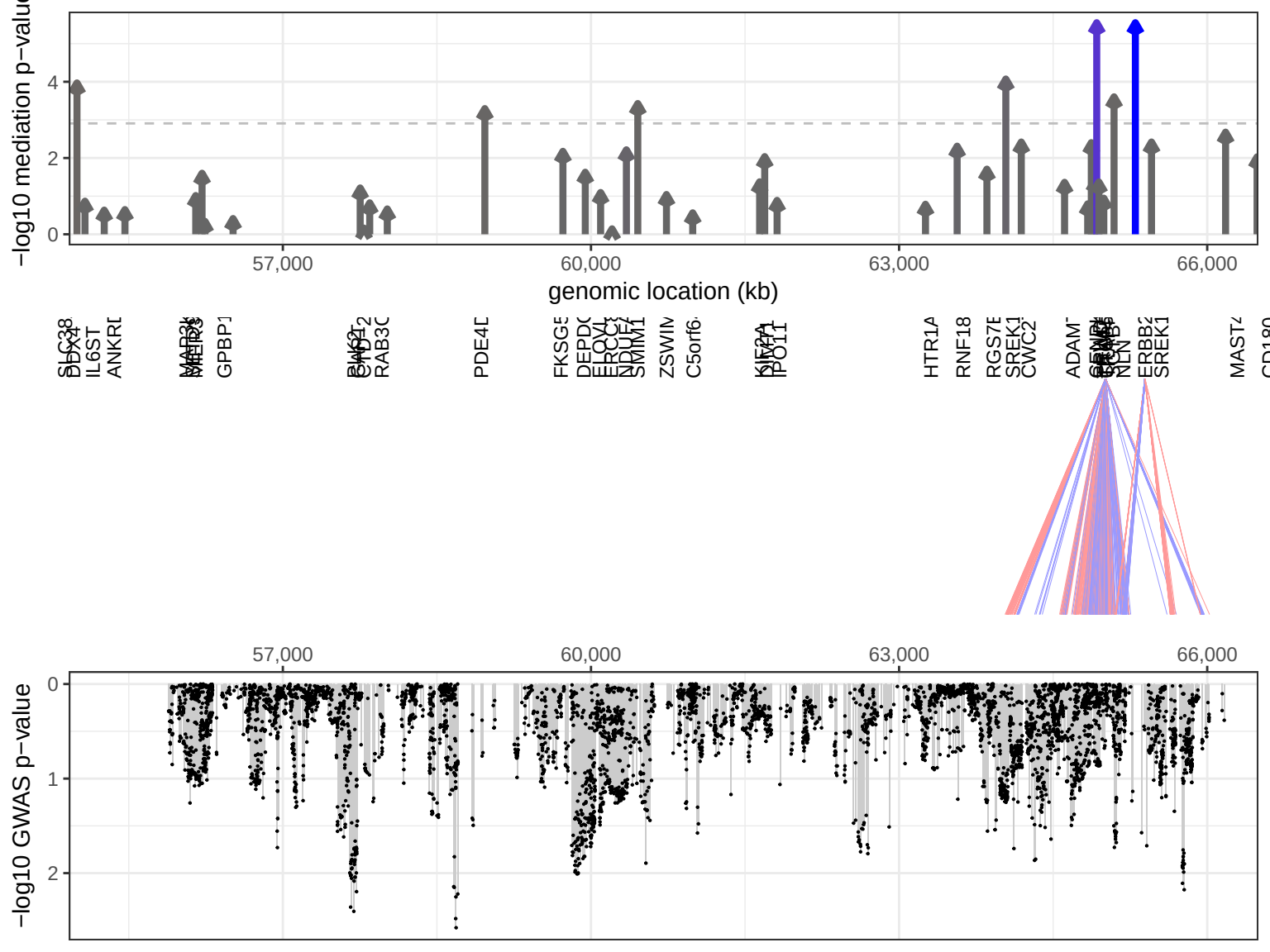
−log10 mediation p−value
0
2
4
57,000
60,000
63,000
66,000
genomic location (kb)
IL6ST
ANKRD
GPBP1
RAB3C
PDE4D
ZSWIM
C5orf6
HTR1A
RNF18
RGS7B
SREK1
CWC2
ADAM
NLN
ERBB2
SREK1
MAST4
−log10 GWAS p−value
0
1
2