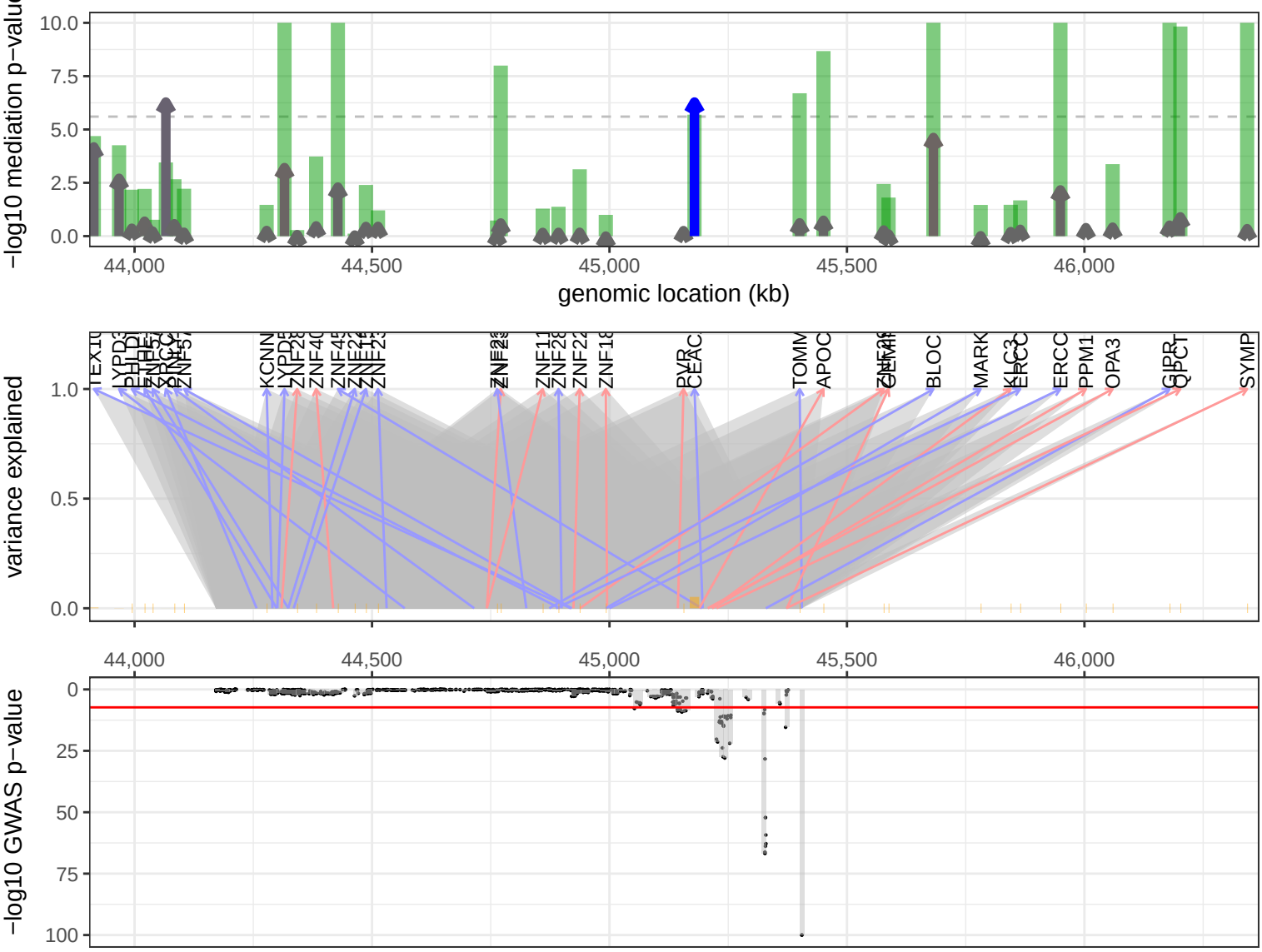
−log10 mediation p−value
10.0
7.5
5.0
2.5
0.0
44,000
44,500
45,000
45,500
46,000
genomic location (kb)
variance explained
1.0
0.5
0.0
KCNN
ZNF40
ZNF45
ZNF11
ZNF28
ZNF22
ZNF18
PVR
CEAC
TOMM
APOC
BLOC
MARK
ERCC
PPM1
OPA3
SYMP
−log10 GWAS p−value
0
25
50
75
100