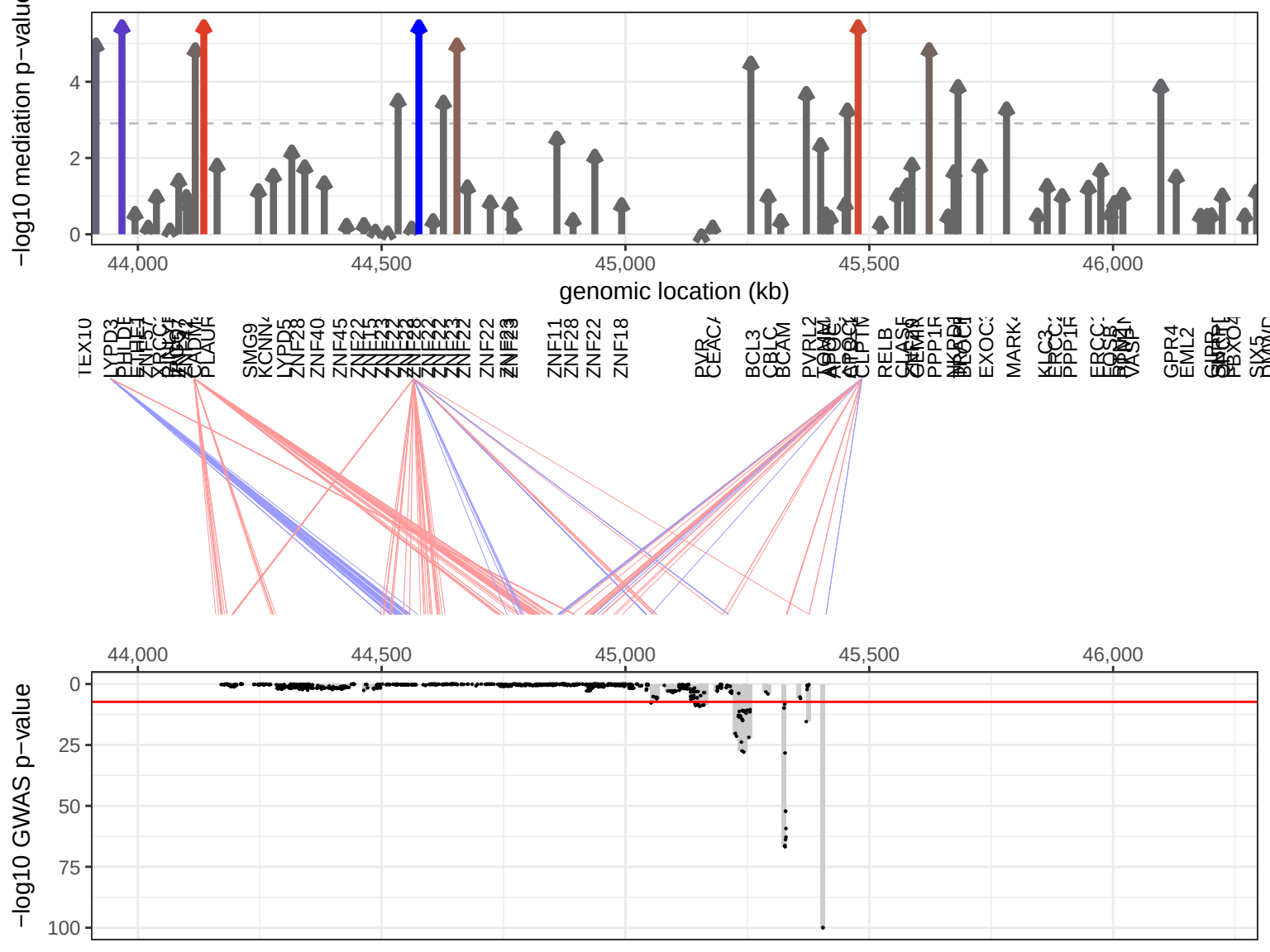
−log10 mediation p−value
0
2
4
44,000
44,500
45,000
45,500
46,000
genomic location (kb)
TEX10
LYPD3
SMG9
ZNF28
ZNF40
ZNF45
ZNF22
ZNF18
ZNF11
ZNF28
ZNF22
BCL3
CBLC
BCAM
RELB
EXOC
MARK
GPR4
EML2
−log10 GWAS p−value
0
25
50
75
100