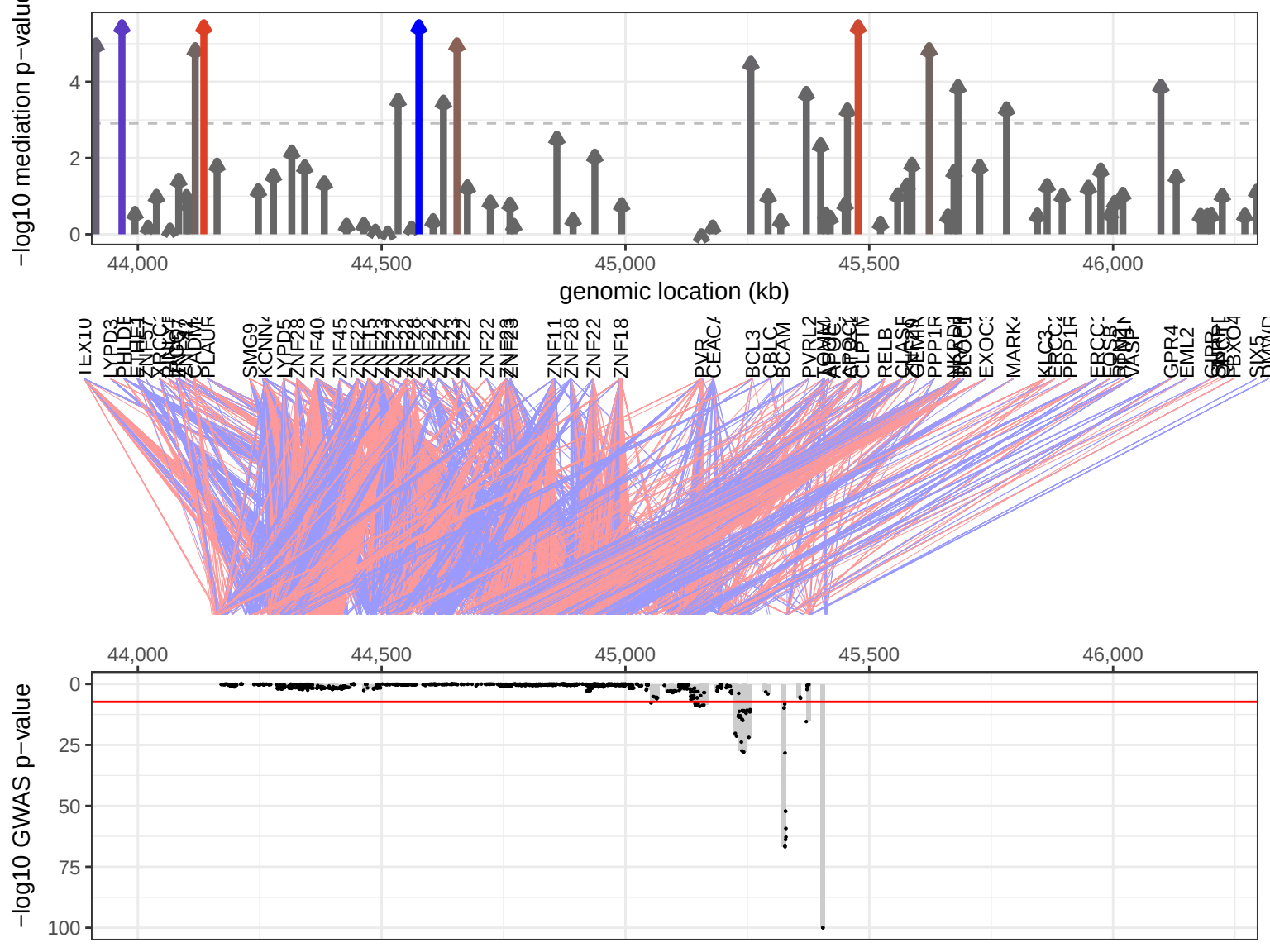
−log10 mediation p−value
0
2
4
44,000
44,500
45,000
45,500
46,000
genomic location (kb)
TEX10
LYPD3
SMG9
LYPD5
ZNF28
ZNF40
ZNF45
ZNF22
ZNF11
ZNF28
ZNF22
ZNF18
BCL3
CBLC
BCAM
RELB
EXOC
MARK
PPP1R
GPR4
EML2
SIX5
−log10 GWAS p−value
0
25
50
75
100