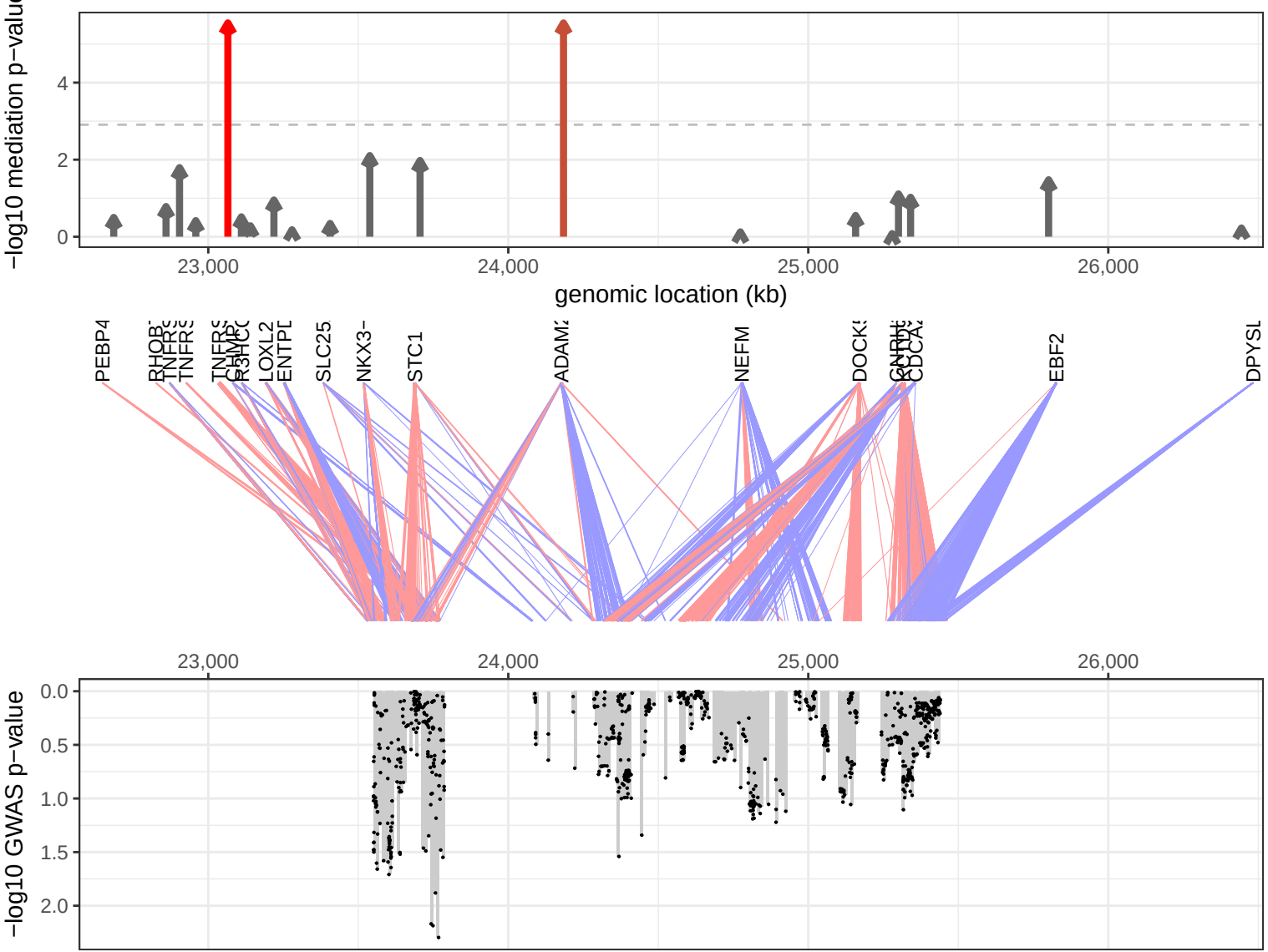
−log10 mediation p−value
0
2
4
23,000
24,000
25,000
26,000
genomic location (kb)
PEBP4
LOXL2
SLC25
STC1
NEFM
EBF2
DPYSL
−log10 GWAS p−value
0.0
0.5
1.0
1.5
2.0