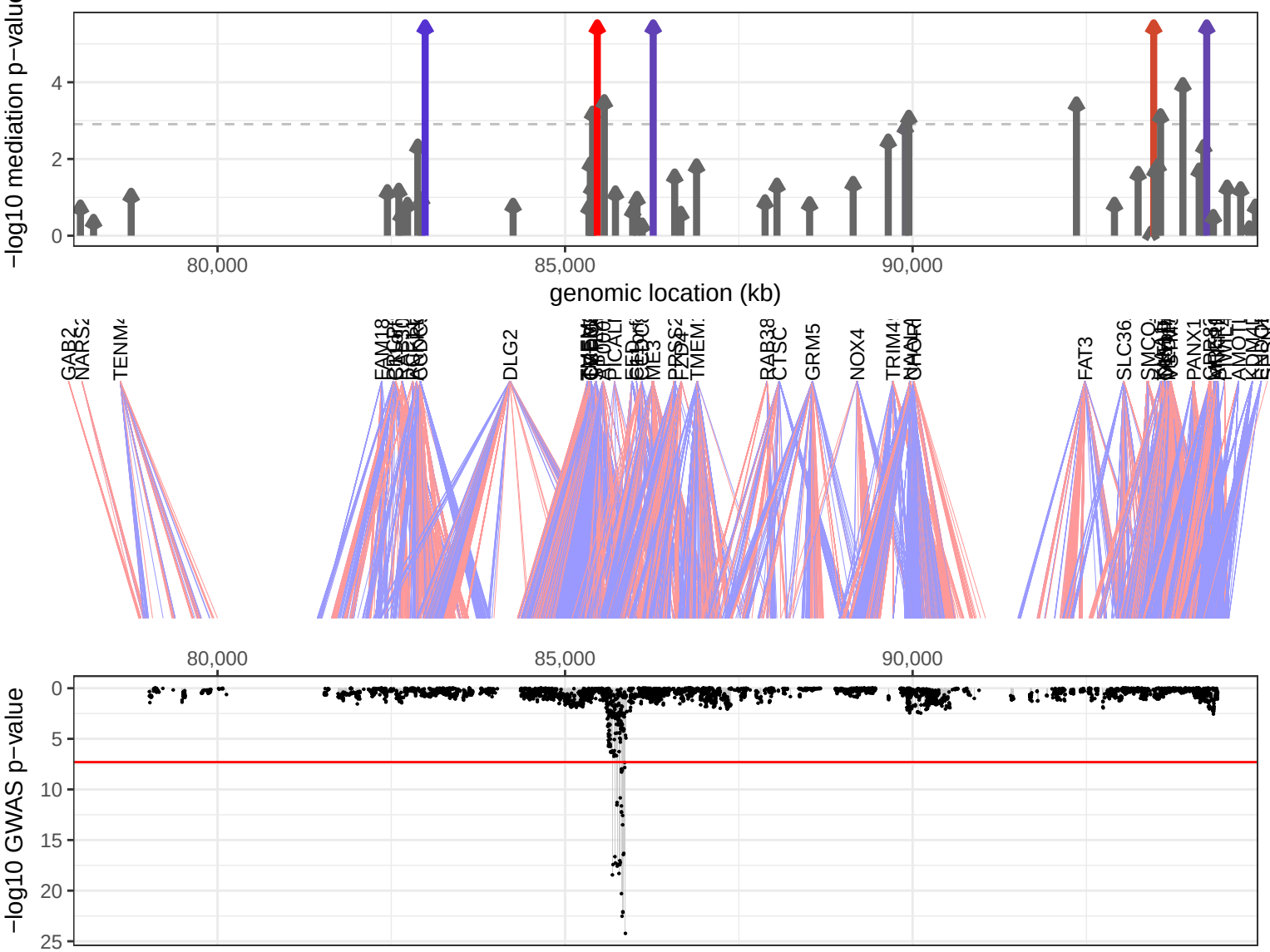
−log10 mediation p−value
0
2
4
80,000
85,000
90,000
genomic location (kb)
GAB2
TENM4
DLG2
ME3
TMEM
RAB38
CTSC
GRM5
NOX4
TRIM4
FAT3
SLC36
PANX1
80,000
85,000
90,000
−log10 GWAS p−value
0
5
10
15
20
25