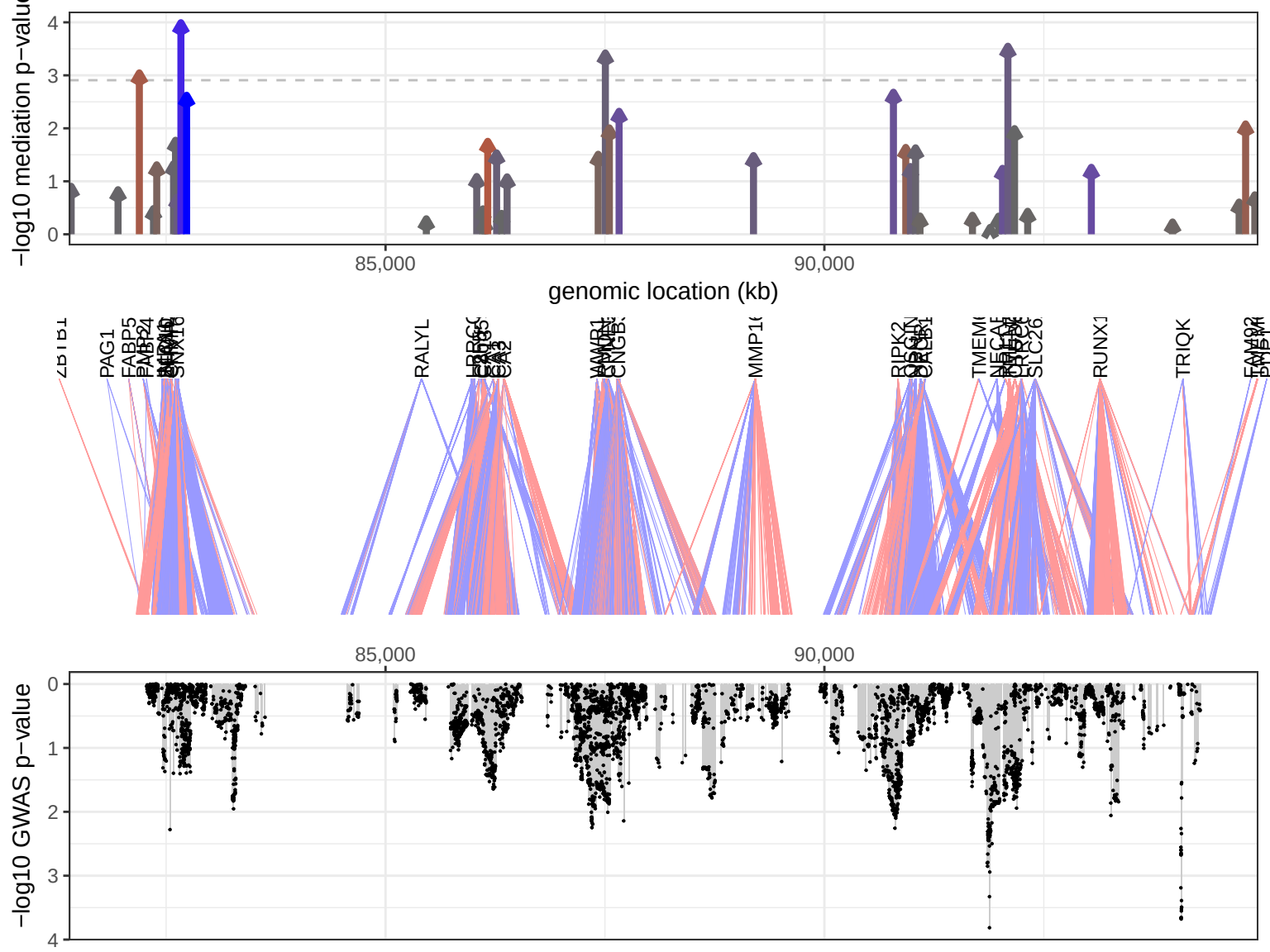
−log10 mediation p−value
0
1
2
3
4
85,000
90,000
genomic location (kb)
PAG1
RALYL
MMP16
RIPK2
TMEM
SLC26
RUNX1
TRIQK
85,000
90,000
−log10 GWAS p−value
0
1
2
3
4