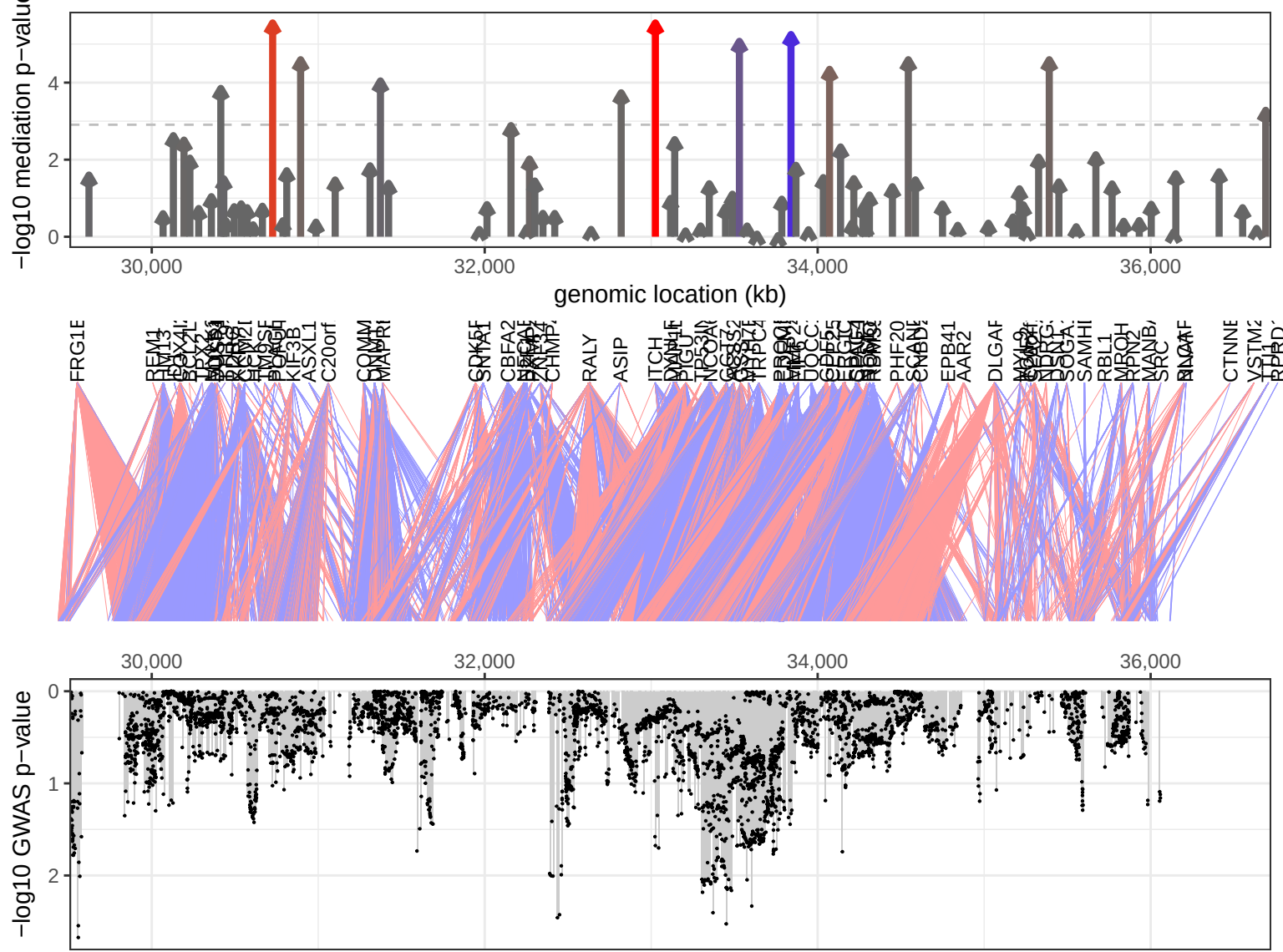
−log10 mediation p−value
0
2
4
30,000
32,000
34,000
36,000
genomic location (kb)
FRG1E
ASXL1
C20orf
RALY
ASIP
ITCH
EPB41
AAR2
DLGAP
SAMHD
RBL1
CTNNB
−log10 GWAS p−value
0
1
2