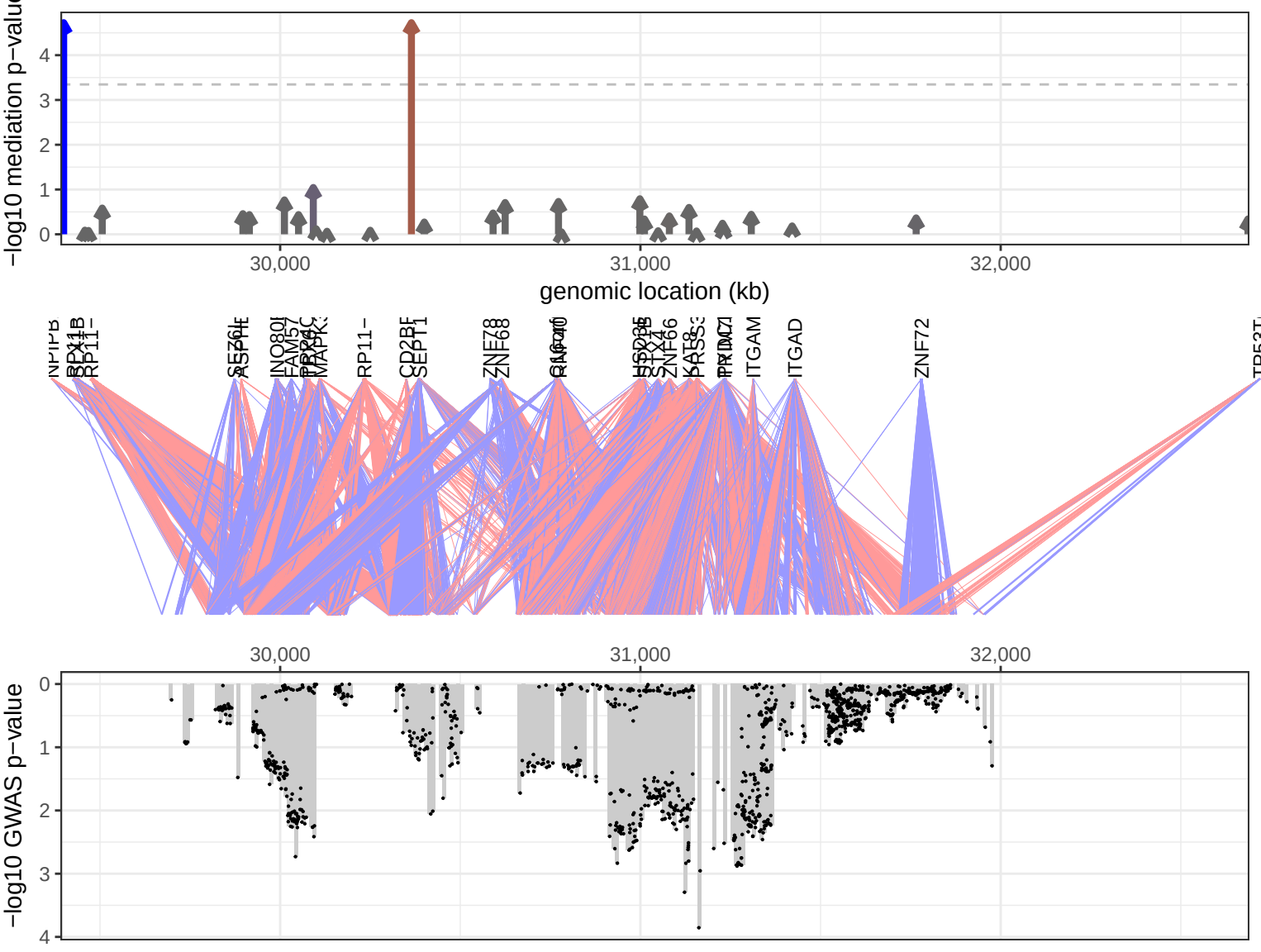
−log10 mediation p−value
0
1
2
3
4
30,000
31,000
32,000
genomic location (kb)
ITGAM
ITGAD
ZNF72
−log10 GWAS p−value
0
1
2
3
4
30,000
31,000
32,000