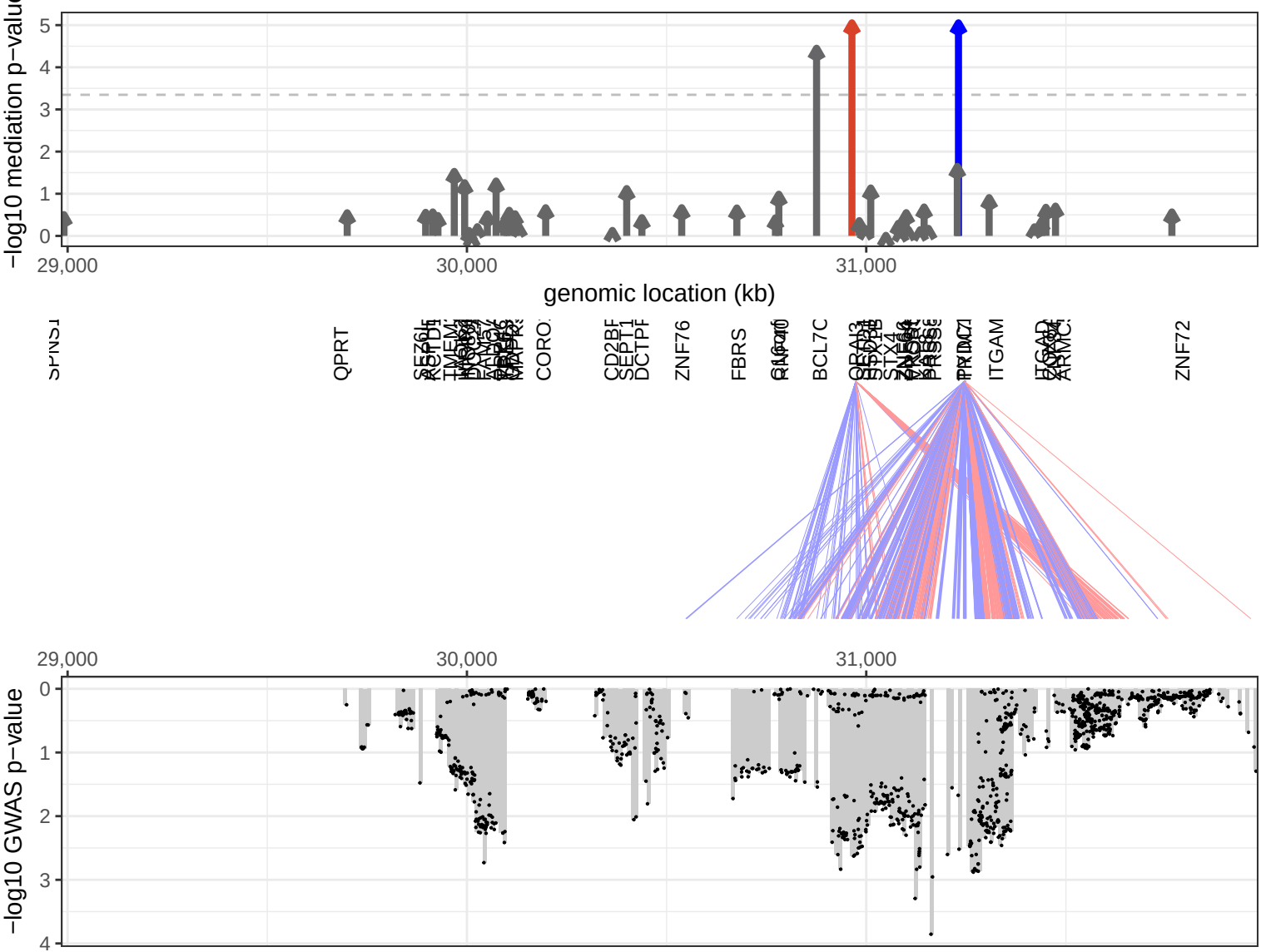
−log10 mediation p−value
5
4
3
2
1
0
29,000
30,000
31,000
genomic location (kb)
QPRT
CORO
CD2BP
SEPT1
DCTPP
ZNF76
FBRS
BCL7C
ITGAM
ZNF72
29,000
30,000
31,000
−log10 GWAS p−value
0
1
2
3
4