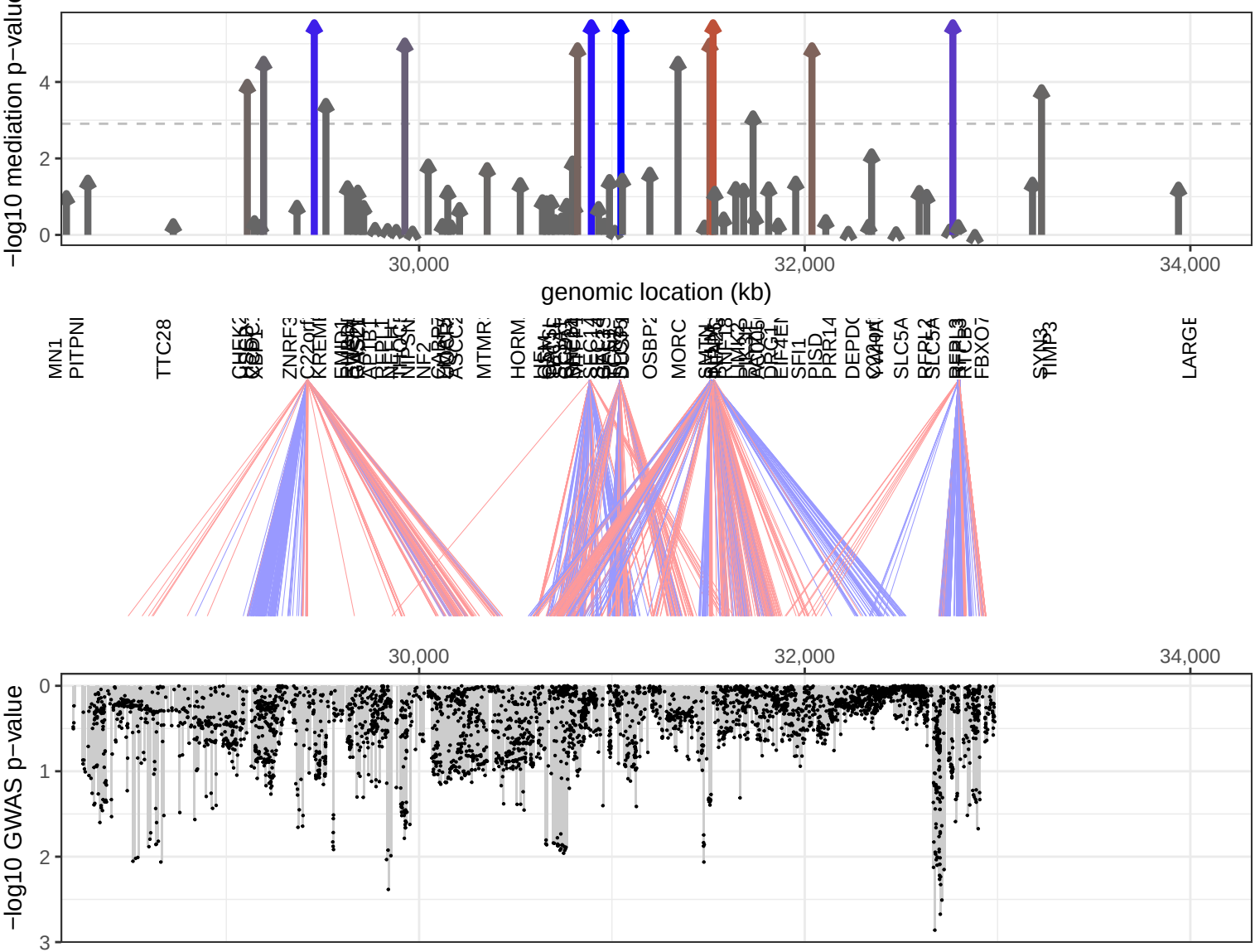
−log10 mediation p−value
0
2
4
30,000
32,000
34,000
genomic location (kb)
MN1
PITPNI
TTC28
ZNRF3
MTMR
HORM
OSBP2
MORC
SFI1
PRR14
DEPDC
SLC5A
FBXO7
SYN3
TIMP3
LARGE
30,000
32,000
34,000
−log10 GWAS p−value
0
1
2
3