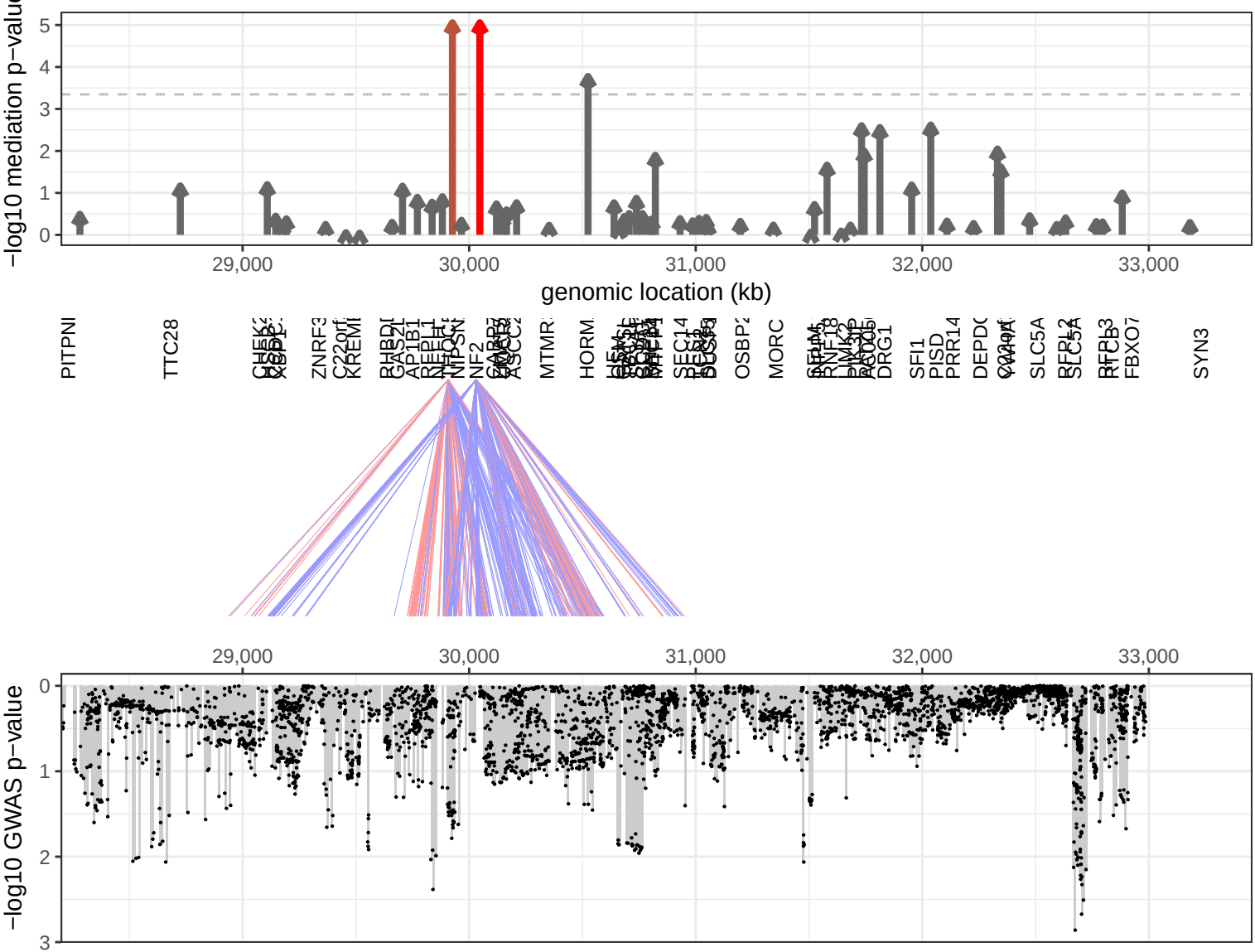
−log10 mediation p−value
genomic location (kb)
29,000
30,000
31,000
32,000
33,000
PITPNI
TTC28
ZNRF3
KREMI
AP1B1
NF2
MTMR
HORM
SEC14
OSBP2
MORC
DRG1
SFI1
PISD
PRR14
DEPDC
SLC5A
FBXO7
SYN3
−log10 GWAS p−value