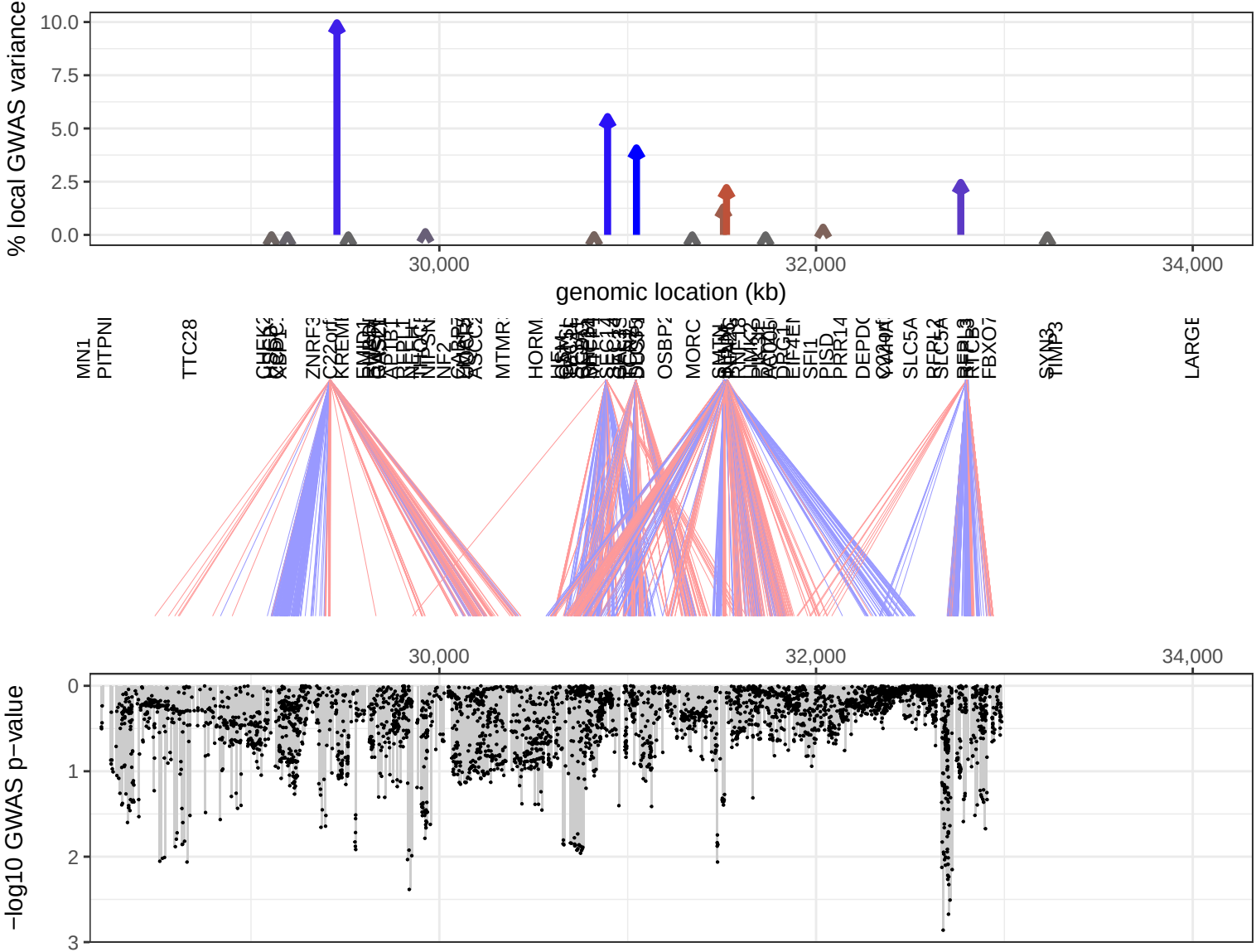
% local GWAS variance
10.0
7.5
5.0
2.5
0.0
30,000
32,000
34,000
genomic location (kb)
MN1
PITPNI
TTC28
ZNRF3
MTMR
HORM
OSBP
MORC
SFI1
PISD
PRR14
DEPDC
SLC5A
FBXO7
SYN3
TIMP3
LARGE
−log10 GWAS p−value
0
1
2
3