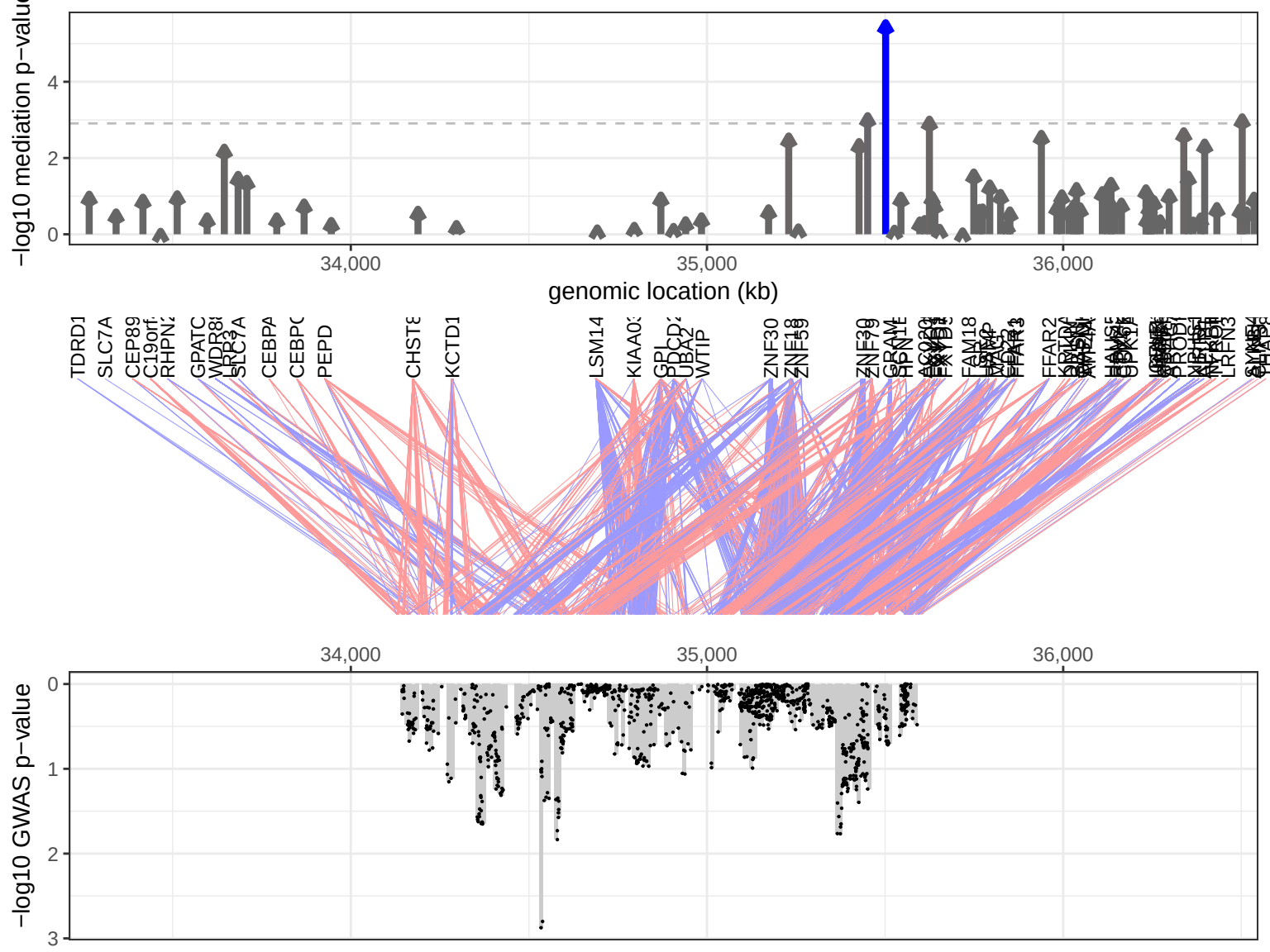
−log10 mediation p−value
0
2
4
34,000
35,000
36,000
genomic location (kb)
TDRD1
SLC7A
CEP89
C19orf
RHPN2
GPATC
WDR8
LRP3
SLC7A
CEBPA
CEBPC
PEPD
CHSTE
KCTD1
LSM14
KIAA0
GPI
PDCD
UBA2
WTIP
ZNF30
ZNF18
ZNF59
ZNF79
GRAM
HPN
FAM18
FFAR2
LRFN3
34,000
35,000
36,000
−log10 GWAS p−value
0
1
2
3