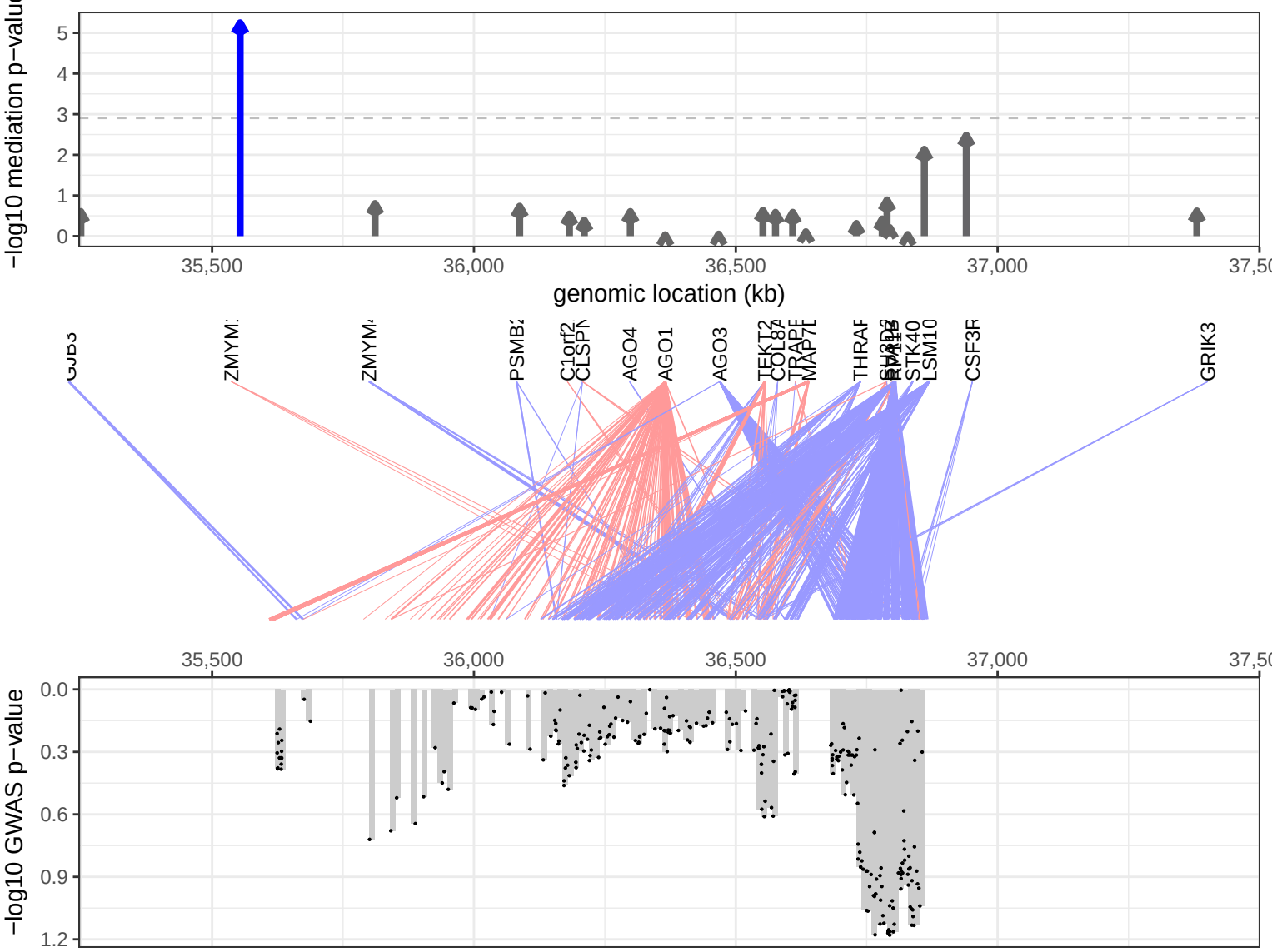
−log10 mediation p−value
genomic location (kb)
35,500
36,000
36,500
37,000
ZMYM
PSMB
C1orf2
CLSPN
AGO4
AGO1
AGO3
TEKT2
COL8A
TRAPPC
MAP7D
THRAP
STK40
LSM10
CSF3R
GRIK3
−log10 GWAS p−value