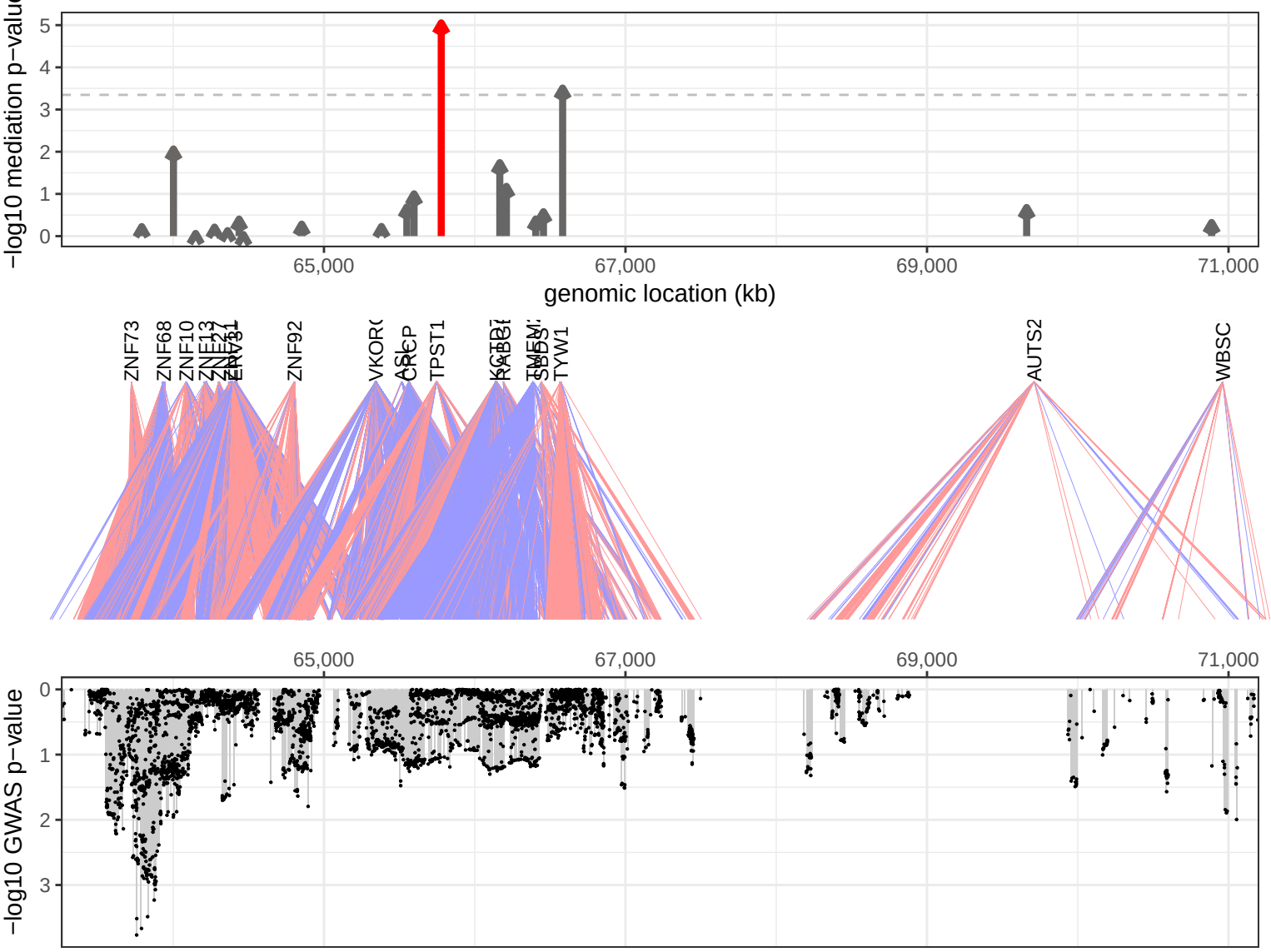
−log10 mediation p−value
5
4
3
2
1
0
65,000
67,000
69,000
71,000
genomic location (kb)
ZNF73
ZNF68
ZNF10
ZNF92
TPST1
TYW1
AUTS2
WBSC
65,000
67,000
69,000
71,000
−log10 GWAS p−value
0
1
2
3