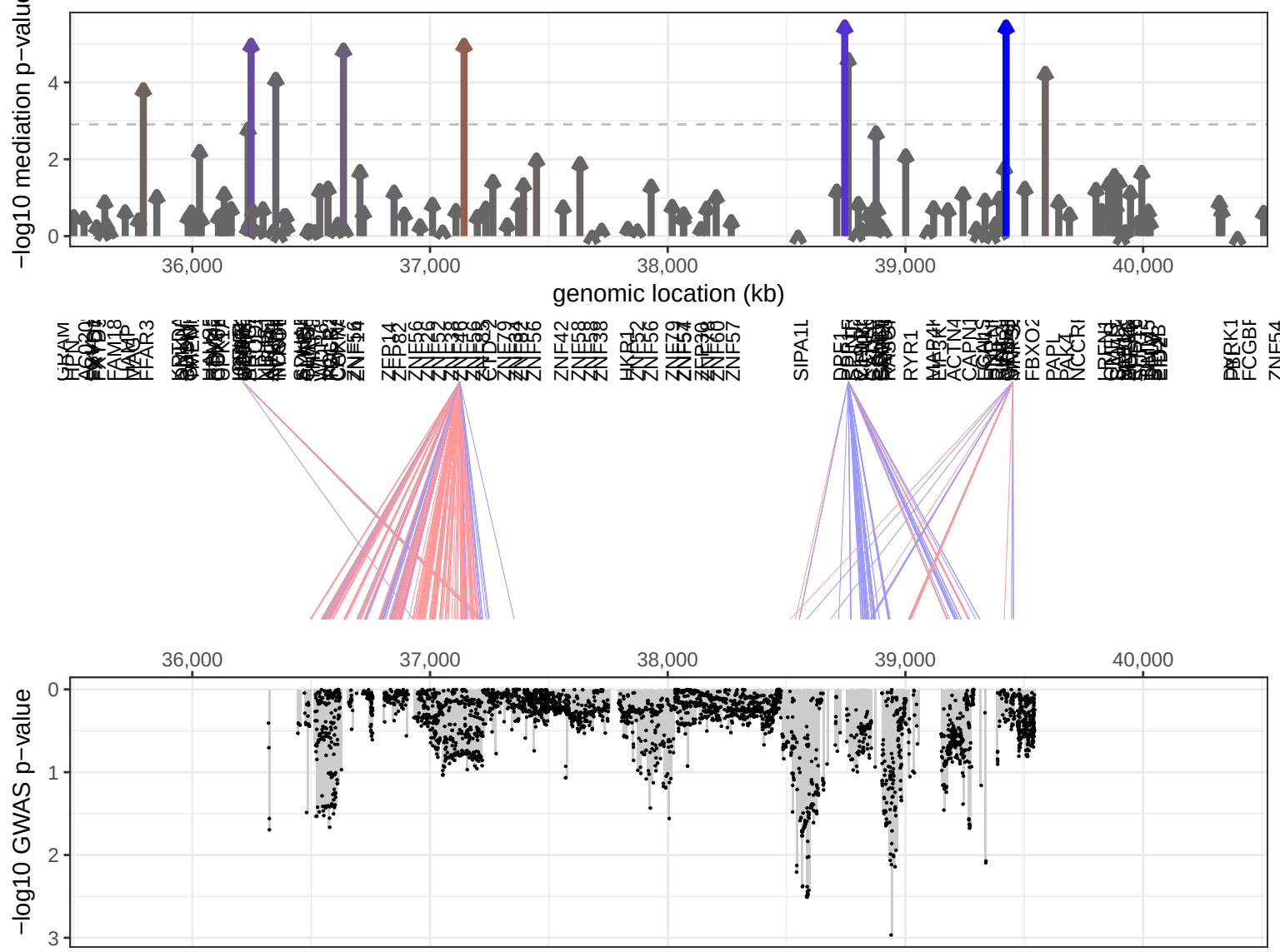
−log10 mediation p−value
0
2
4
36,000
37,000
38,000
39,000
40,000
genomic location (kb)
SIPA1L
RYR1
ACTN4
FBXO2
PAP1
NCCRP1
ZNF42
ZNF58
ZNF38
ZNF56
ZNF57
ZEB1
ZNF56
ZNF79
ZNF56
FFAR3
PARK1
FCGBP
ZNF54
−log10 GWAS p−value
0
1
2
3