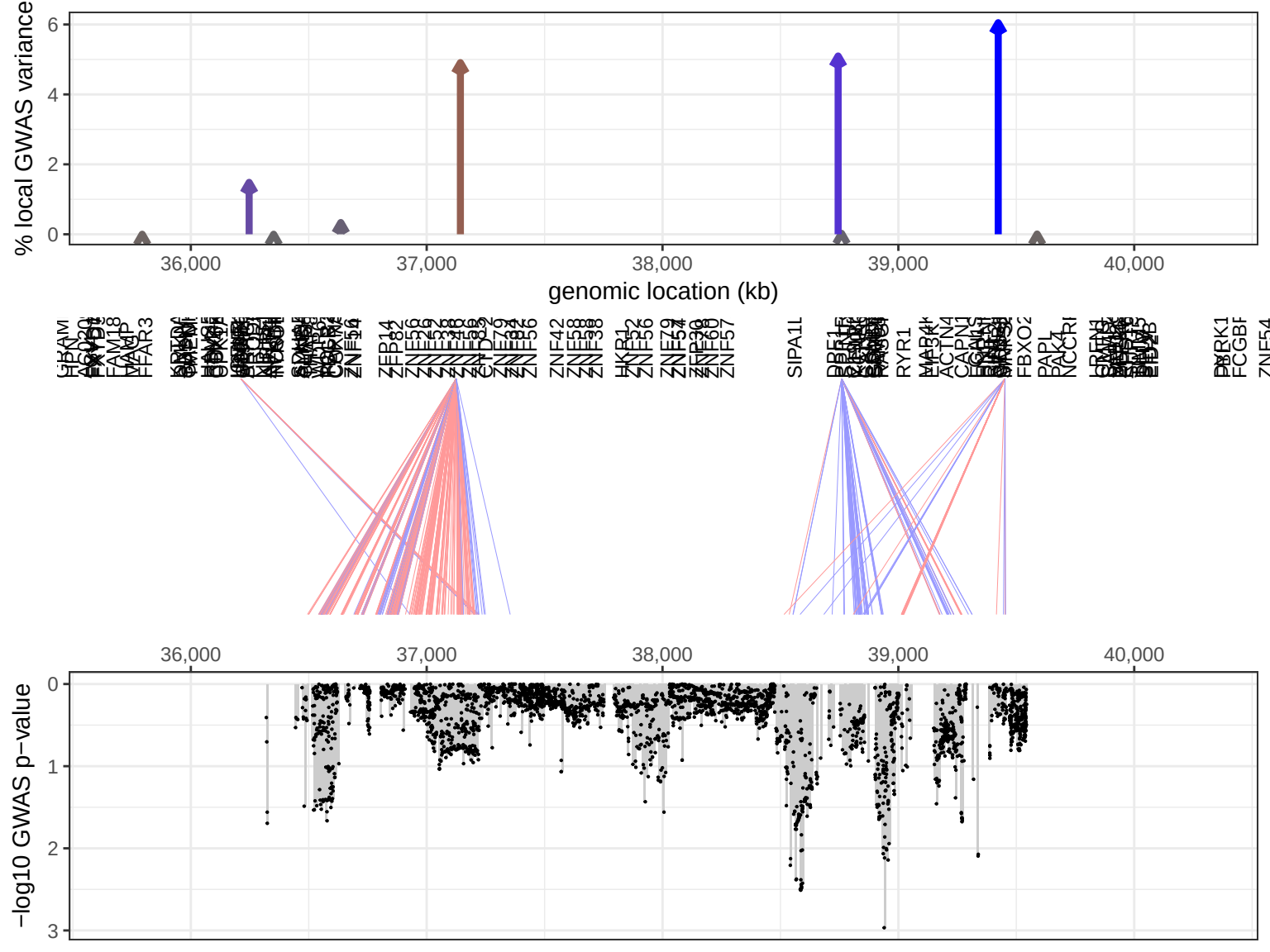
% local GWAS variance
0
2
4
6
36,000
37,000
38,000
39,000
40,000
genomic location (kb)
SIPA1L
RYR1
FBXO2
RAP1
NCCRP1
ACTN4
ZNF42
ZNF58
ZNF38
ZNF56
ZNF57
ZNF79
ZNF54
ZNF55
ZFP14
ZNF56
ZNF56
ZNF57
FFAR3
PARK1
FCGBP
ZNF54
−log10 GWAS p−value
0
1
2
3