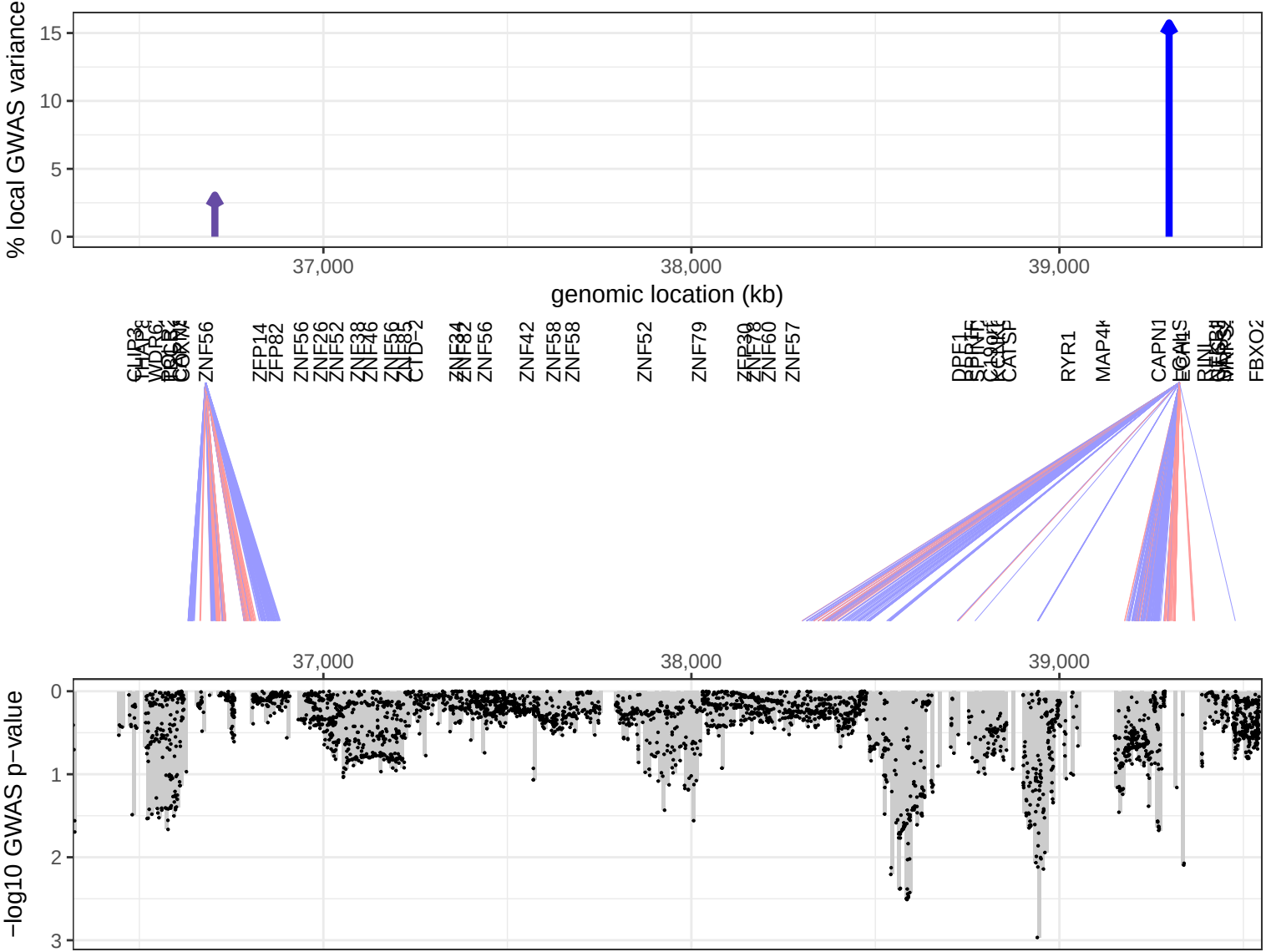
% local GWAS variance
15
10
5
0
37,000
38,000
39,000
genomic location (kb)
ZNF56
ZFP14
ZFP82
ZNF56
ZNF26
ZNF52
ZNF42
ZNF58
ZNF58
ZNF52
ZNF79
ZNF57
RYR1
MAP4
CAPN1
FBXO2
37,000
38,000
39,000
−log10 GWAS p−value
0
1
2
3