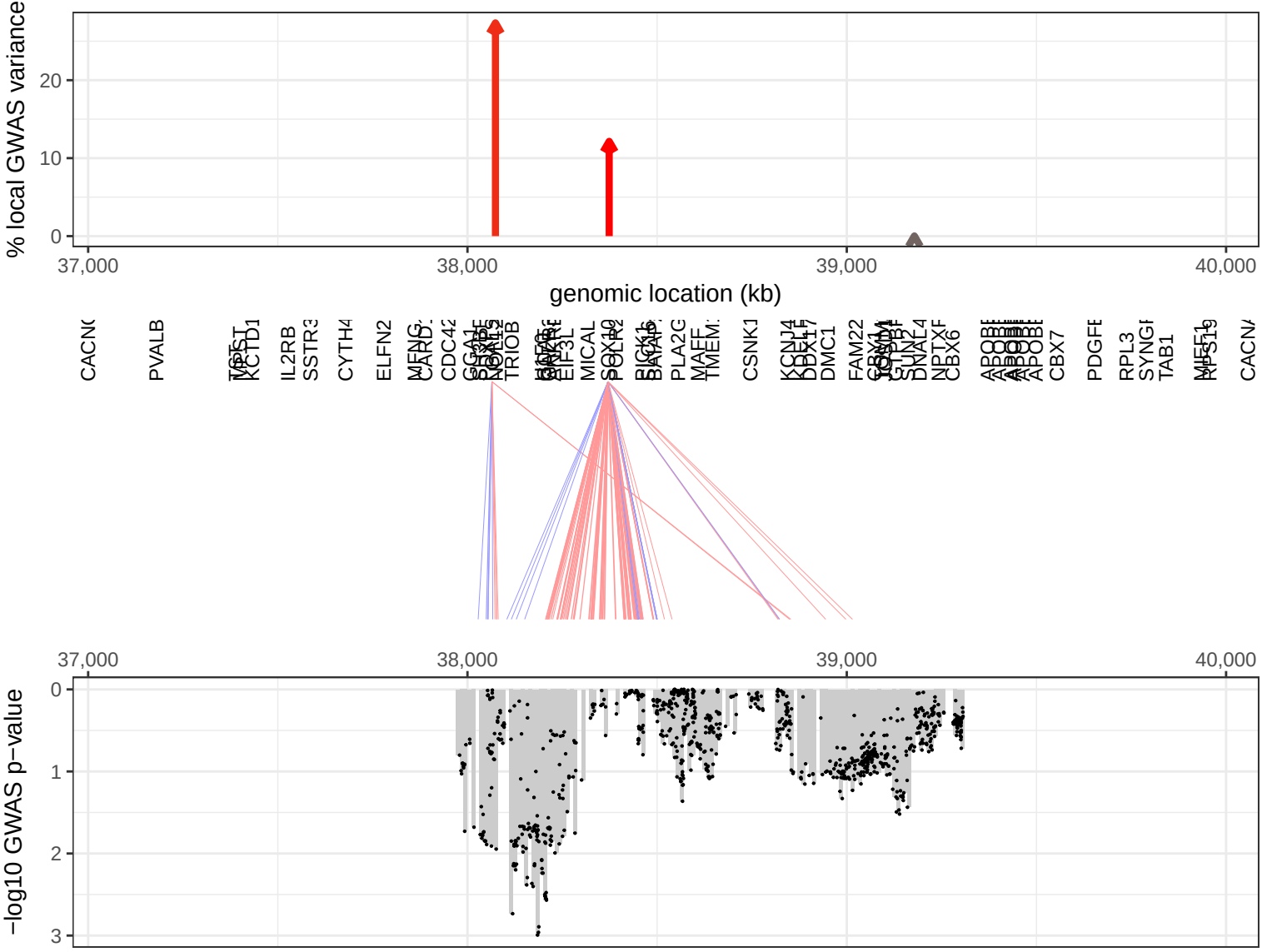
% local GWAS variance
0
10
20
37,000
38,000
39,000
40,000
genomic location (kb)
CACN
PVALB
IL2RB
CYTH4
ELFN2
TRIOB
MICAL
PLA2G
CSNK1
DMC1
FAM22
CBX7
PDGFE
RPL3
TAB1
CACN
−log10 GWAS p−value
0
1
2
3