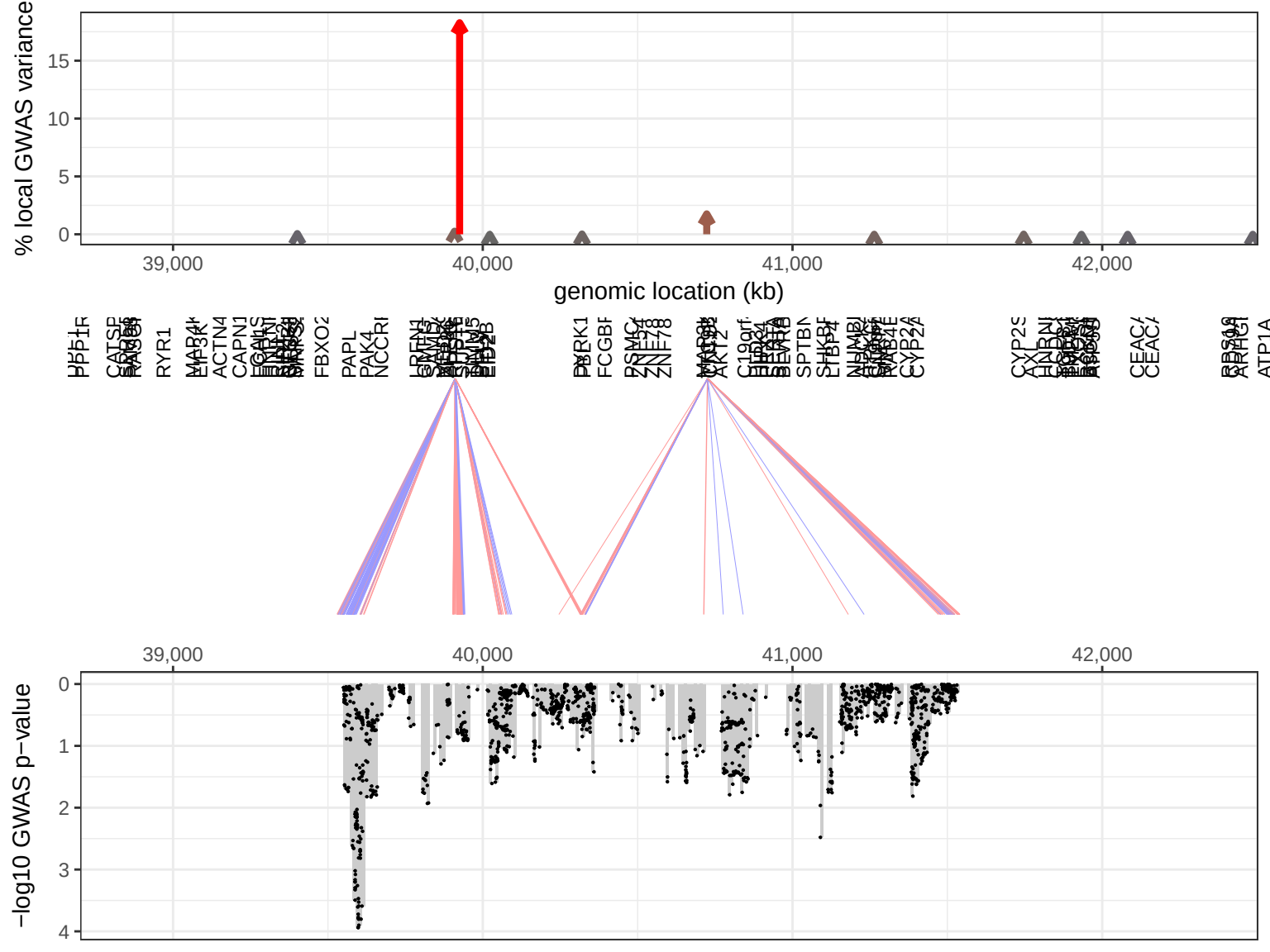
% local GWAS variance
15
10
5
0
39,000
40,000
41,000
42,000
genomic location (kb)
RYR1
ACTN4
CAPN1
FBXO2
PAPL
PAK4
NCCRI
FCGBP
SPTBN
CYP2A
CEAC
ATP1A
−log10 GWAS p−value
0
1
2
3
4