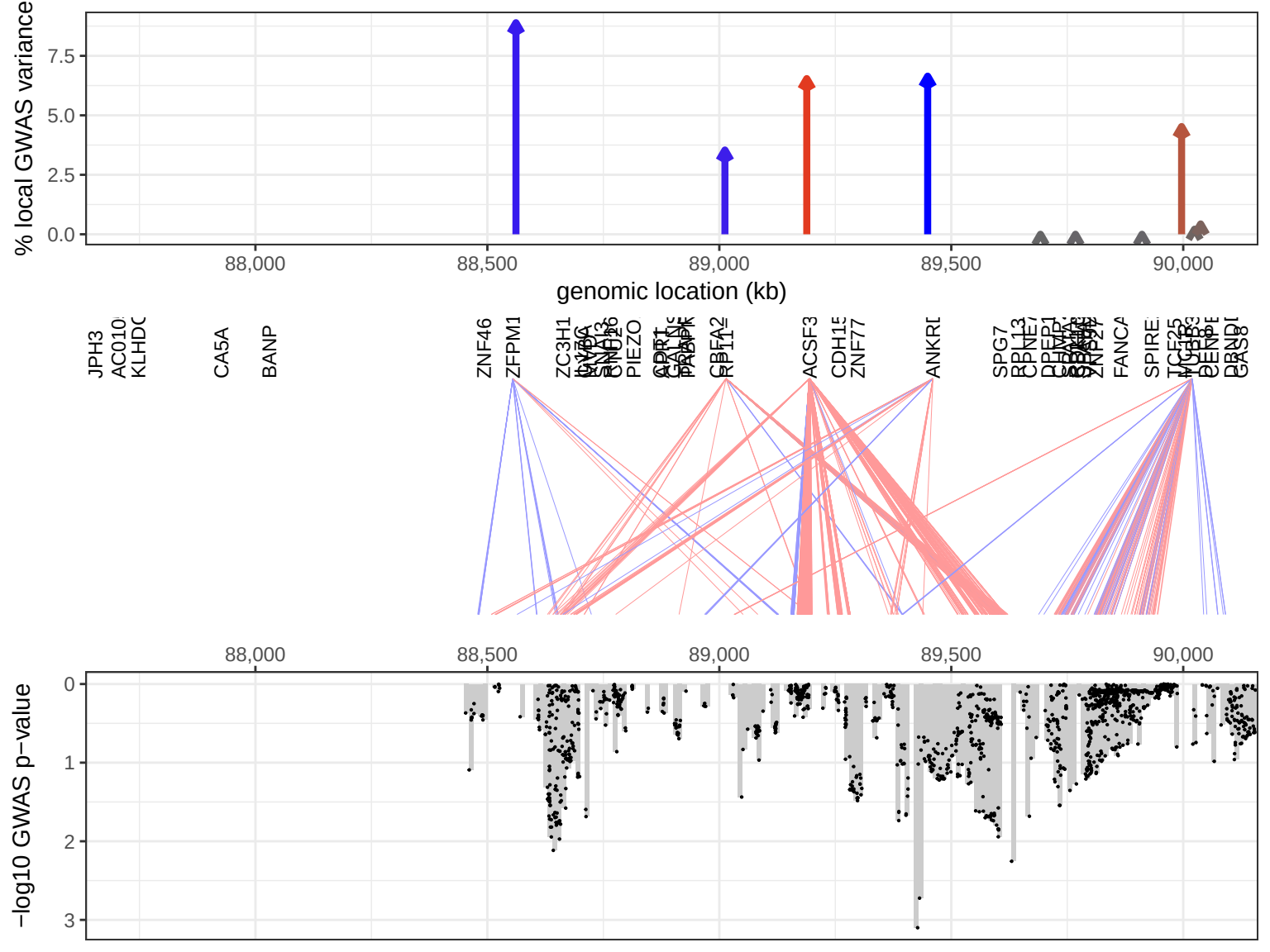
% local GWAS variance
7.5
5.0
2.5
0.0
88,000
88,500
89,000
89,500
90,000
genomic location (kb)
JPH3
CA5A
BANP
ZNF46
ZFPM1
ZC3H1
PIEZO
ACSF3
ANKRD
SPG7
FANCA
SPIRE
−log10 GWAS p−value
0
1
2
3