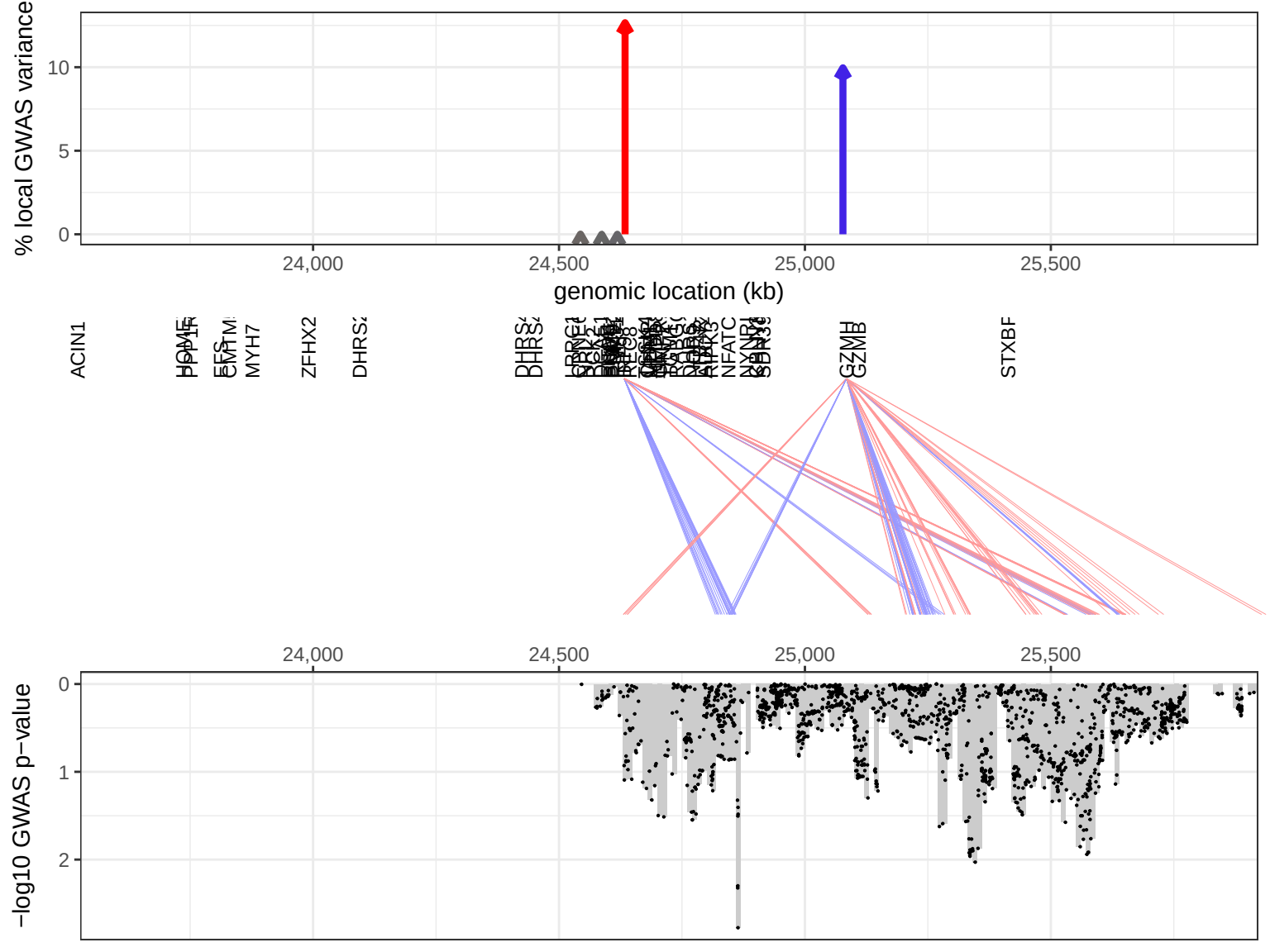
% local GWAS variance
10
5
0
24,000
24,500
25,000
25,500
genomic location (kb)
ACIN1
MYH7
ZFHX2
NFATC
GZMB
STXBF
24,000
24,500
25,000
25,500
−log10 GWAS p−value
0
1
2