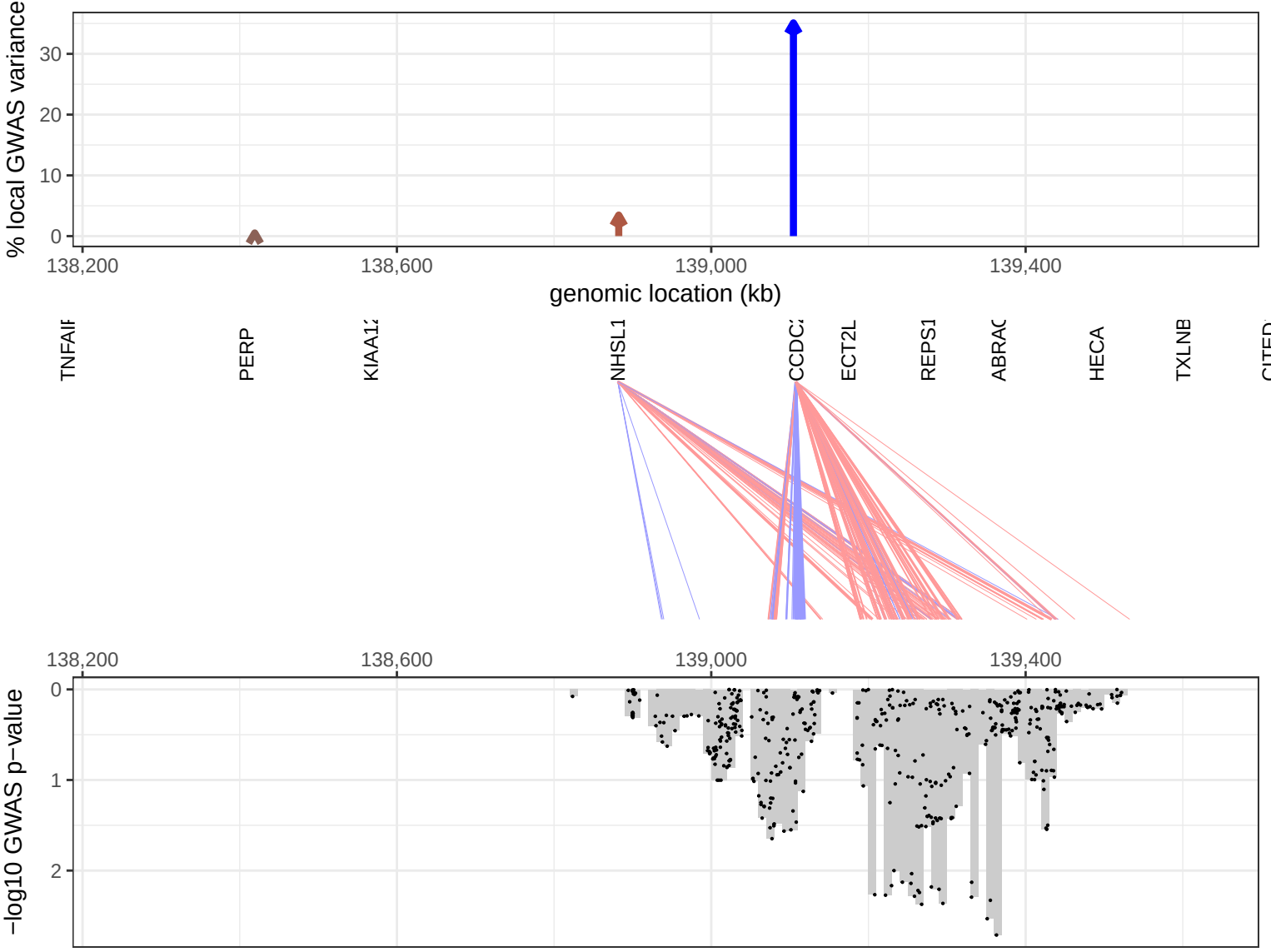
% local GWAS variance
30
20
10
0
138,200
138,600
139,000
139,400
genomic location (kb)
TNFAIP
PERP
KIAA12
NHSL1
CCDC
ECT2L
REPS1
ABRAC
HECA
TXLNB
CITED
138,200
138,600
139,000
139,400
−log10 GWAS p−value
0
1
2