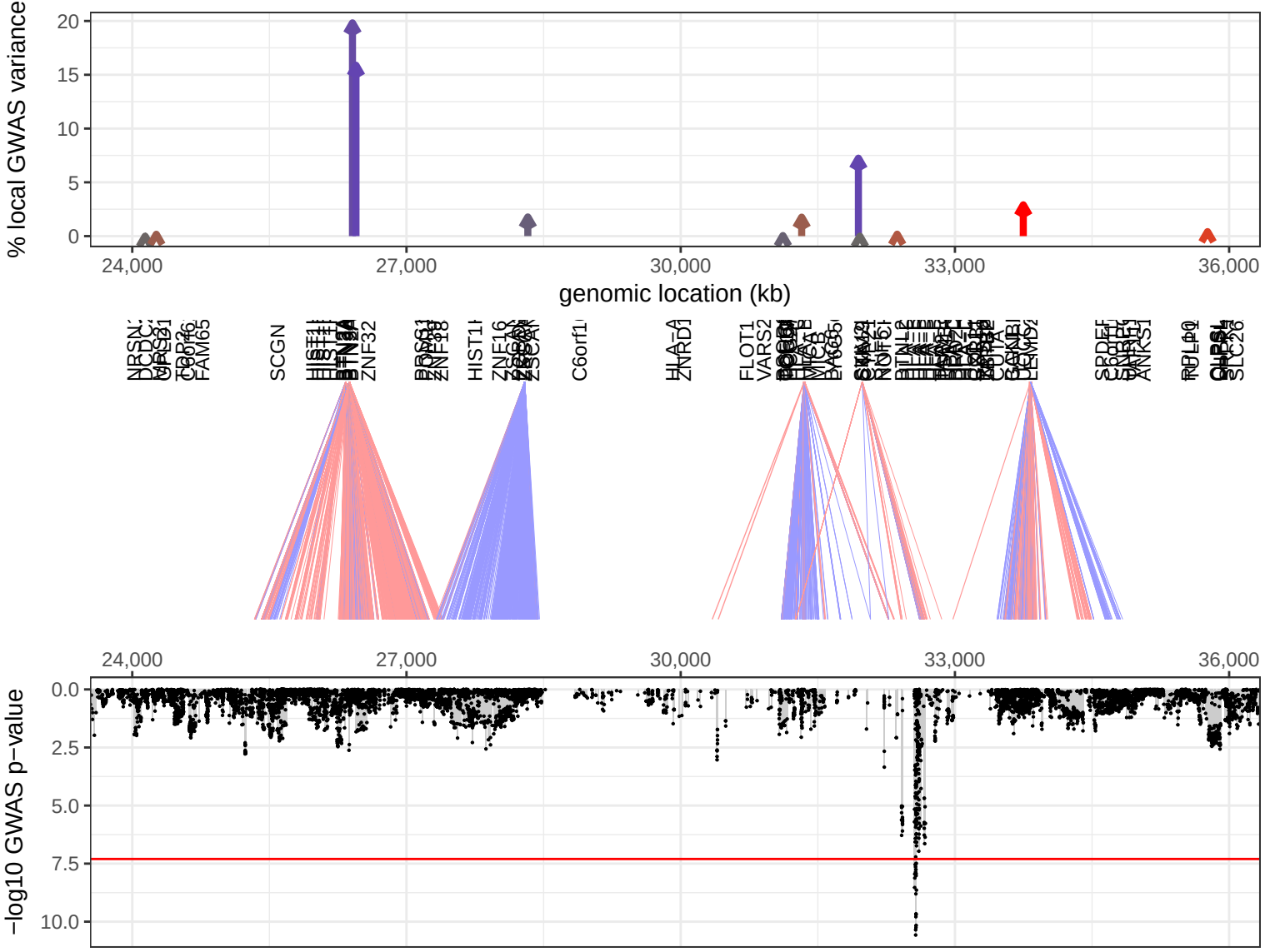
% local GWAS variance
20
15
10
5
0
24,000
27,000
30,000
33,000
36,000
genomic location (kb)
SCGN
ZNF32
HIST1
C6orf1
HLA-A
FLOT1
VARS2
−log10 GWAS p−value
0.0
2.5
5.0
7.5
10.0