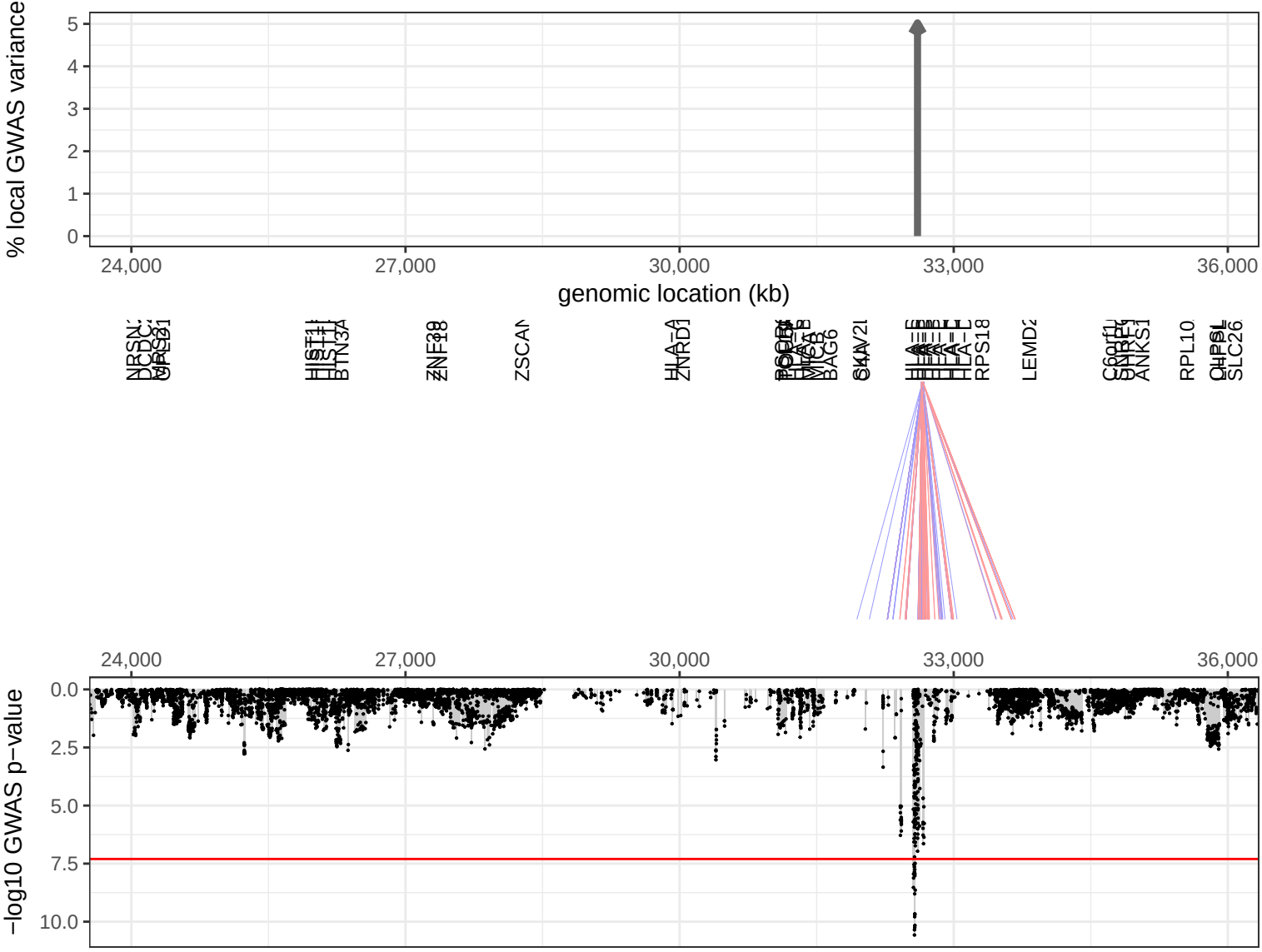
% local GWAS variance
5
4
3
2
1
0
24,000
27,000
30,000
33,000
36,000
genomic location (kb)
ZSCAN
ZNRD1
RPS18
LEMD2
ANKS1
RPL10
SLC26
−log10 GWAS p−value
0.0
2.5
5.0
7.5
10.0