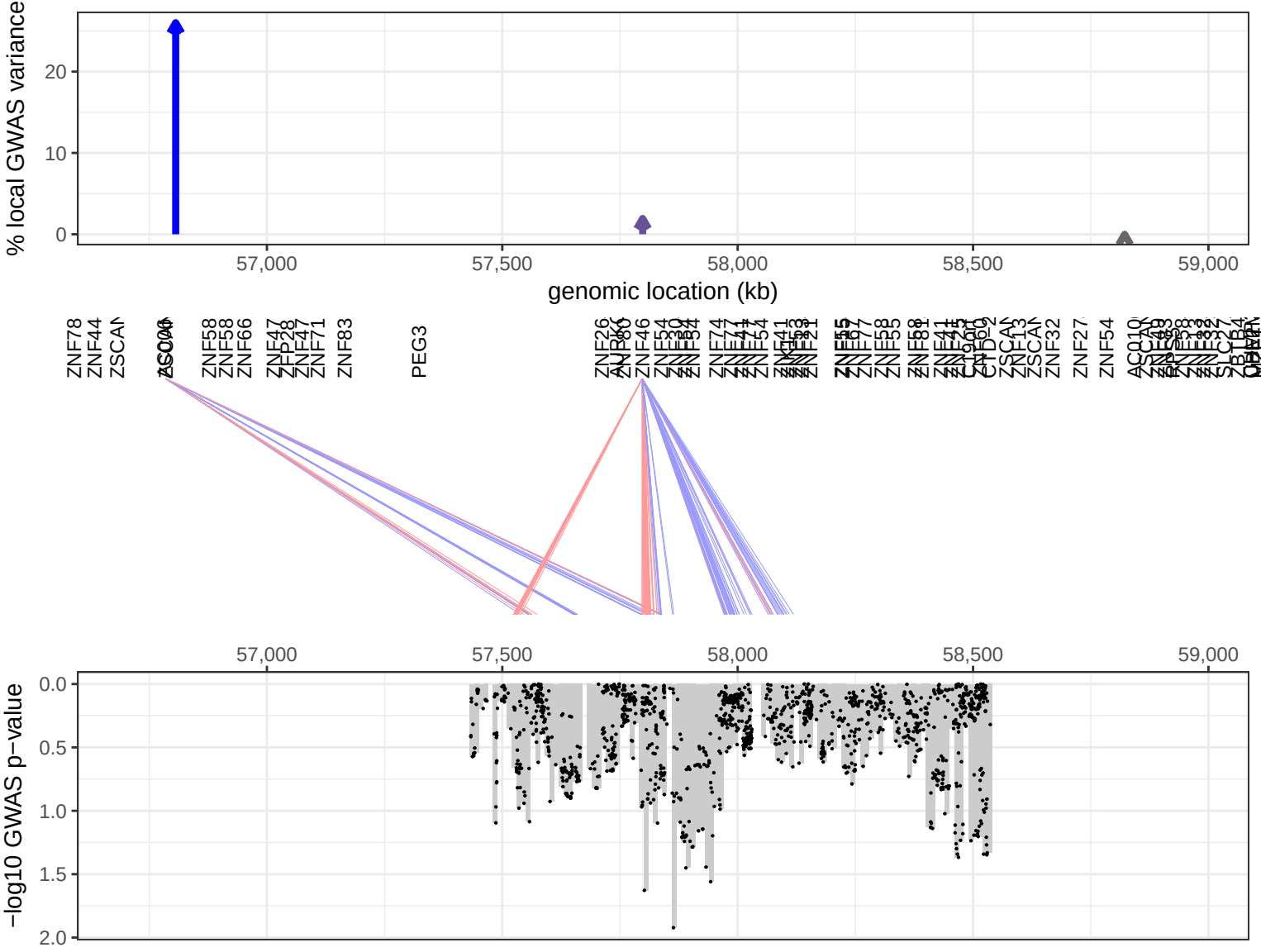
% local GWAS variance
0
10
20
57,000
57,500
58,000
58,500
59,000
genomic location (kb)
ZNF78
ZNF44
ZNF58
ZNF58
ZNF66
ZNF71
ZNF83
PEG3
ZNF46
ZNF32
ZNF27
ZNF54
−log10 GWAS p−value
0.0
0.5
1.0
1.5
2.0