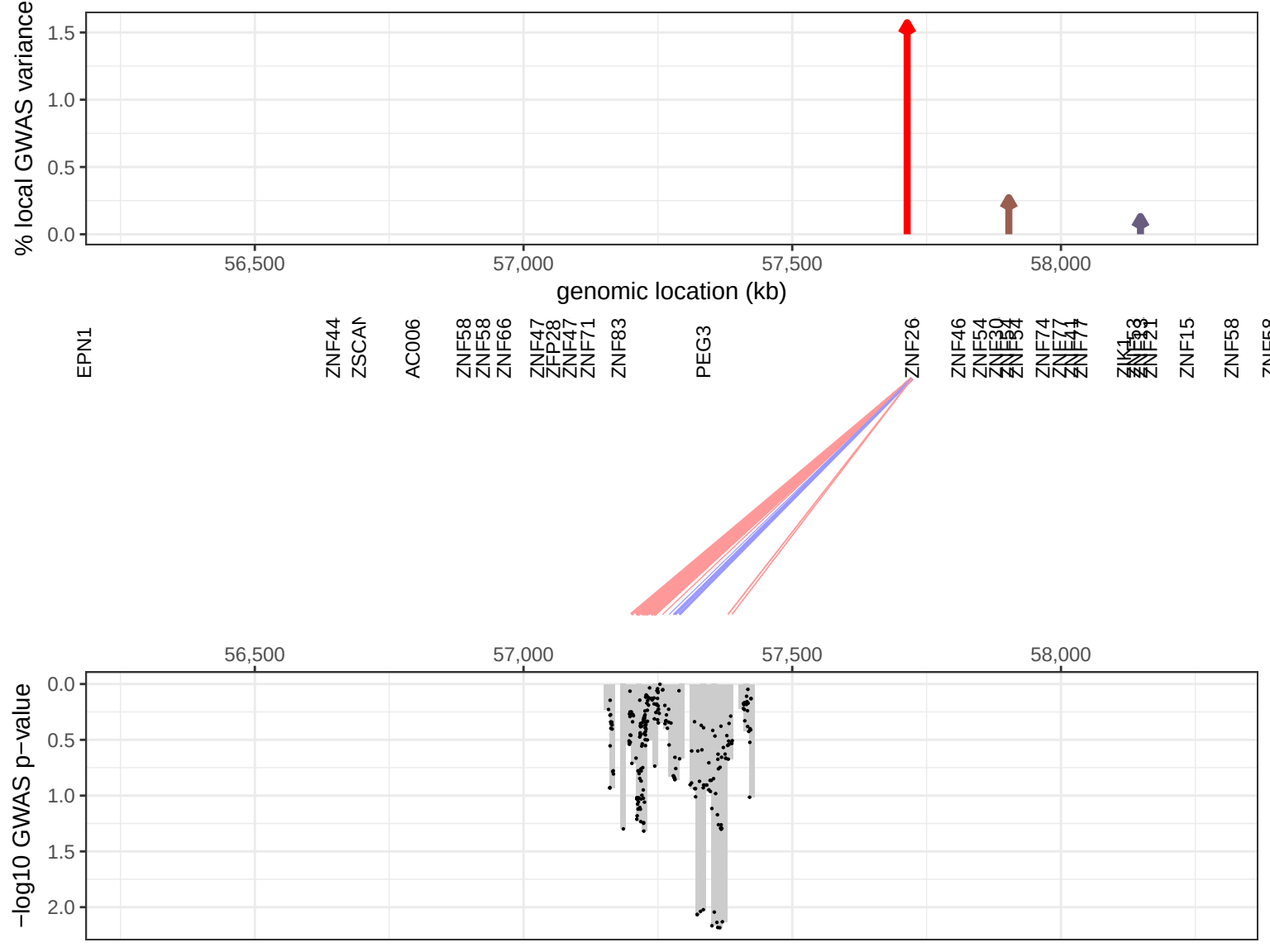
% local GWAS variance
1.5
1.0
0.5
0.0
56,500
57,000
57,500
58,000
genomic location (kb)
EPN1
ZNF44
ZSCAN
AC006
ZNF58
ZNF58
ZNF66
ZNF47
ZFP28
ZNF47
ZNF71
ZNF83
PEG3
ZNF26
ZNF46
ZNF54
ZNF74
ZNF15
ZNF58
56,500
57,000
57,500
58,000
−log10 GWAS p−value
0.0
0.5
1.0
1.5
2.0