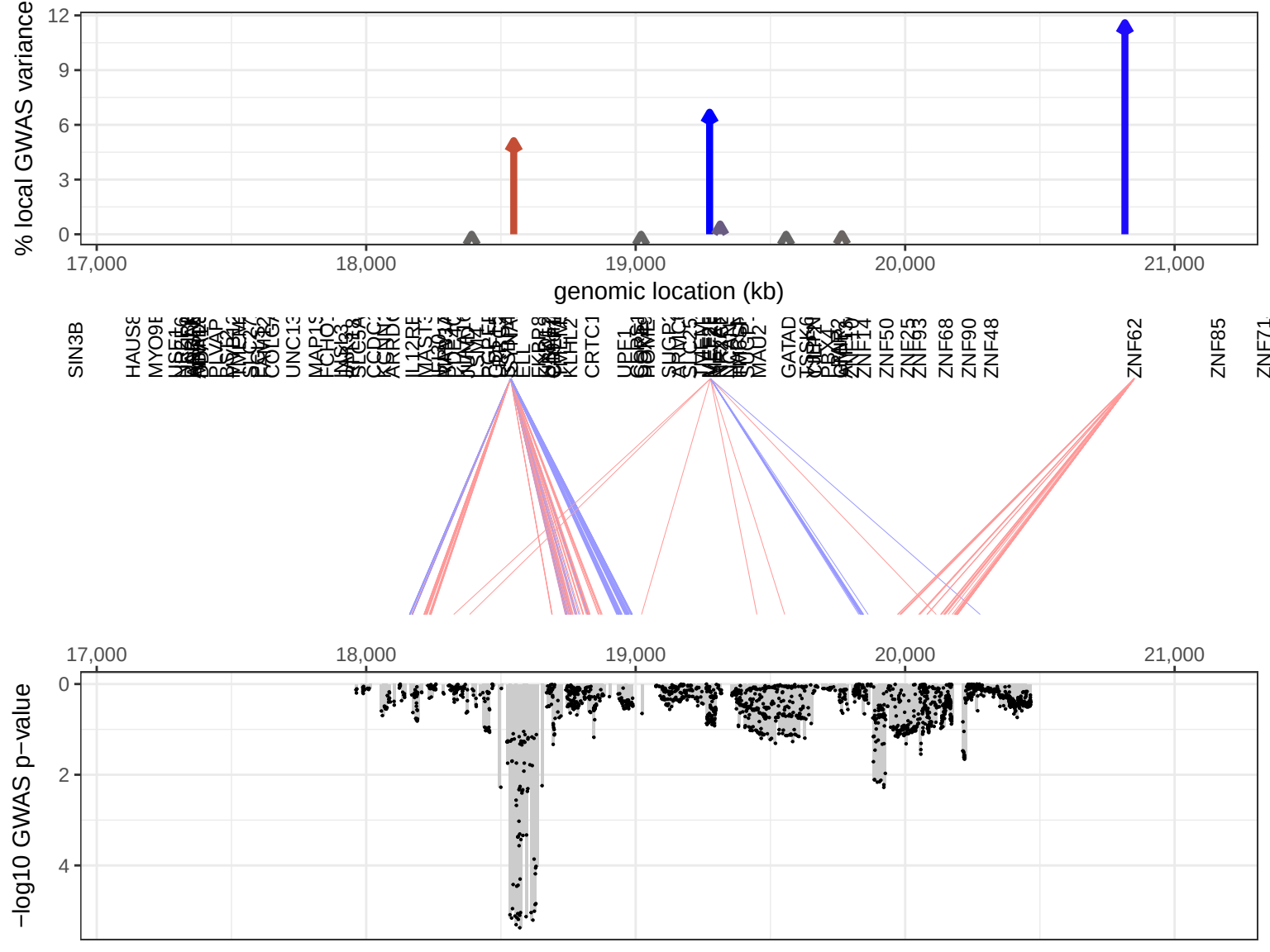
% local GWAS variance
12
9
6
3
0
17,000
18,000
19,000
20,000
21,000
genomic location (kb)
SIN3B
CRTC1
GATAD
ZNF50
ZNF25
ZNF93
ZNF68
ZNF90
ZNF48
ZNF62
ZNF85
−log10 GWAS p−value
0
2
4