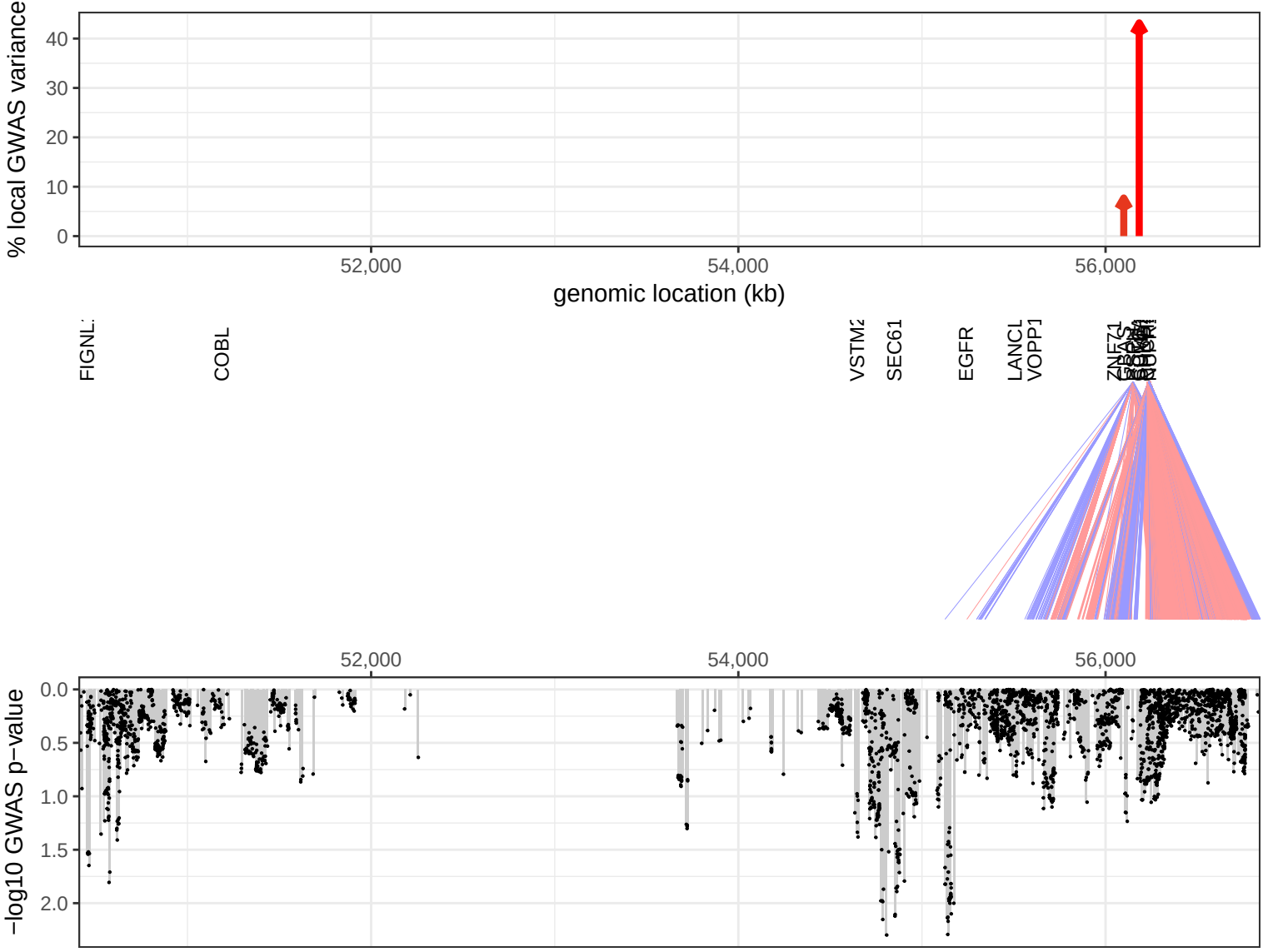
% local GWAS variance
40
30
20
10
0
52,000
54,000
56,000
genomic location (kb)
FIGNL
COBL
VSTM2
SEC61
EGFR
LANCL
VOPP1
ZNF71
52,000
54,000
56,000
−log10 GWAS p−value
0.0
0.5
1.0
1.5
2.0