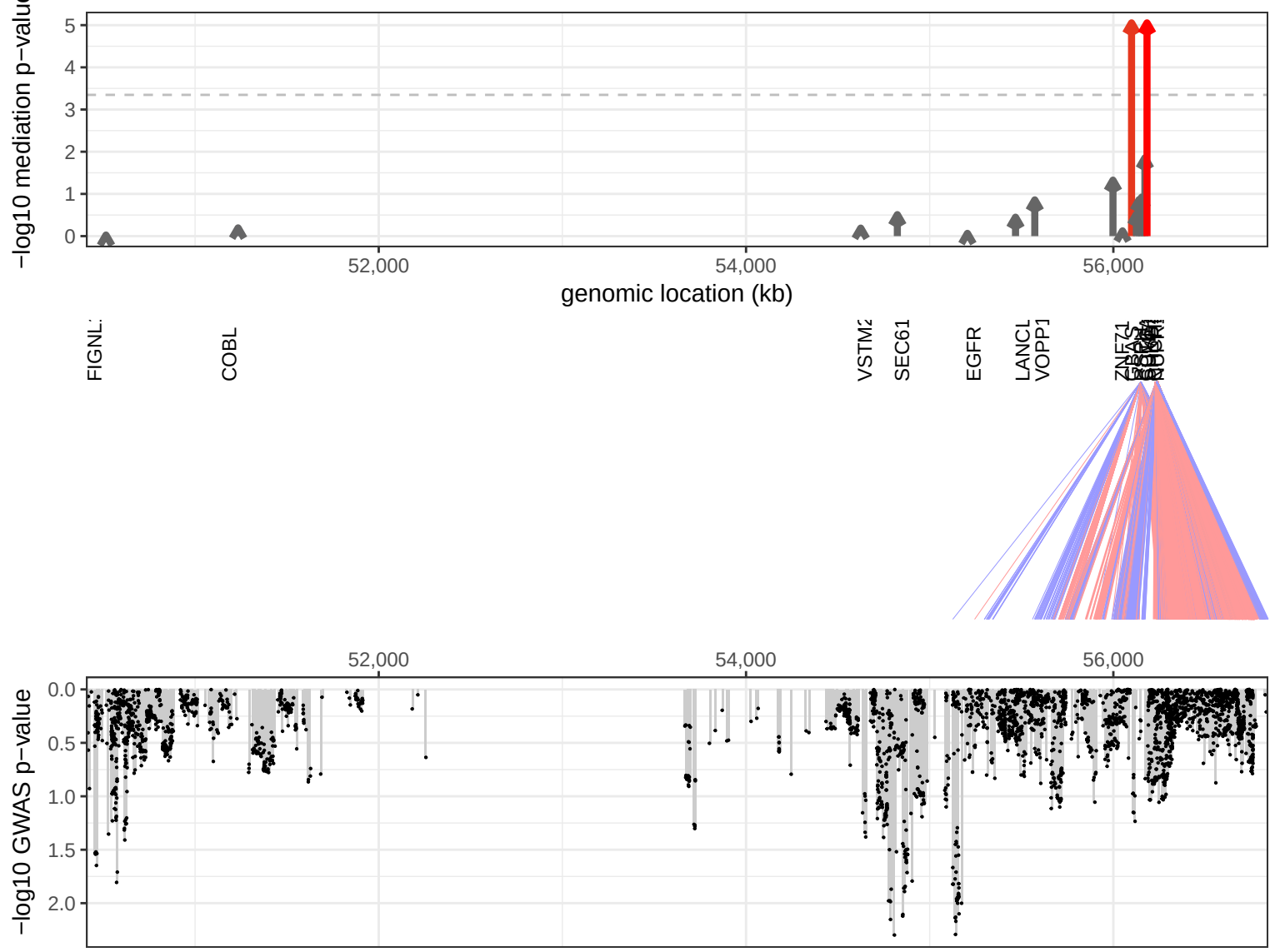
−log10 mediation p−value
5
4
3
2
1
0
52,000
54,000
56,000
genomic location (kb)
FIGNL1
COBL
VSTM2
SEC61
EGFR
LANCL
VOPP1
ZNF71
−log10 GWAS p−value
0.0
0.5
1.0
1.5
2.0