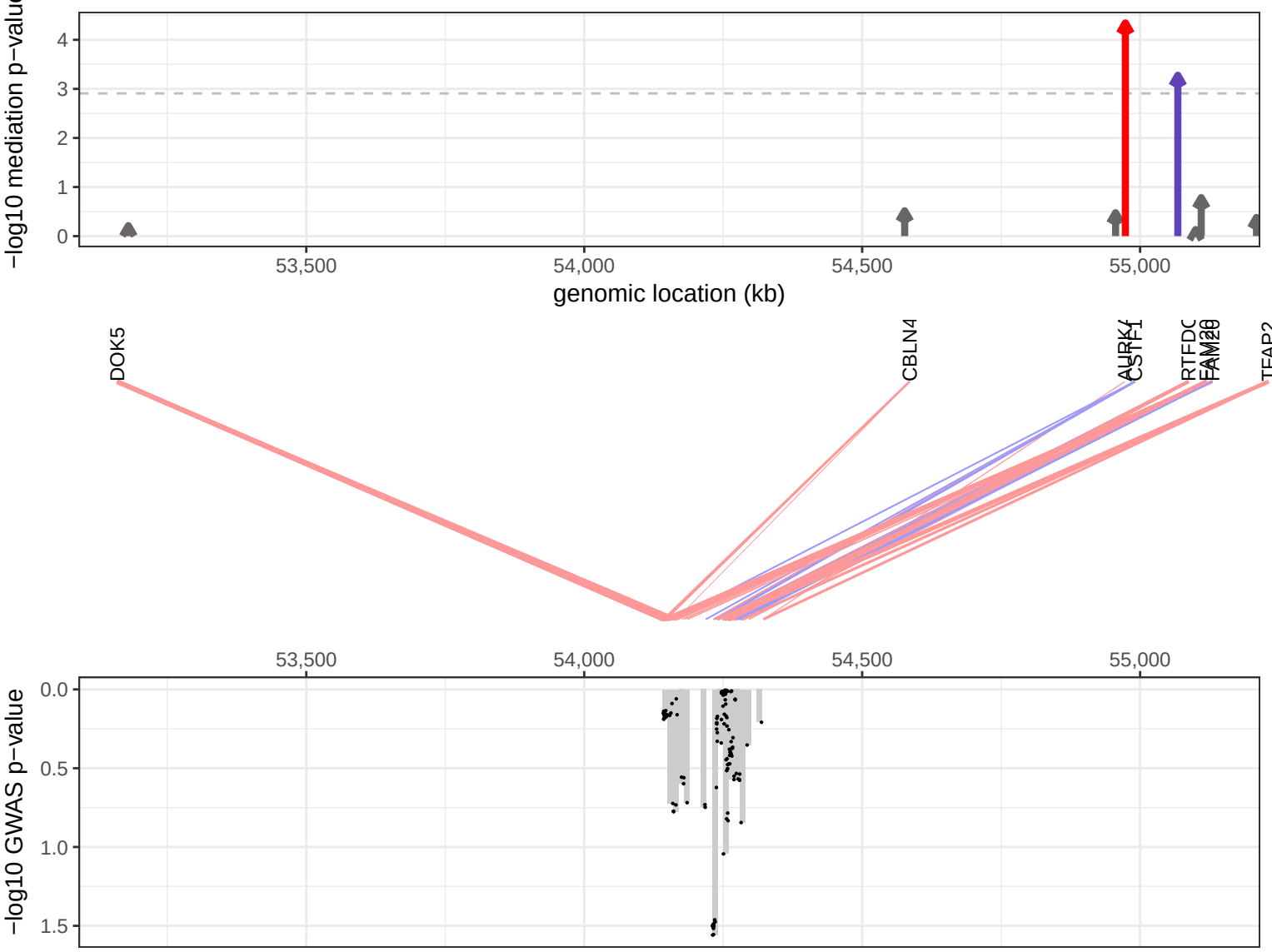
−log10 mediation p−value
4
3
2
1
0
53,500
54,000
54,500
55,000
genomic location (kb)
DOK5
CBLN4
AURKA
CSTF1
RTFDC
FAM20
TEAD2
53,500
54,000
54,500
55,000
−log10 GWAS p−value
0.0
0.5
1.0
1.5