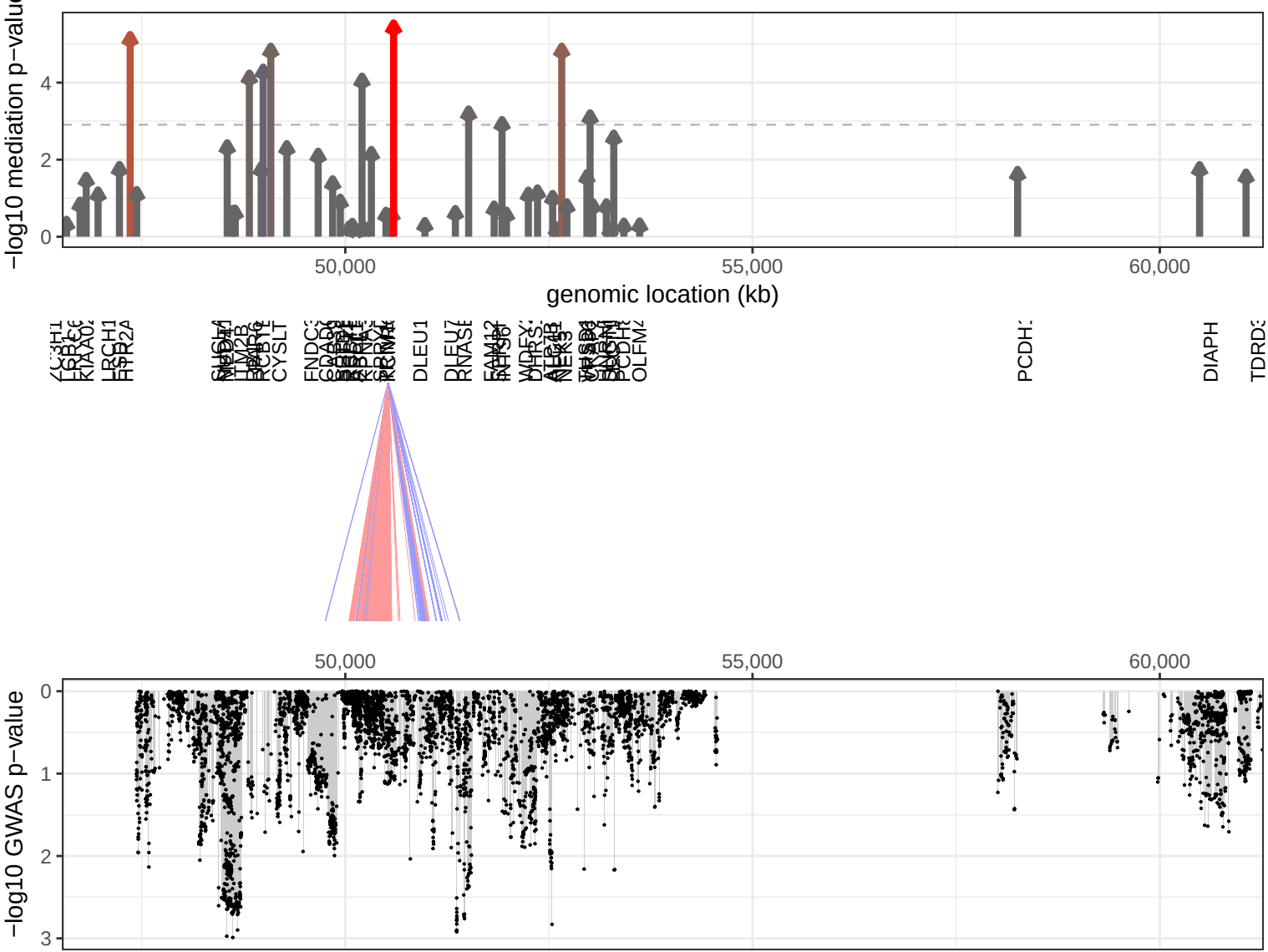
−log10 mediation p−value
0
2
4
50,000
55,000
60,000
genomic location (kb)
CYSLT
DLEU1
PCDH
DIAPH
TDRD
−log10 GWAS p−value
0
1
2
3
50,000
55,000
60,000