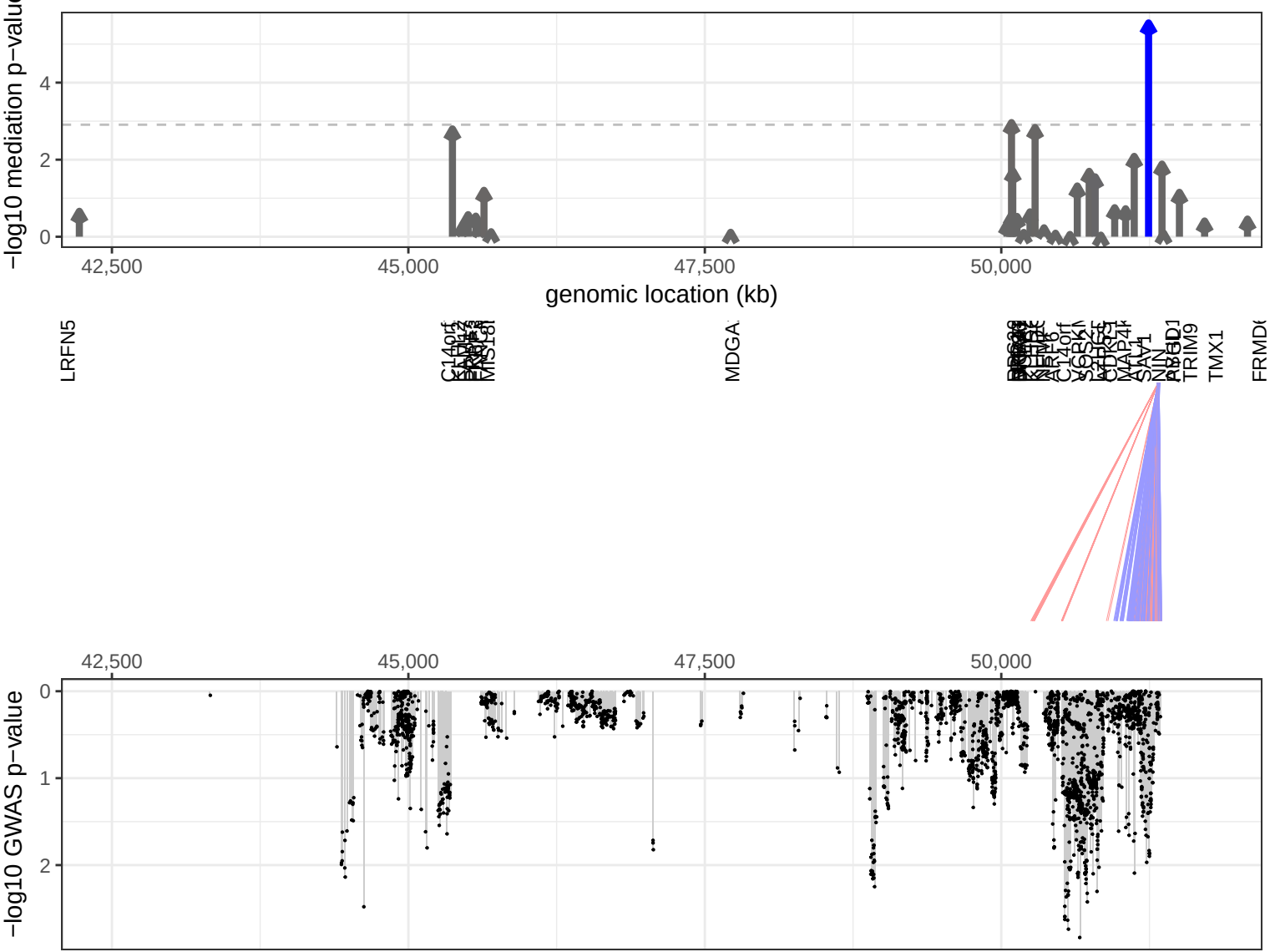
−log10 mediation p−value
0
2
4
42,500
45,000
47,500
50,000
genomic location (kb)
LRFN5
MDGA
TRIM9
TMX1
FRMD
−log10 GWAS p−value
0
1
2