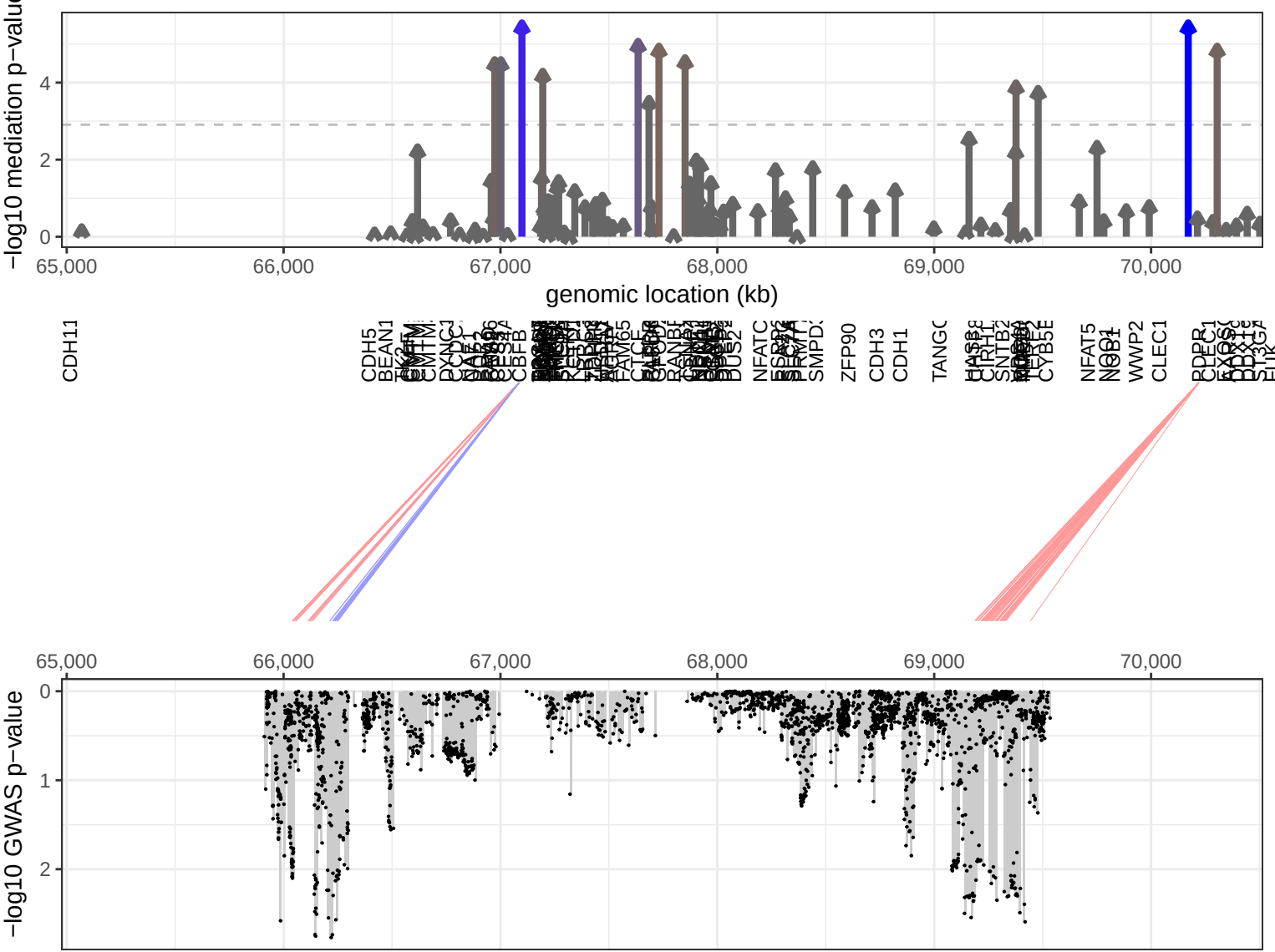
−log10 mediation p−value
0
2
4
65,000
66,000
67,000
68,000
69,000
70,000
genomic location (kb)
CDH11
CDH5
BEAN1
NFATC
ZFP90
CDH3
CDH1
TANGO
NFAT5
WWP2
CLEC1
−log10 GWAS p−value
0
1
2