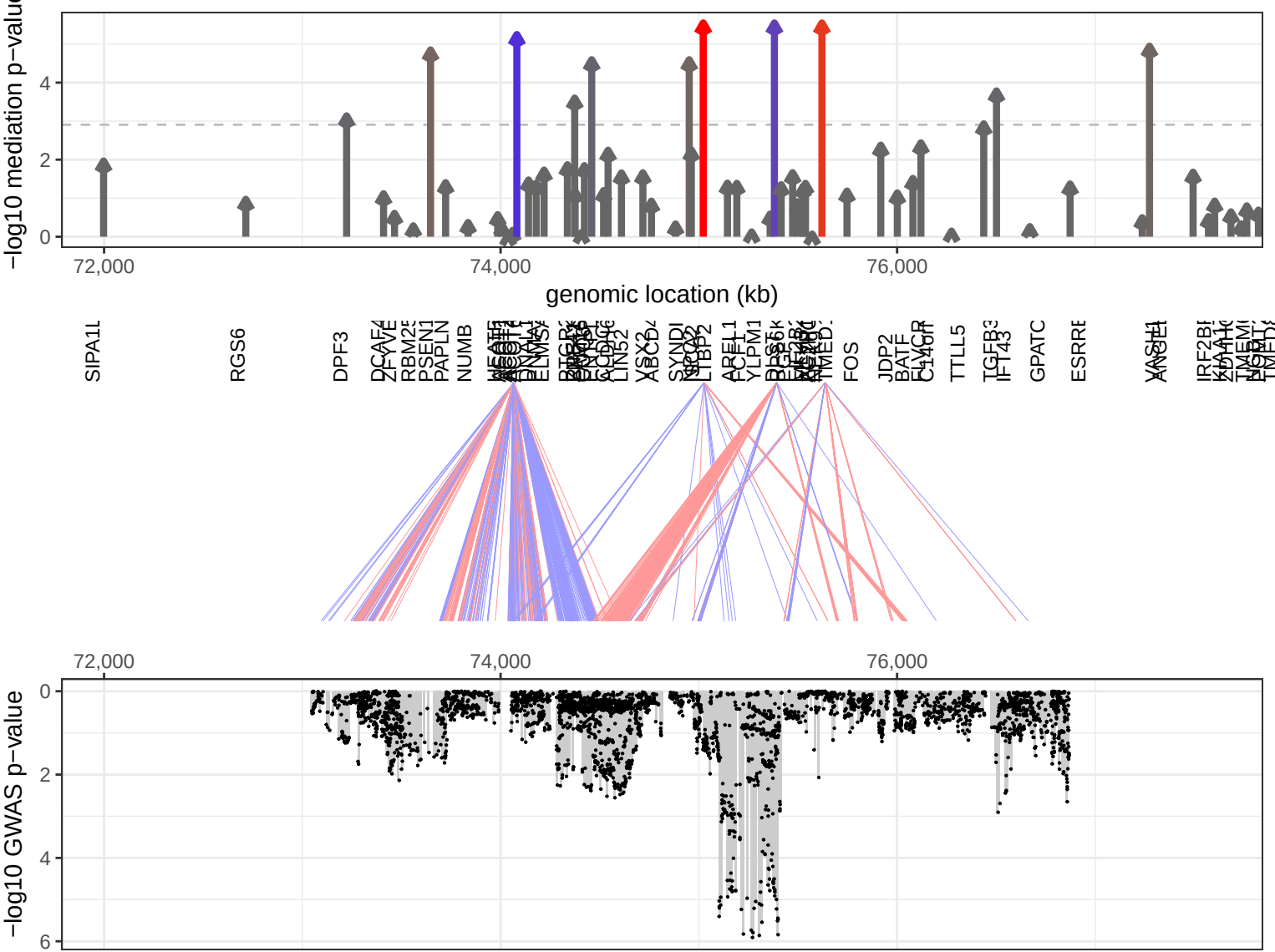
−log10 mediation p−value
0
2
4
72,000
74,000
76,000
genomic location (kb)
SIPA1L
RGS6
DPF3
NUMB
LIN52
LTBP2
YLPM1
FOS
JDP2
BATF
TTLL5
GPATC
ESRRE
−log10 GWAS p−value
0
2
4
6
72,000
74,000
76,000