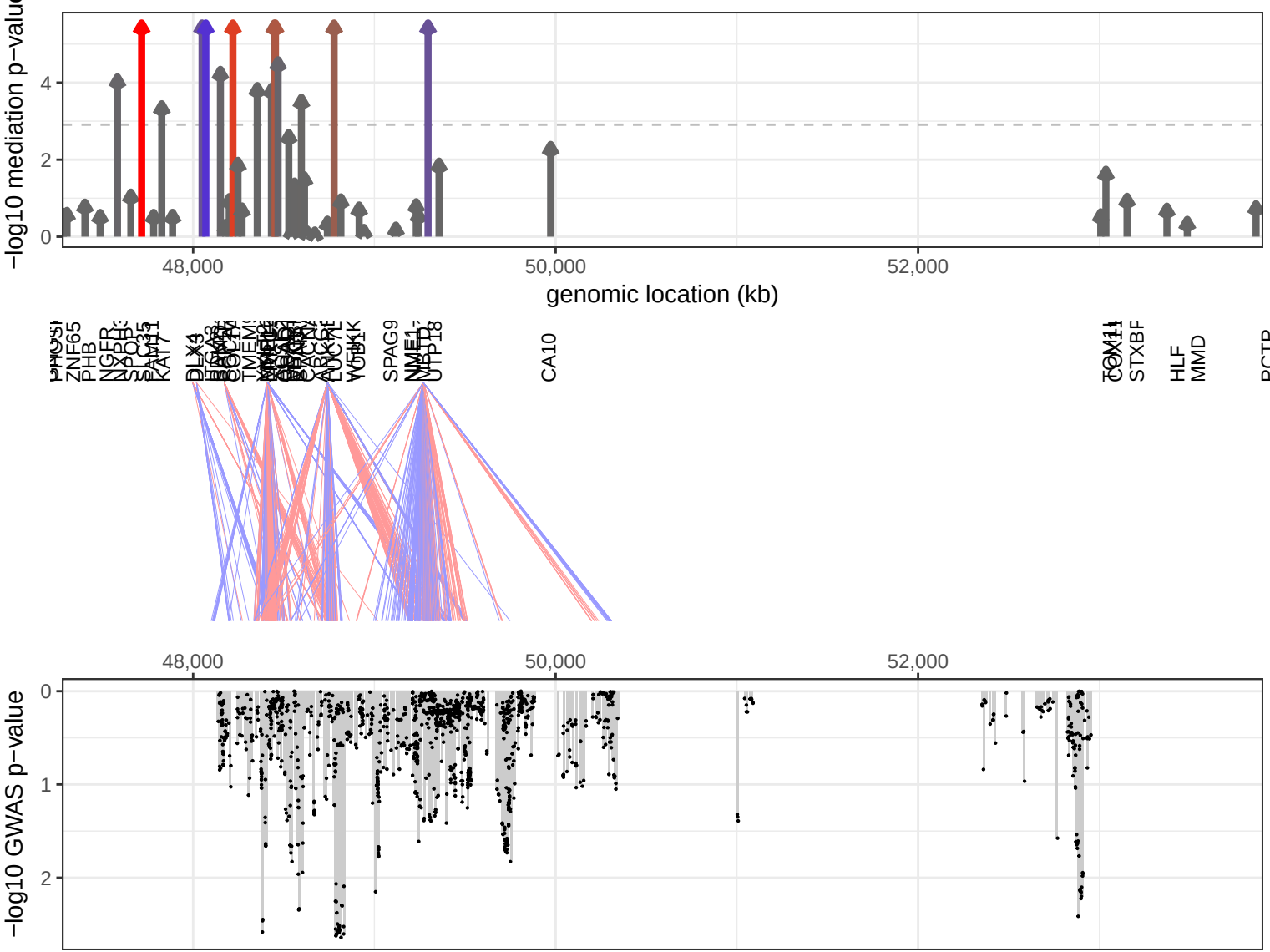
−log10 mediation p−value
0
2
4
48,000
50,000
52,000
genomic location (kb)
ZNF65
PHB
NGFR
KAT7
DLX3
SPAG9
CA10
STXBF
HLF
MMD
PCTP
48,000
50,000
52,000
−log10 GWAS p−value
0
1
2