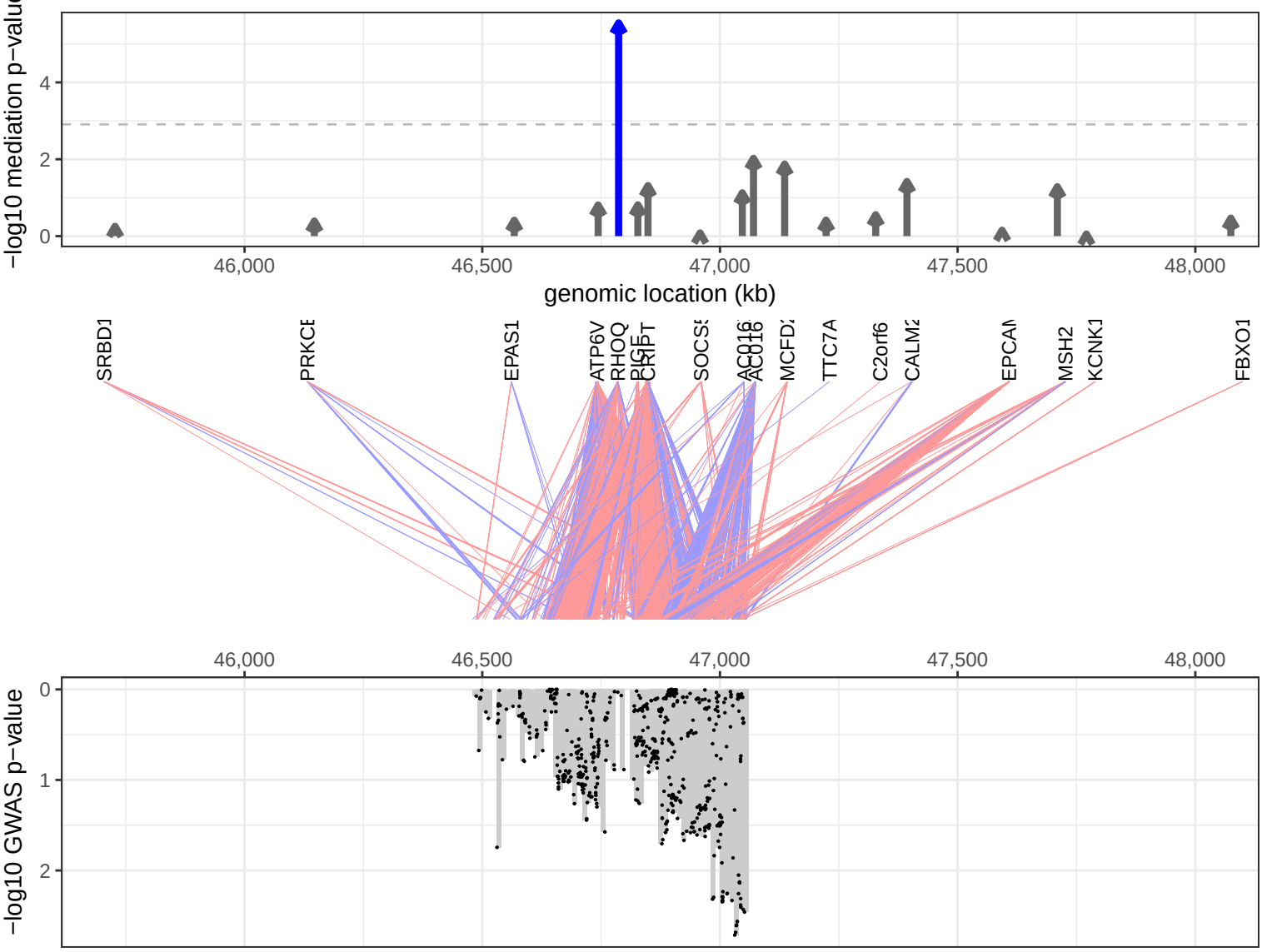
−log10 mediation p−value
0
2
4
46,000
46,500
47,000
47,500
48,000
genomic location (kb)
SRBD1
PRKCE
EPAS1
ATP6V
RHOQ
CRIPT
SOCS
MCFD
TTC7A
C2orf6
CALM
EPCAM
MSH2
KCNK1
FBXO1
46,000
46,500
47,000
47,500
48,000
−log10 GWAS p−value
0
1
2