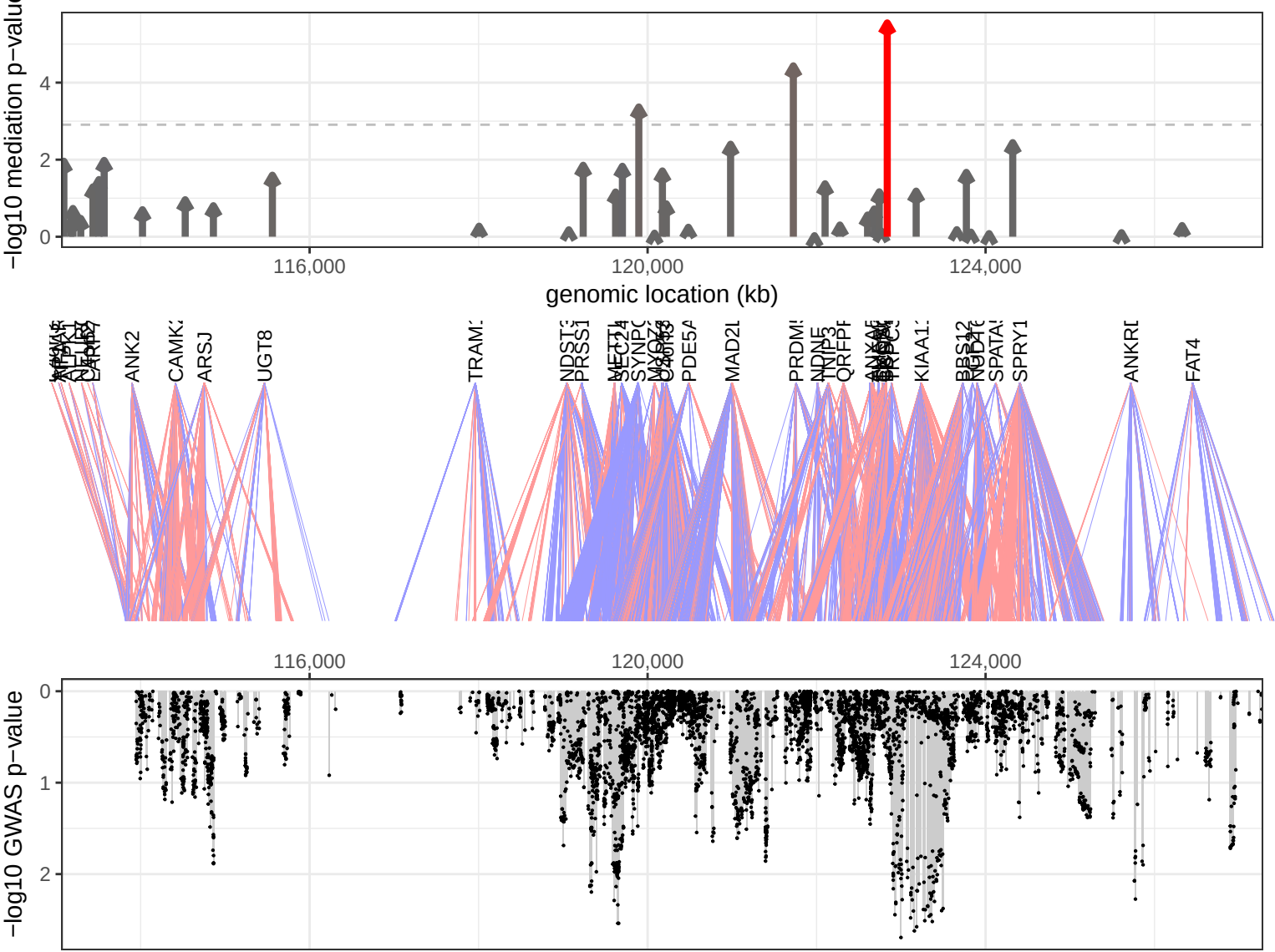
−log10 mediation p−value
0
2
4
116,000
120,000
124,000
genomic location (kb)
ANK2
CAMK2
ARSJ
UGT8
TRAM2
NDST3
PRSS1
PDE5A
MAD2L
PRDM1
KIAA1
SPATA1
SPRY1
ANKRD
FAT4
116,000
120,000
124,000
−log10 GWAS p−value
0
1
2