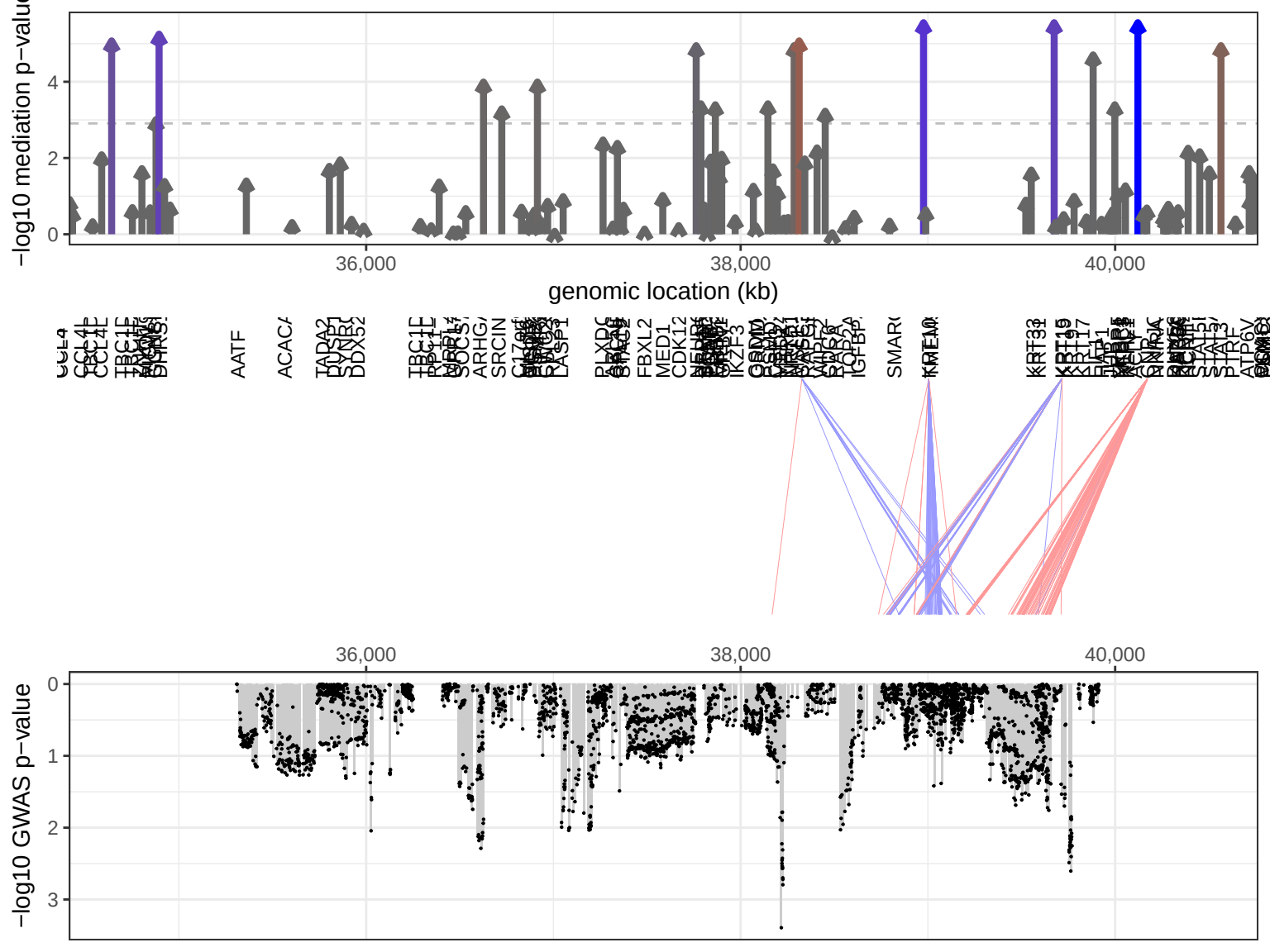
−log10 mediation p−value
0
2
4
36,000
38,000
40,000
genomic location (kb)
AATF
ACAC
TADA2
DDX52
ARHG
SRCIN
FBXL2
MED1
CDK12
SMAR
KRT33
−log10 GWAS p−value
0
1
2
3
36,000
38,000
40,000