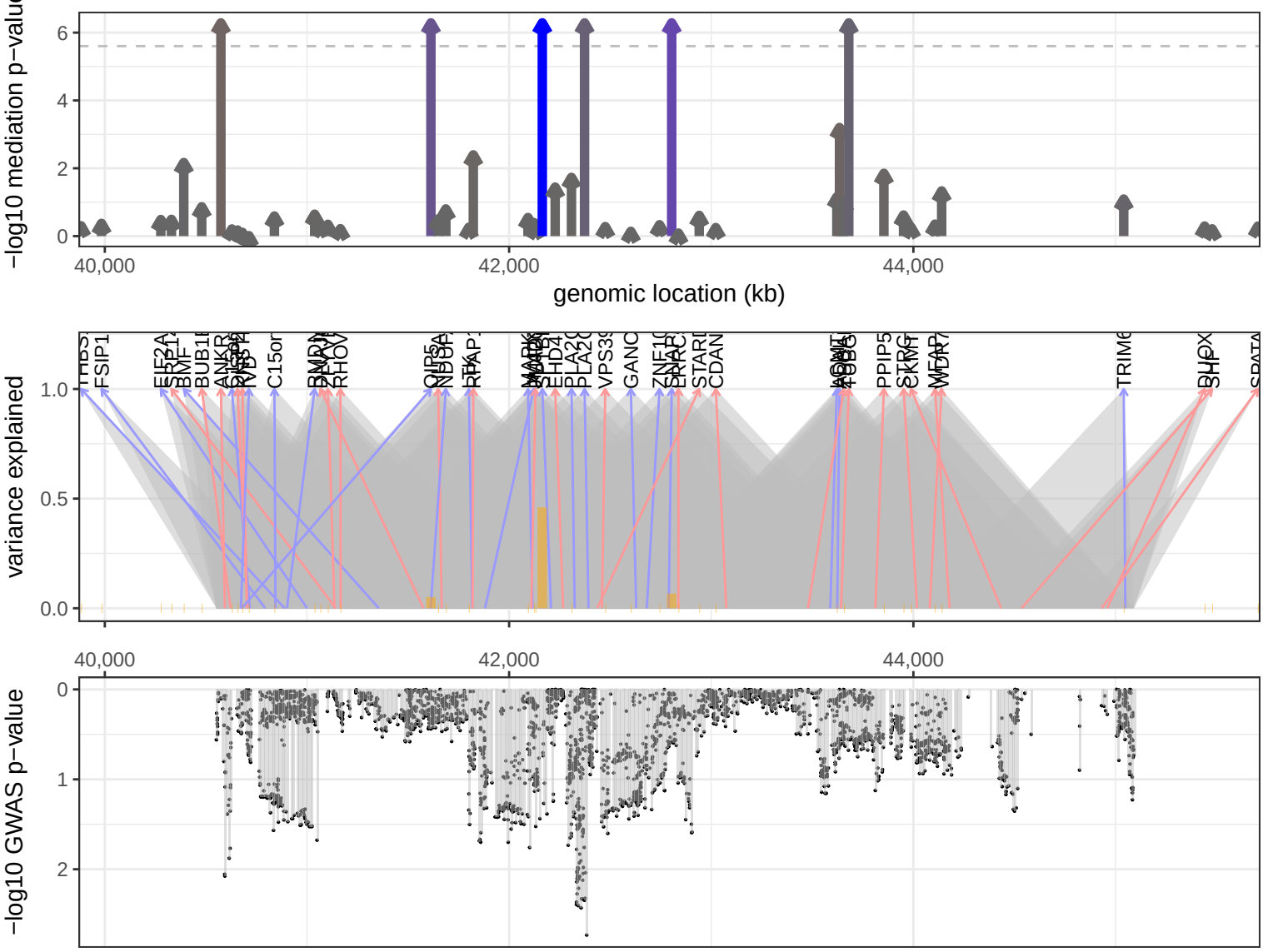
−log10 mediation p−value
6
4
2
0
40,000
42,000
44,000
genomic location (kb)
variance explained
1.0
0.5
0.0
FSIP1
C15orf
RHOV
GANC
STARD
CDAN
PPIP5
TRIM6
40,000
42,000
44,000
−log10 GWAS p−value
0
1
2