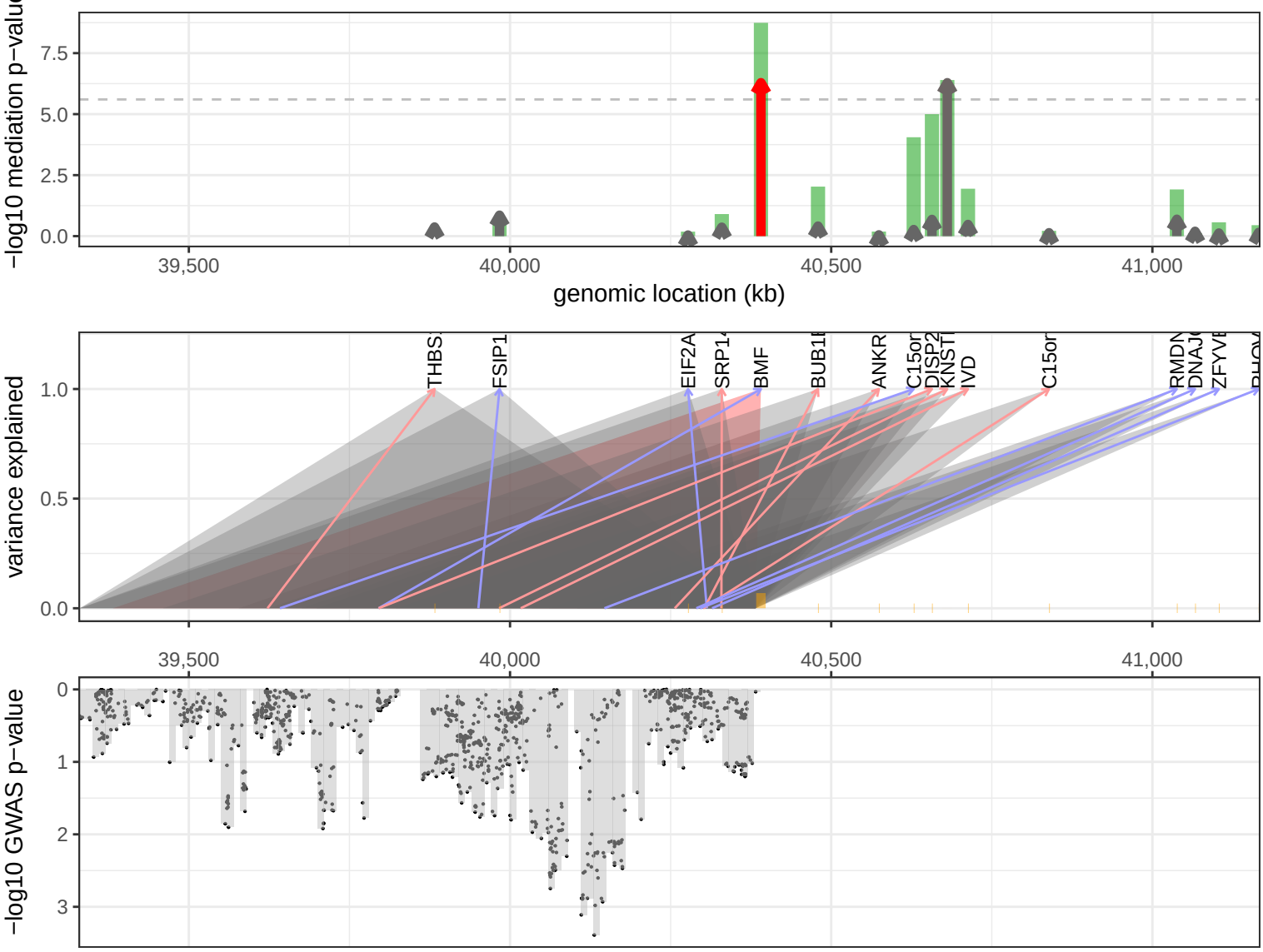
−log10 mediation p−value
7.5
5.0
2.5
0.0
39,500
40,000
40,500
41,000
genomic location (kb)
variance explained
1.0
0.5
0.0
THBS
FSIP1
EIF2A
SRP14
BMF
BUB1B
ANKR
C15or
DISP2
KNST
IVD
C15or
RMDN
DNAJ
ZFYVE
39,500
40,000
40,500
41,000
−log10 GWAS p−value
0
1
2
3