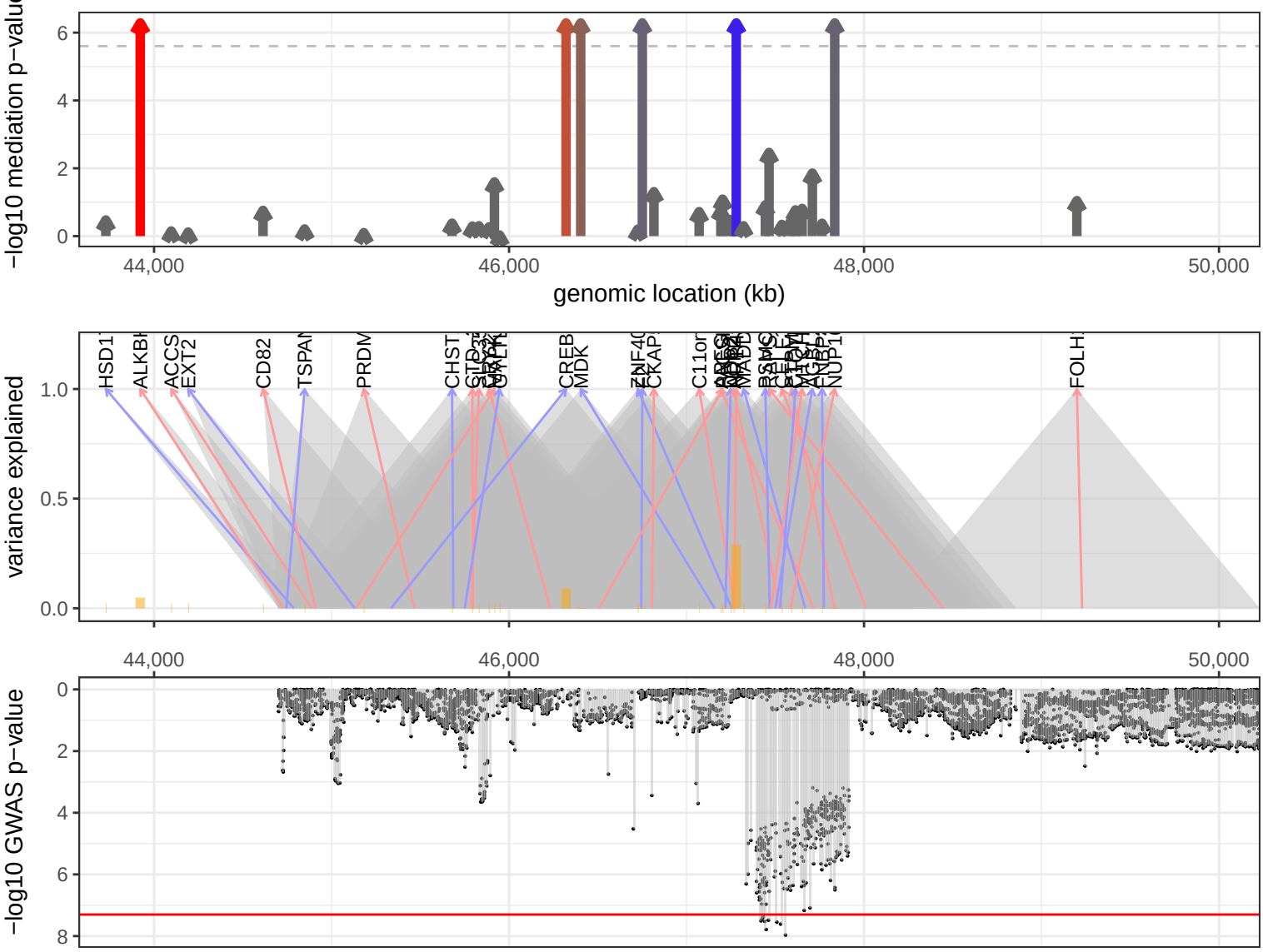
−log10 mediation p−value
6
4
2
0
44,000
46,000
48,000
50,000
genomic location (kb)
variance explained
1.0
0.5
0.0
HSD1
ALKBH
ACCS
EXT2
CD82
TSPAN
PRDM
CHST
CREB
MDK
ZNF40
CKAP
C11or
FOLH
NUP1
−log10 GWAS p−value
0
2
4
6
8