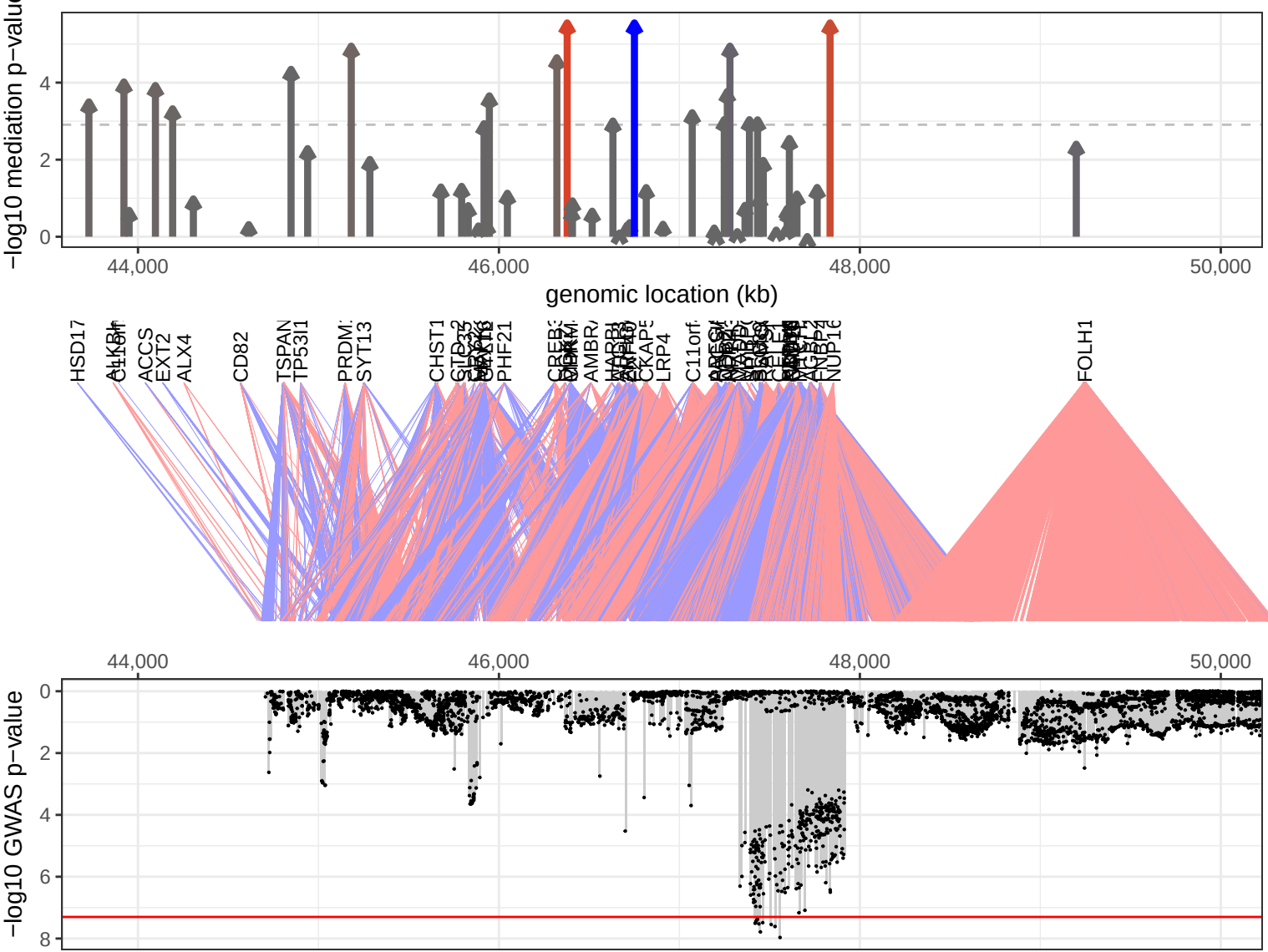
−log10 mediation p−value
0
2
4
44,000
46,000
48,000
50,000
genomic location (kb)
HSD17
ACCS
EXT2
ALX4
CD82
TSPAN
TP53I1
PRDM
SYT13
CHST1
PHF21
AMBR
LRP4
C11orf
FOLH1
−log10 GWAS p−value
0
2
4
6
8