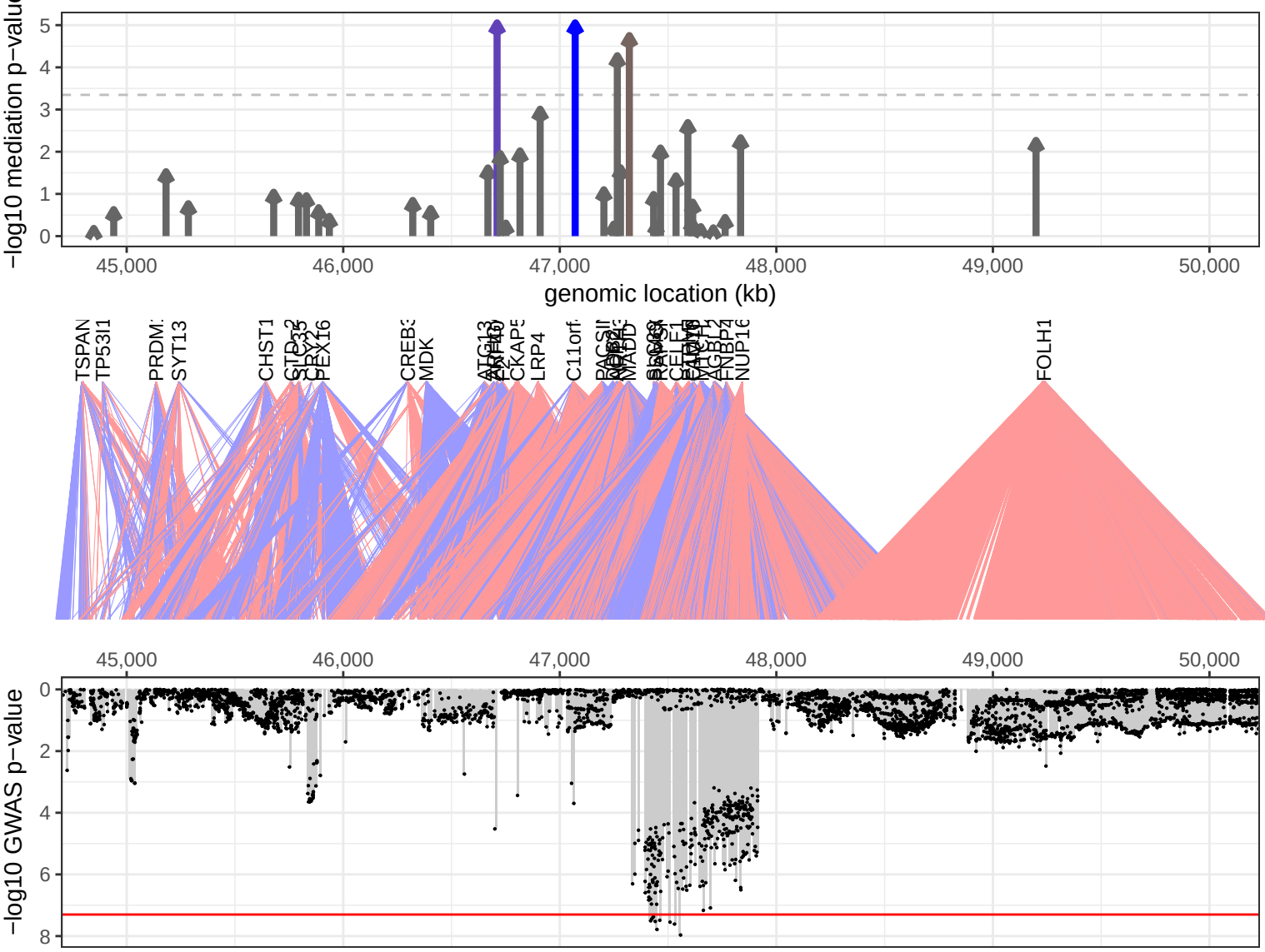
−log10 mediation p−value
genomic location (kb)
45,000
46,000
47,000
48,000
49,000
50,000
TSPAN
TP53I1
PRDM
SYT13
CHST1
PEX16
CREB
MDK
CKAP
LRP4
C11orf
MADD
NUP16
FOLH1
−log10 GWAS p−value