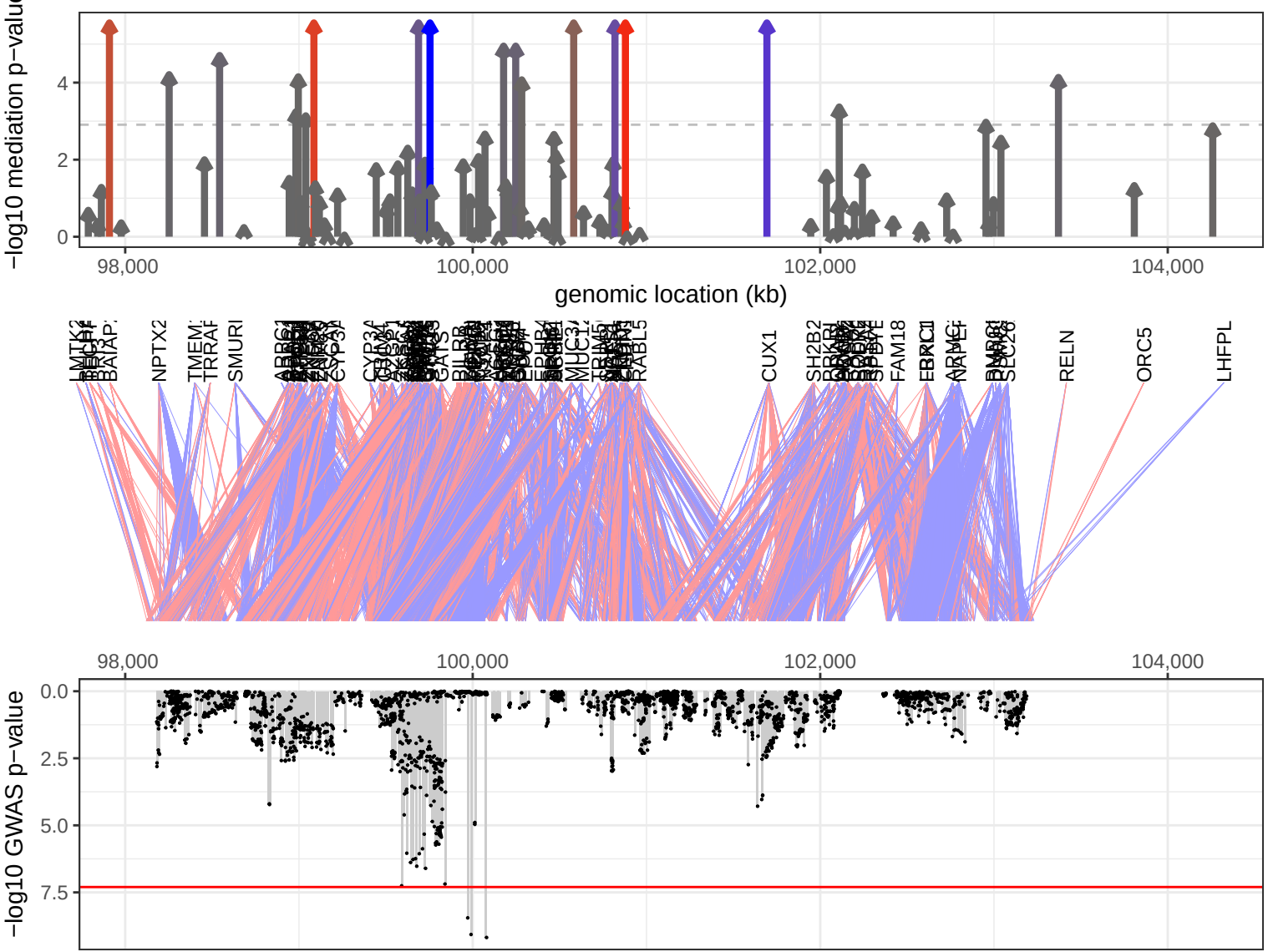
−log10 mediation p−value
0
2
4
98,000
100,000
102,000
104,000
genomic location (kb)
NPTX2
TMEM
TRRAF
SMURI
CYP3
CATS
RABL5
CUX1
SH2B2
FAM18
RELN
ORC5
LHFPL
98,000
100,000
102,000
104,000
−log10 GWAS p−value
0.0
2.5
5.0
7.5