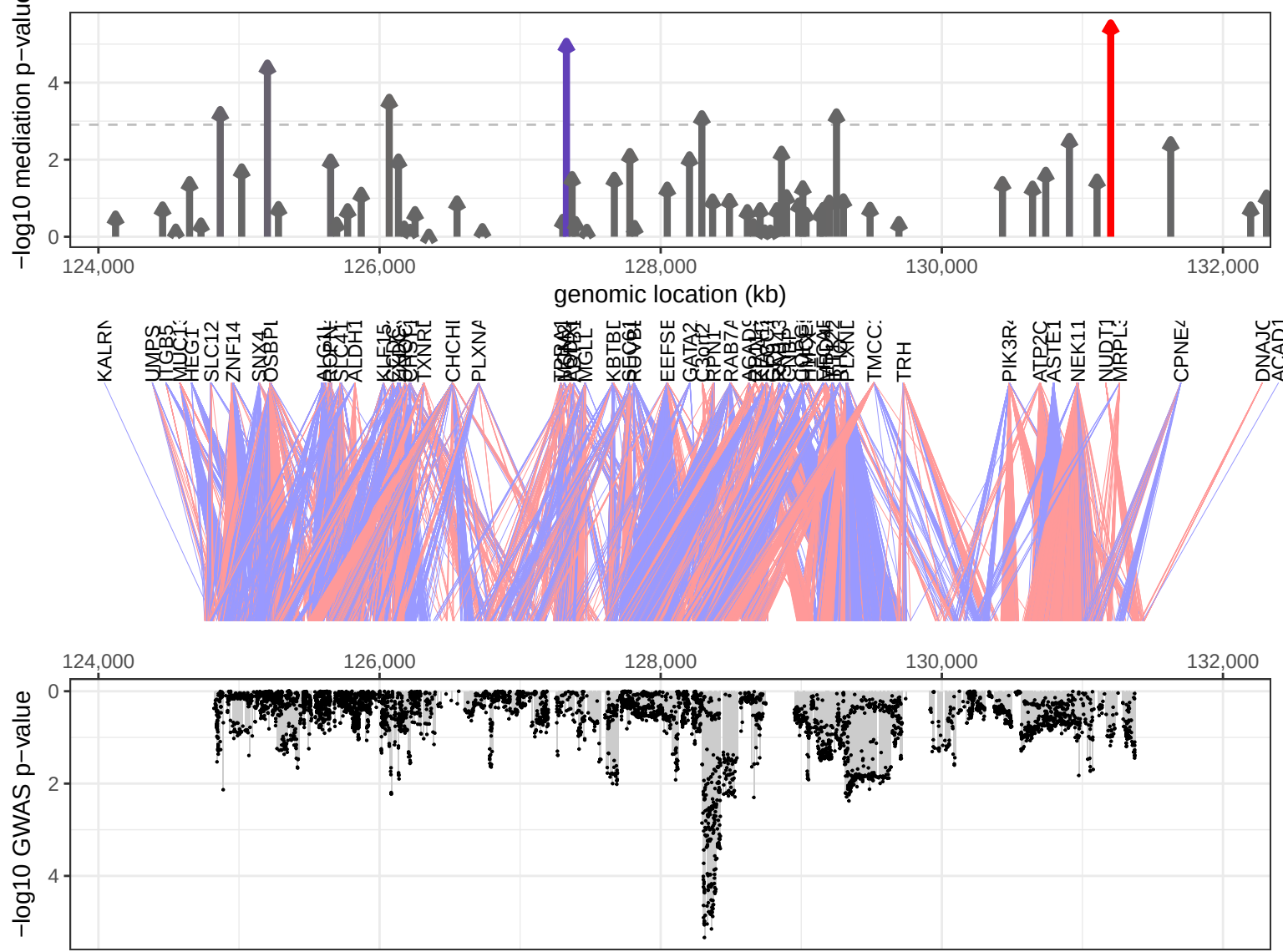
−log10 mediation p−value
0
2
4
124,000
126,000
128,000
130,000
132,000
genomic location (kb)
KALRN
UMPS
SLC12
ZNF14
CHCHI
PLXNA
KBTBD
EEFSE
TMCC
TRH
PIK3R
ASTE1
NEK11
CPNE
−log10 GWAS p−value
0
2
4
124,000
126,000
128,000
130,000
132,000