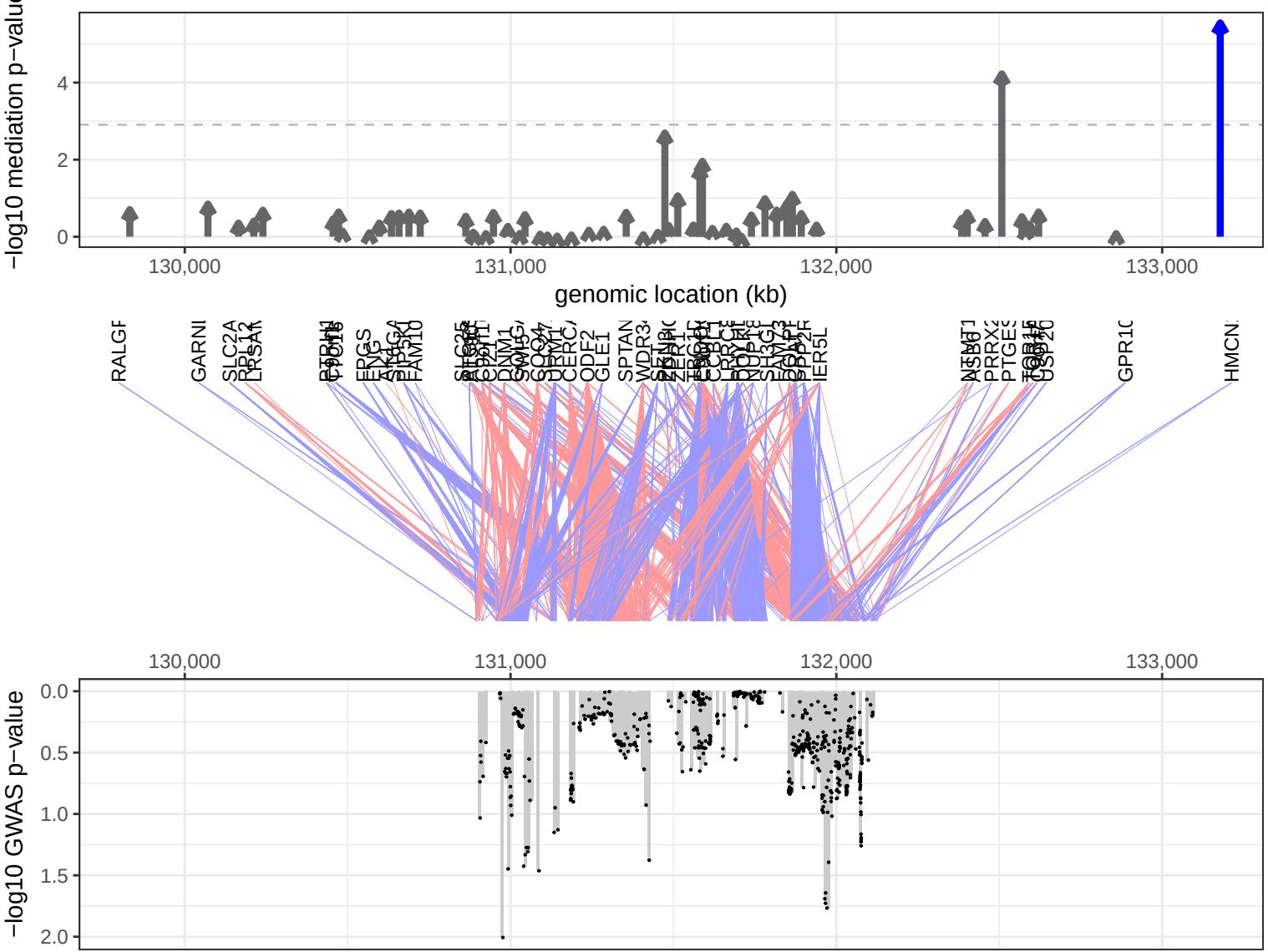
−log10 mediation p−value
0
2
4
130,000
131,000
132,000
133,000
genomic location (kb)
RALGF
GARNI
SLC2A
LRSAR
ERGS
FAM10
SPTAN
WDR3
CERC
GLE1
IER5L
PTGES
GPR1C
HMCN
130,000
131,000
132,000
133,000
−log10 GWAS p−value
0.0
0.5
1.0
1.5
2.0