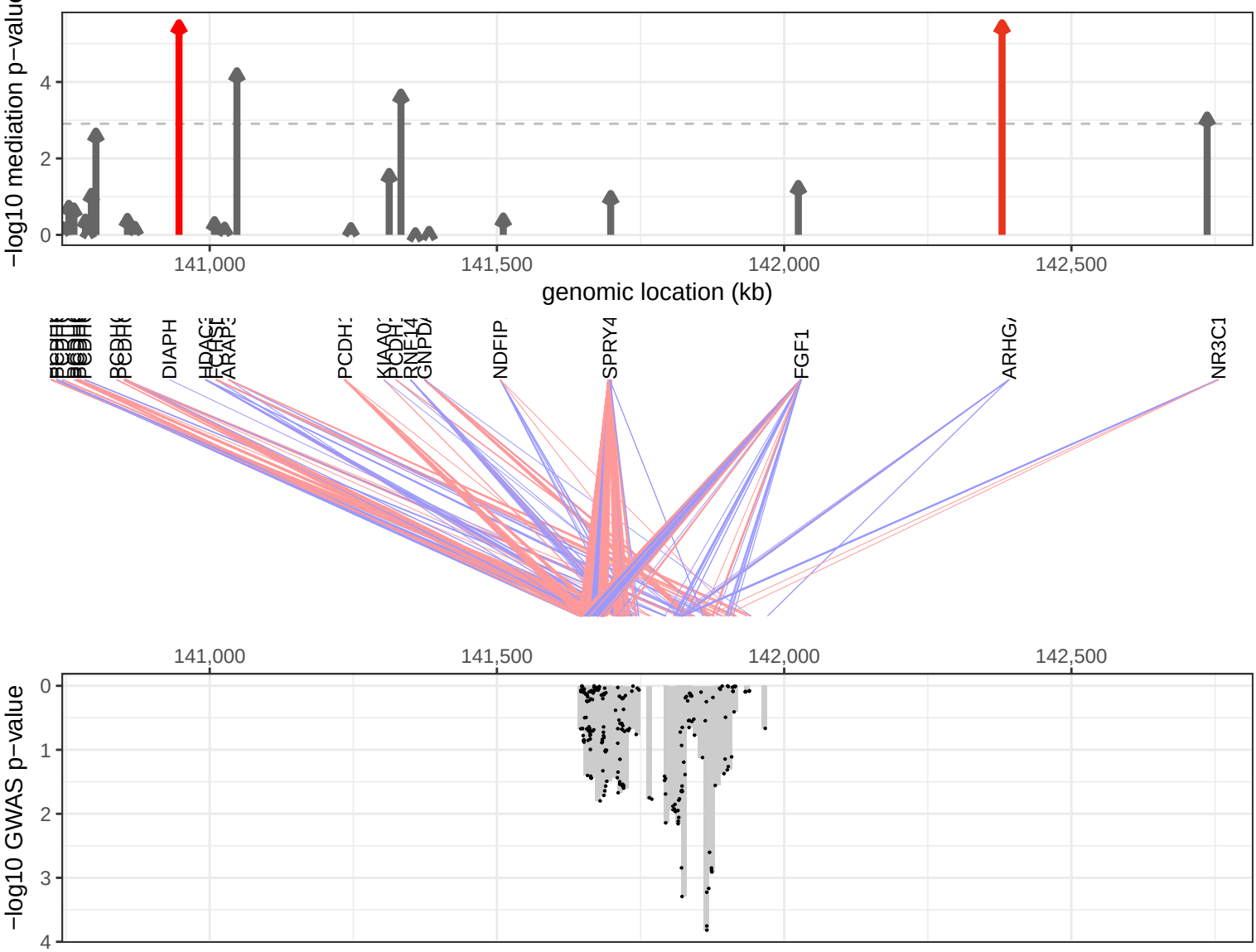
−log10 mediation p−value
0
2
4
141,000
141,500
142,000
142,500
genomic location (kb)
DIAPH
PCDH
NDFIP
SPRY4
FGF1
ARHG
NR3C1
141,000
141,500
142,000
142,500
−log10 GWAS p−value
0
1
2
3
4