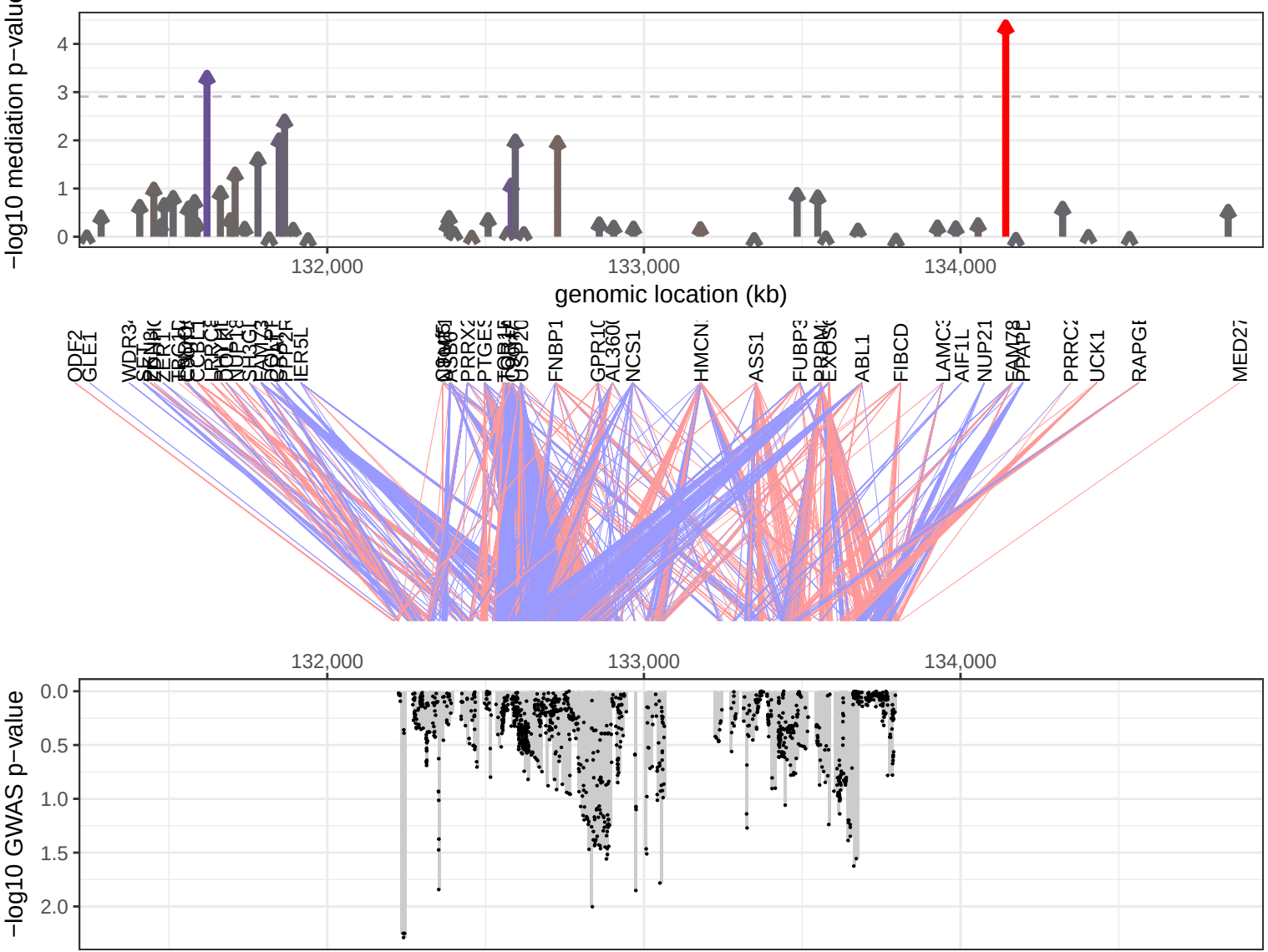
−log10 mediation p−value
4
3
2
1
0
132,000
133,000
134,000
genomic location (kb)
QDE2
GLE1
WDR3
IER5L
FNBP1
NCS1
HMCN
ASS1
FUBP3
ABL1
FIBCD
NUP21
PRRC2
UCK1
RAPGE
MED27
132,000
133,000
134,000
−log10 GWAS p−value
0.0
0.5
1.0
1.5
2.0