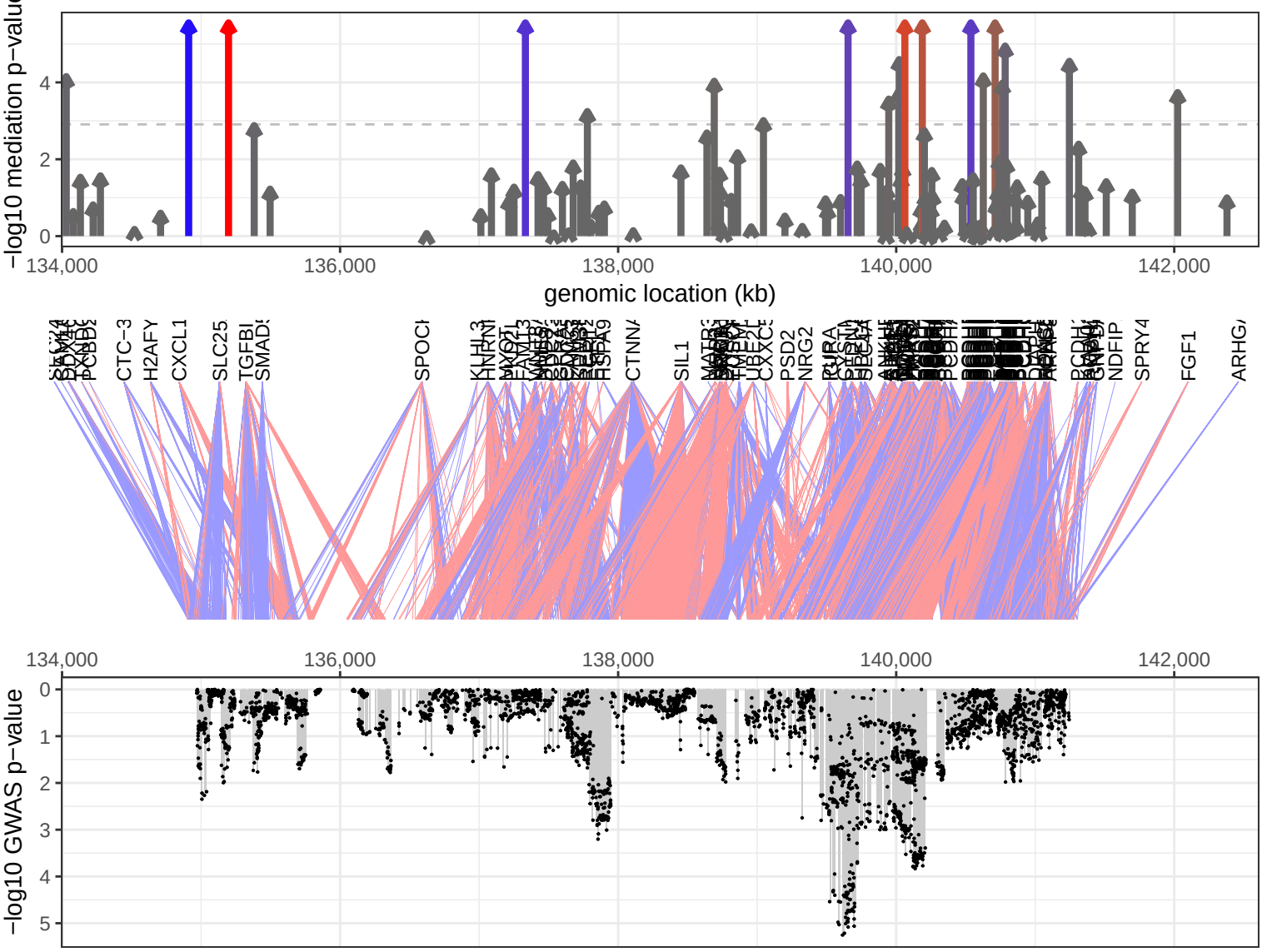
−log10 mediation p−value
0
2
4
134,000
136,000
138,000
140,000
142,000
genomic location (kb)
CTC−3
H2AFY
CXCL1
SLC25
TGFBI
SPOC
CTNN
SIL1
PSD2
NRG2
NDFIP
SPRY4
FGF1
−log10 GWAS p−value
0
1
2
3
4
5