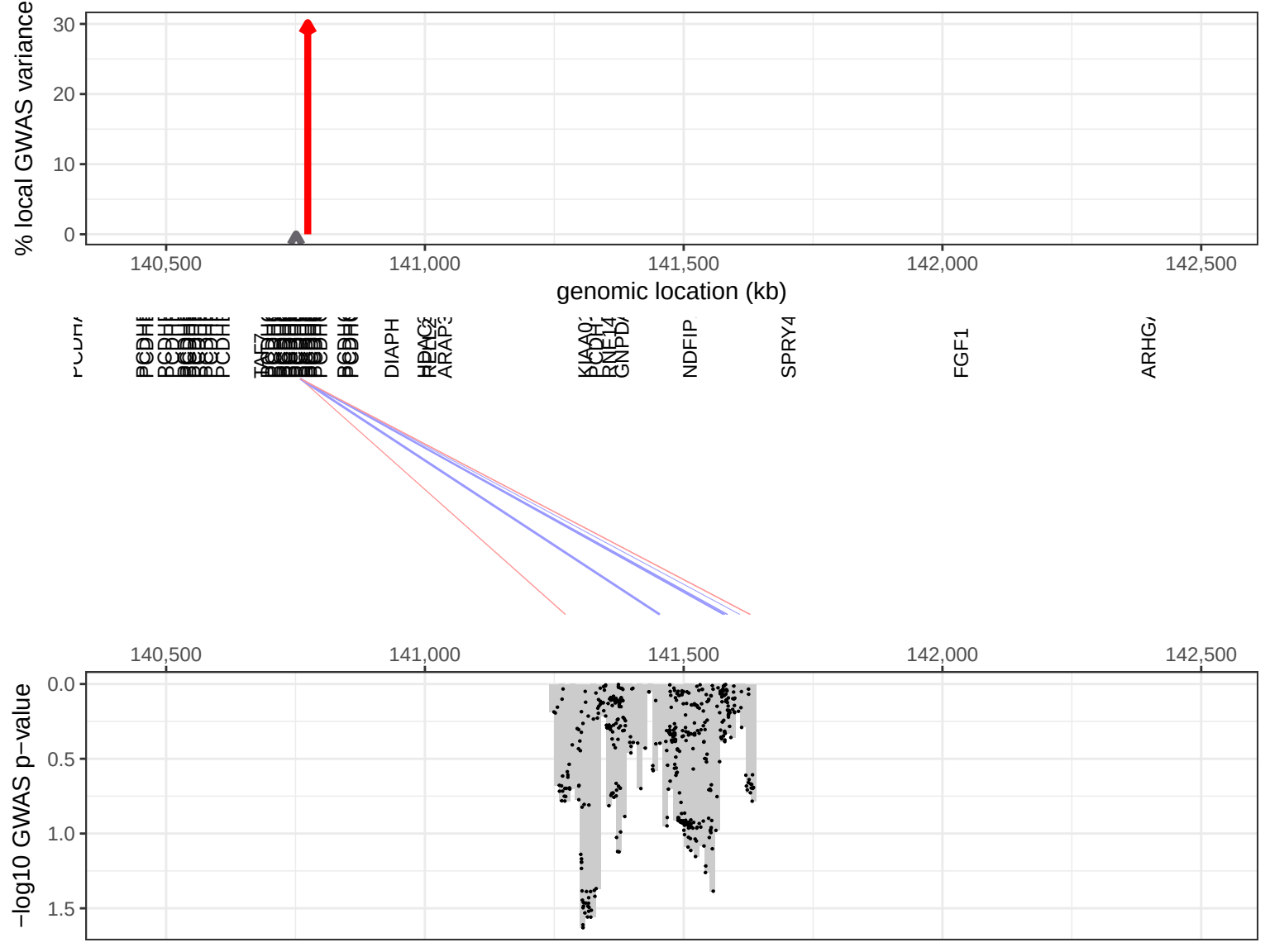
% local GWAS variance
30
20
10
0
140,500
141,000
141,500
142,000
142,500
genomic location (kb)
DIAPH
NDFIP
SPRY4
FGF1
ARHG
140,500
141,000
141,500
142,000
142,500
−log10 GWAS p−value
0.0
0.5
1.0
1.5