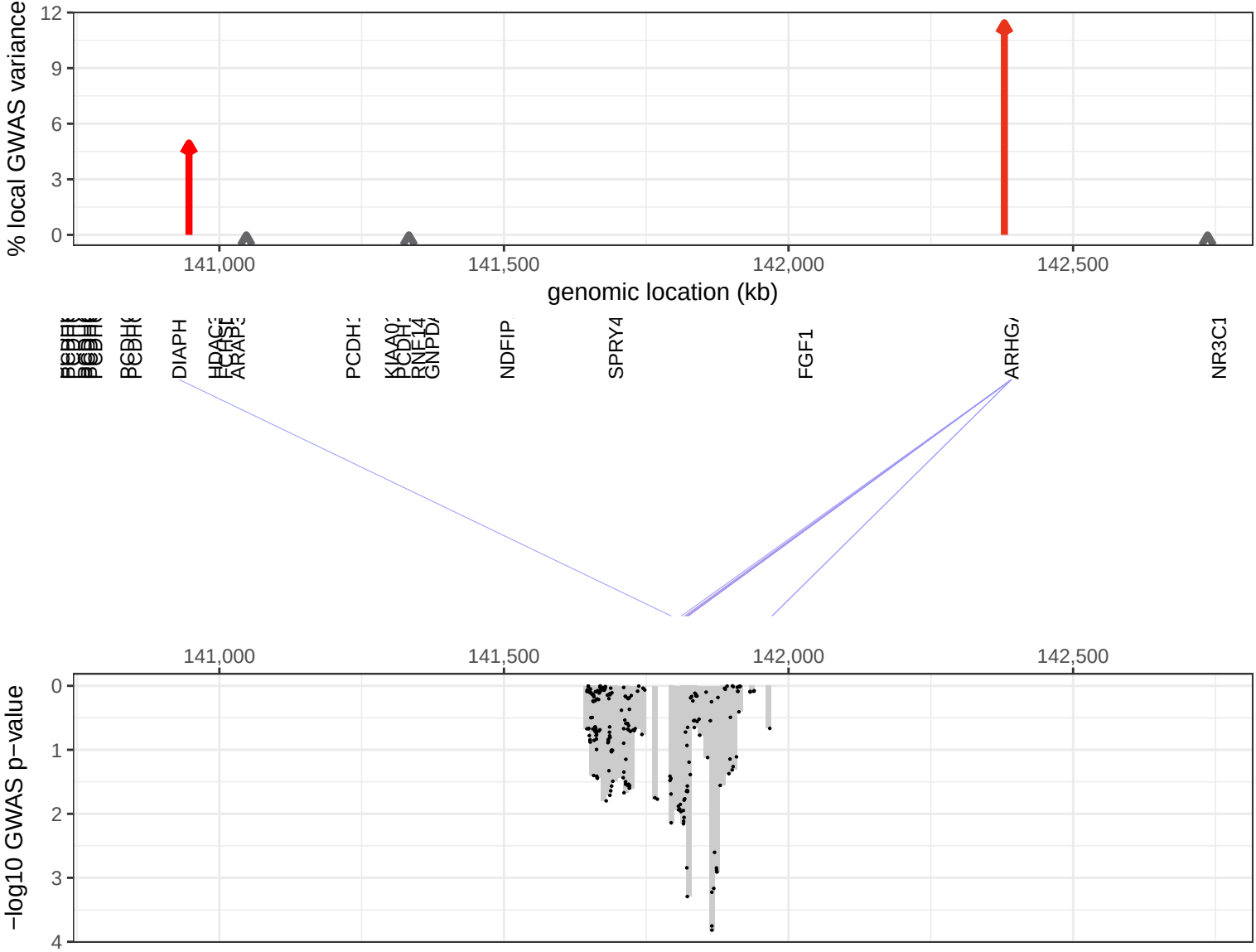
% local GWAS variance
12
9
6
3
0
141,000
141,500
142,000
142,500
genomic location (kb)
DIAPH
NDFIP
SPRY4
FGF1
NR3C1
−log10 GWAS p−value
0
1
2
3
4