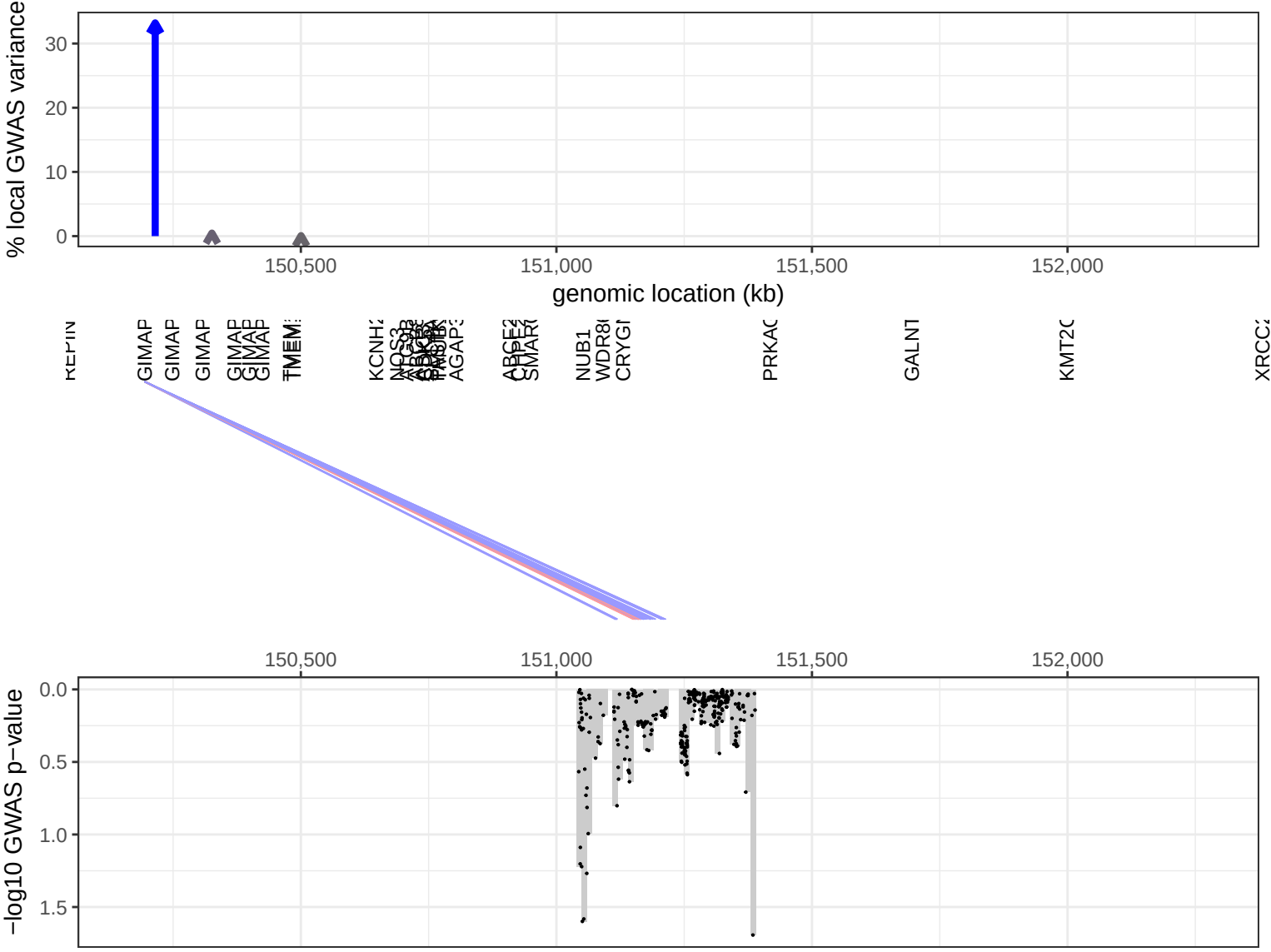
% local GWAS variance
30
20
10
0
150,500
151,000
151,500
152,000
genomic location (kb)
GIMAP
NUB1
WDR8
PRKAC
GALNT
KMT2C
150,500
151,000
151,500
152,000
−log10 GWAS p−value
0.0
0.5
1.0
1.5