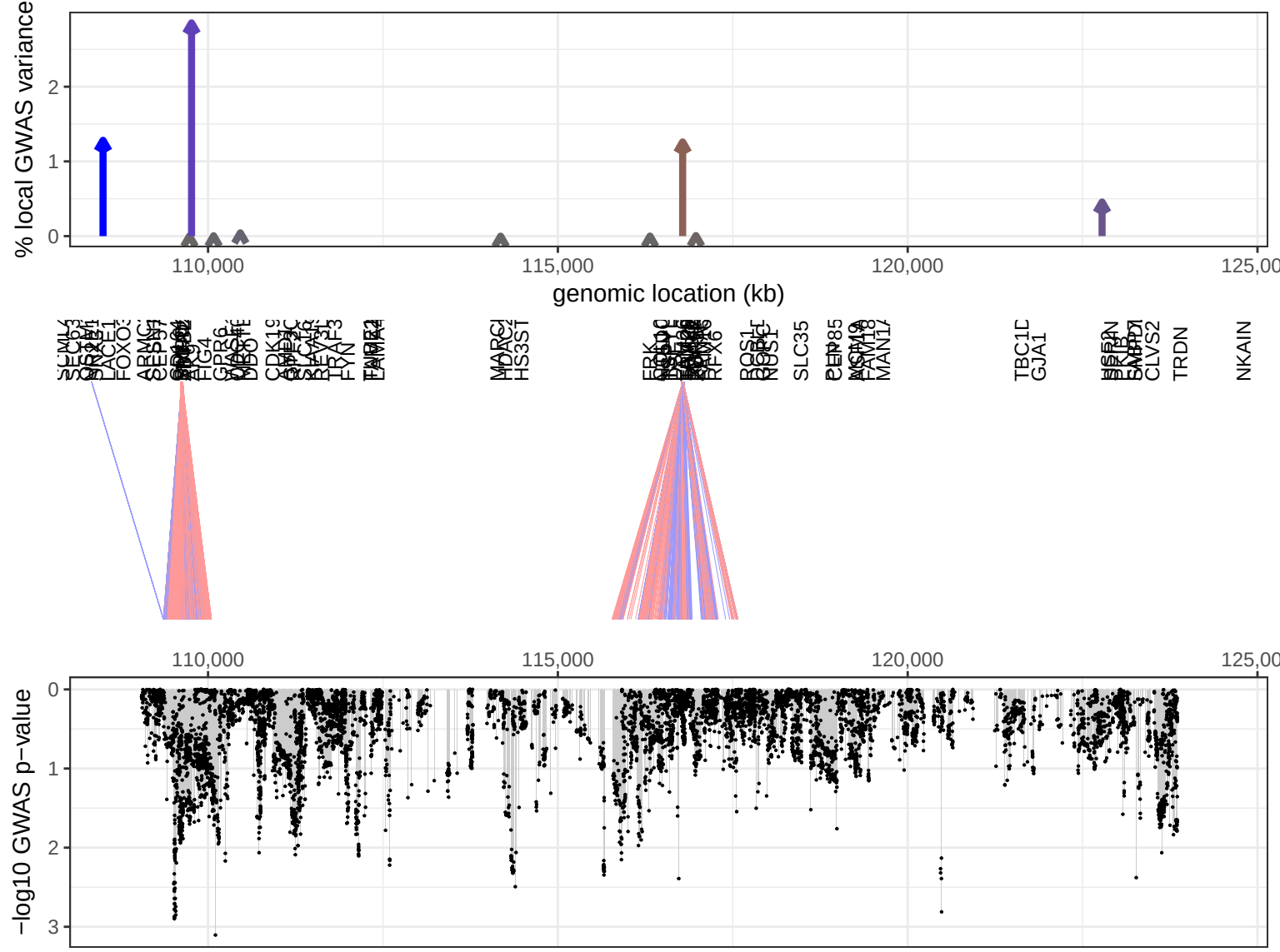
% local GWAS variance
0
1
2
110,000
115,000
120,000
125,00
genomic location (kb)
FOXO1
FIG4
HS3ST1
SLC35
TBC1D
GJA1
CLVS2
TRDN
NKAIN
110,000
115,000
120,000
125,00
−log10 GWAS p−value
0
1
2
3