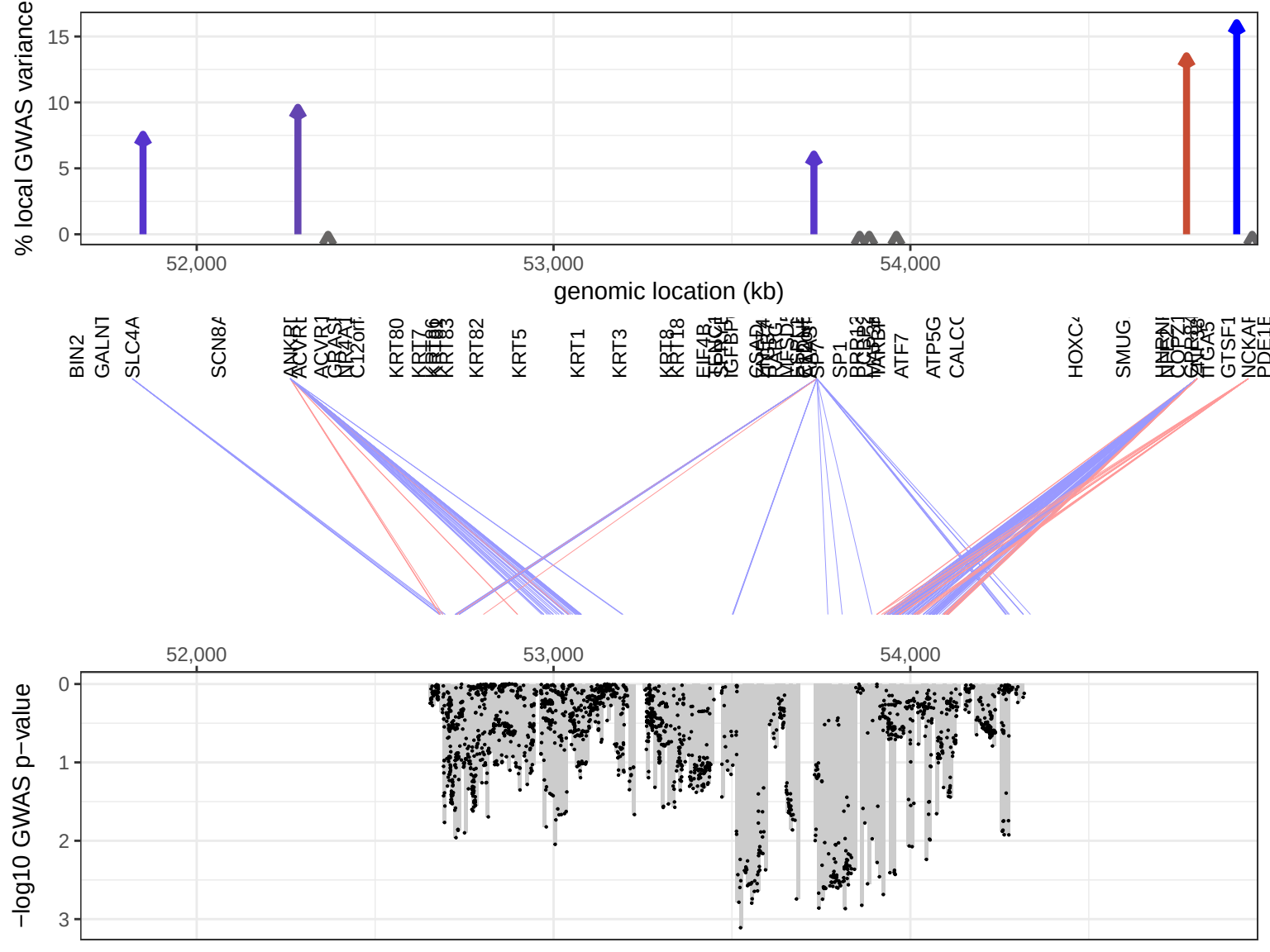
% local GWAS variance
15
10
5
0
52,000
53,000
54,000
genomic location (kb)
BIN2
GALNT
SLC4A
SCN8A
ACVR1
KRT80
KRT82
KRT5
KRT1
KRT3
KRT8
KRT18
SP1
ATF7
ATP5G
CALCC
HOXC4
SMUG
GTSF1
52,000
53,000
54,000
−log10 GWAS p−value
0
1
2
3