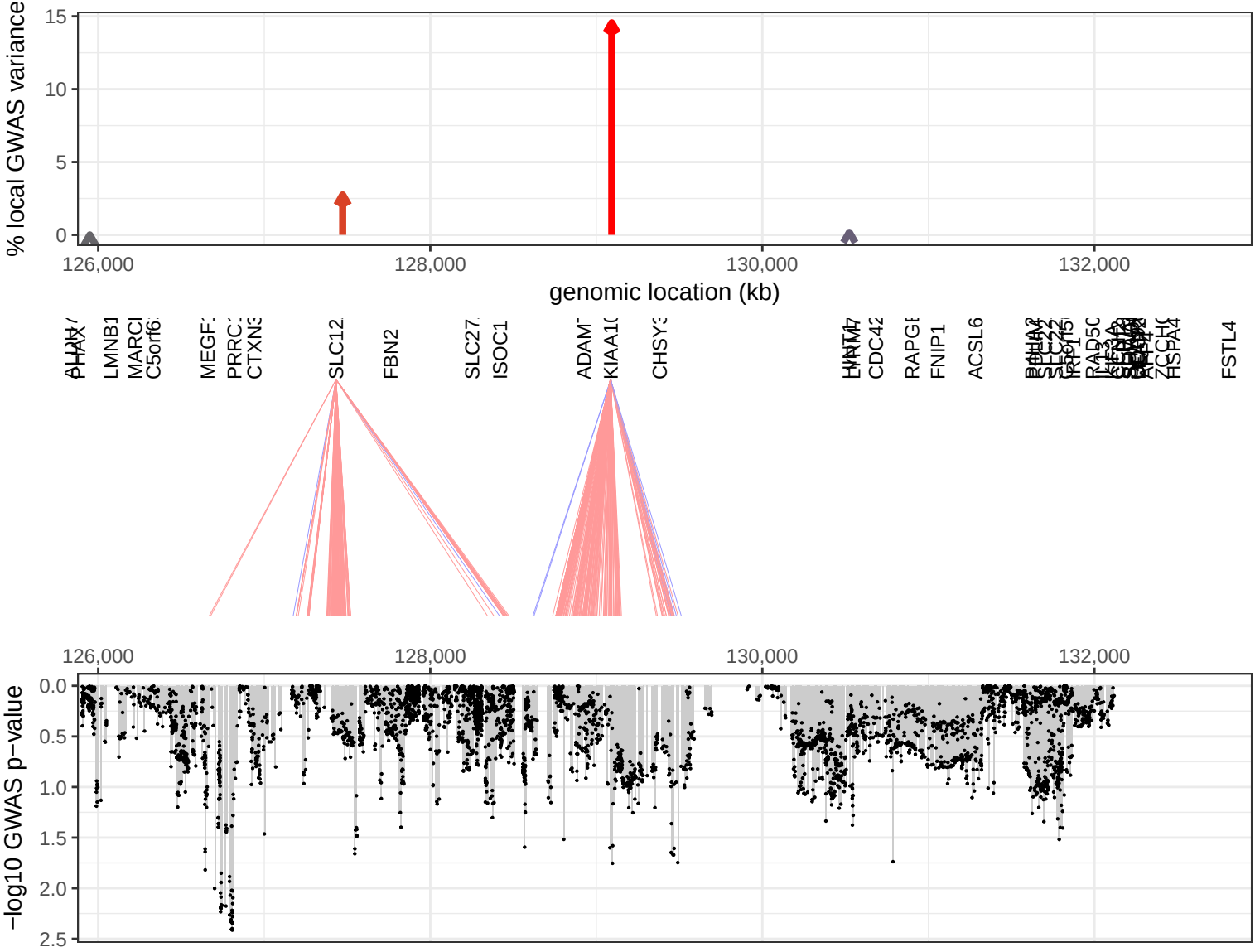
% local GWAS variance
15
10
5
0
126,000
128,000
130,000
132,000
genomic location (kb)
LMNB1
MARCH
C5orf6
MEGF
PRRC
CTXN
SLC12
FBN2
SLC27
ISOC1
ADAM
KIAA1
CHSY
CDC42
RAPGE
FNIP1
ACSL6
RAD50
FSTL4
126,000
128,000
130,000
132,000
−log10 GWAS p−value
0.0
0.5
1.0
1.5
2.0
2.5