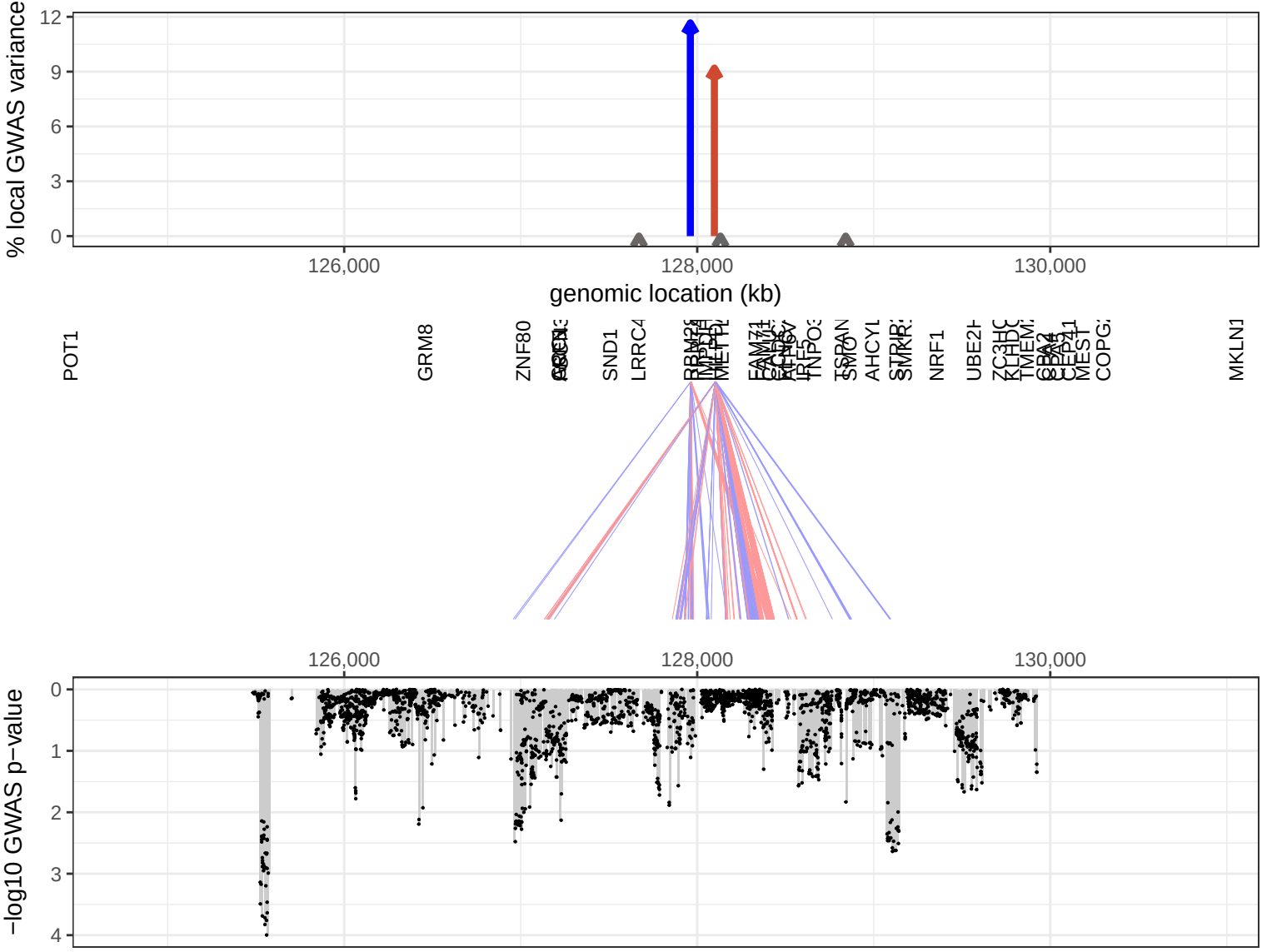
% local GWAS variance
12
9
6
3
0
126,000
128,000
130,000
genomic location (kb)
POT1
GRM8
ZNF80
SND1
LRRC4
AHCY1
NRF1
UBE2I
MKLN1
−log10 GWAS p−value
0
1
2
3
4