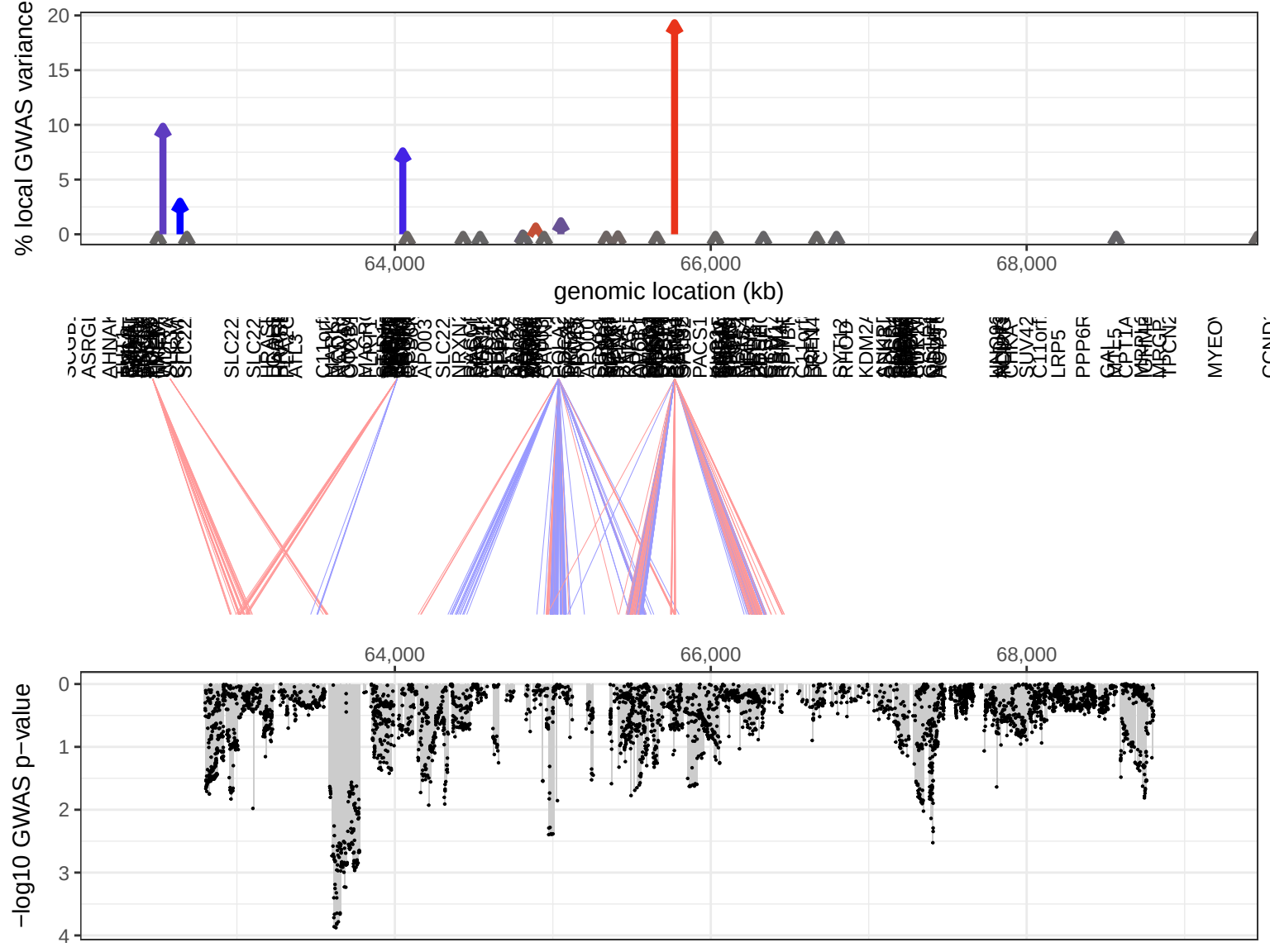
% local GWAS variance
20
15
10
5
0
64,000
66,000
68,000
genomic location (kb)
SLC22
LRP5
PPP6R
MYEO
−log10 GWAS p−value
0
1
2
3
4