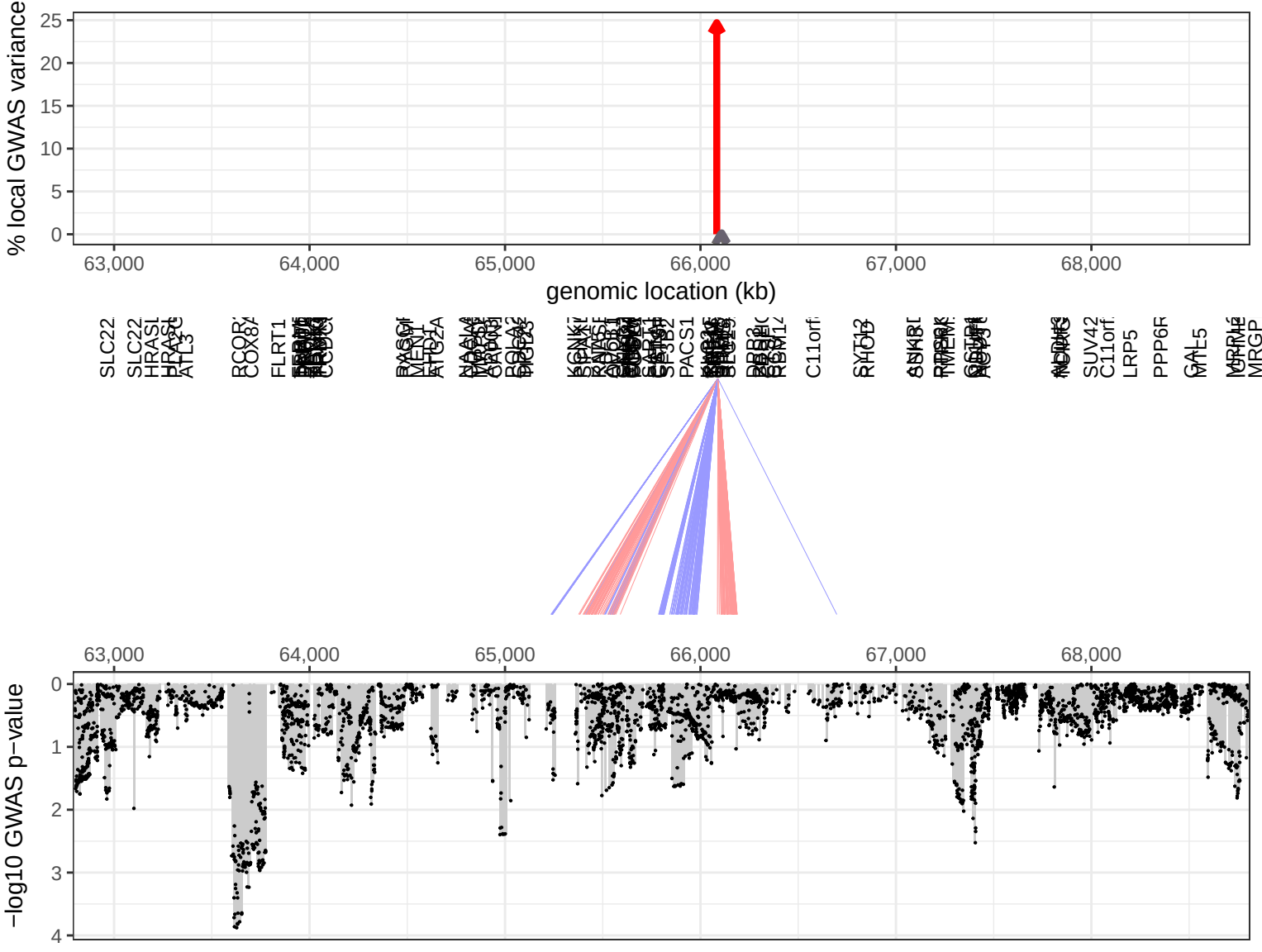
% local GWAS variance
25
20
15
10
5
0
63,000
64,000
65,000
66,000
67,000
68,000
genomic location (kb)
SLC22
SLC22
HRASL
ATL3
FLRT1
PACS1
C11orf
SUV42
C11orf
LRP5
PPP6F
−log10 GWAS p−value
0
1
2
3
4