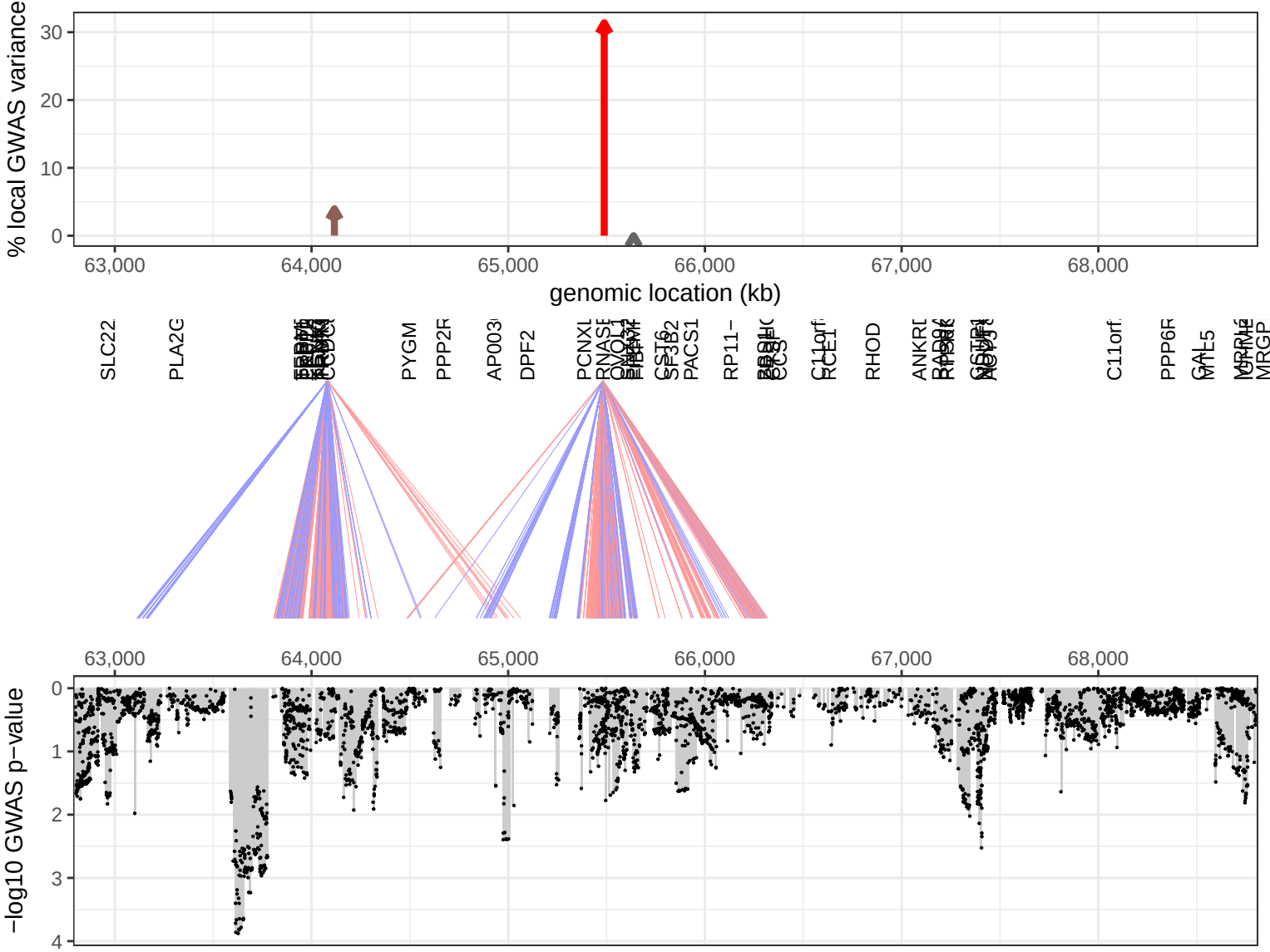
% local GWAS variance
30
20
10
0
63,000
64,000
65,000
66,000
67,000
68,000
genomic location (kb)
SLC22
PLA2G
PYGM
PPP2R
AP003
DPF2
PCNXL
SF3B2
PACS1
RP11−
C11orf
RCE1
RHOD
ANKRD
C11orf
PPP6R
−log10 GWAS p−value
0
1
2
3
4