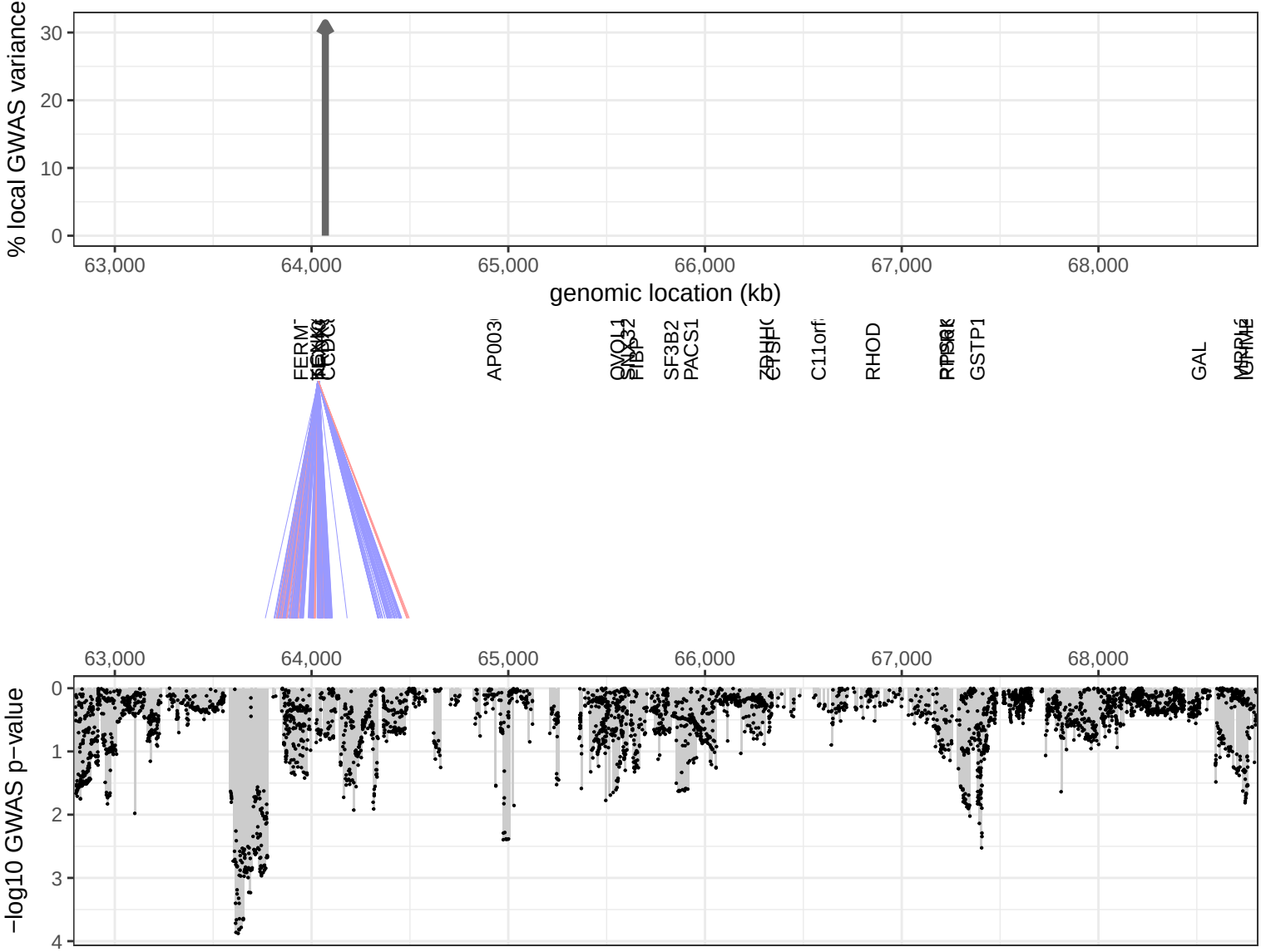
% local GWAS variance
30
20
10
0
63,000
64,000
65,000
66,000
67,000
68,000
genomic location (kb)
AP003
SF3B2
PACS1
C11orf
RHOD
GSTP1
GAL
−log10 GWAS p−value
0
1
2
3
4