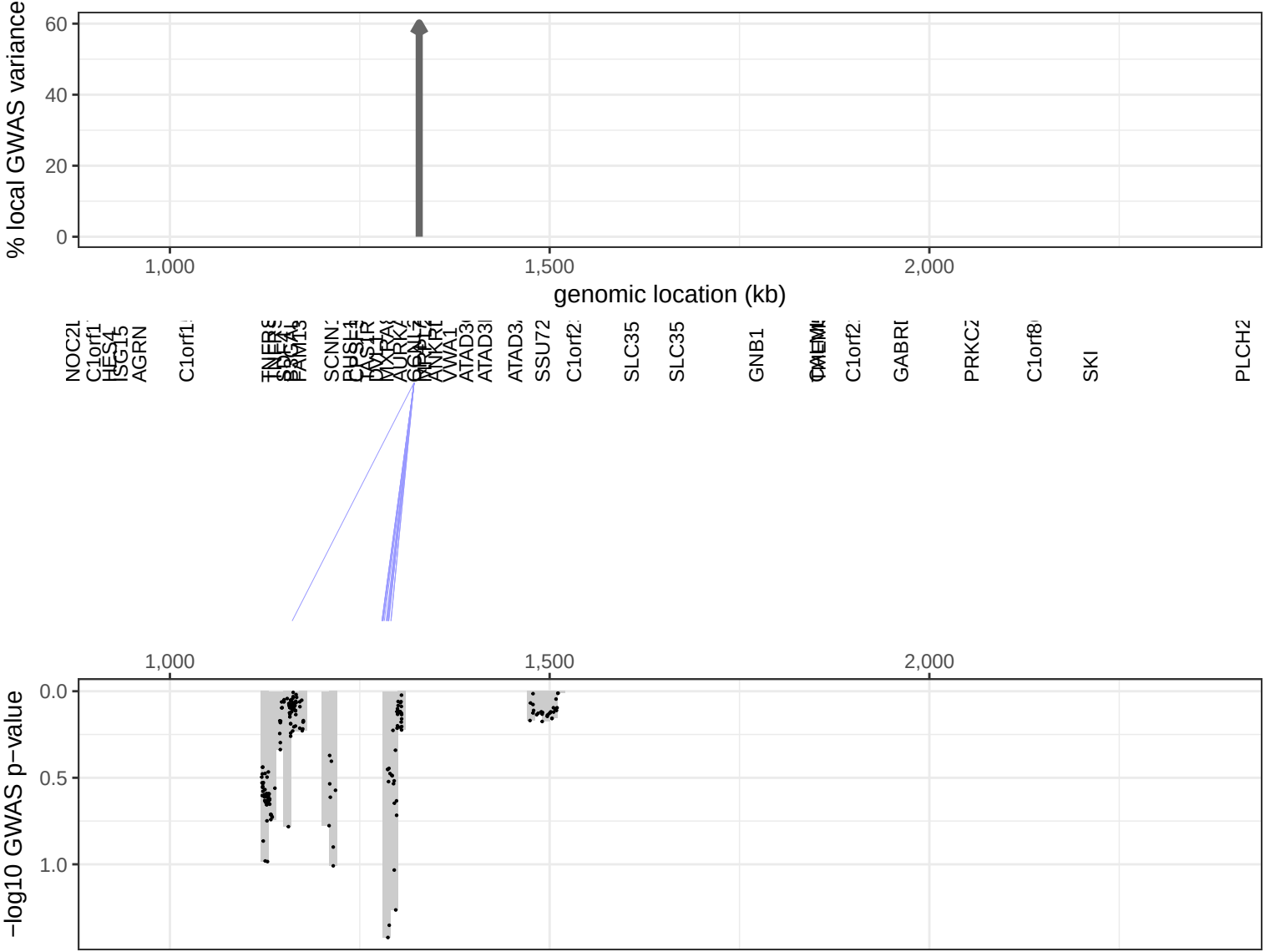
% local GWAS variance
60
40
20
0
1,000
1,500
2,000
genomic location (kb)
AGRN
C1orf1
SCNN
VWA1
ATAD3
ATAD3
SSU72
C1orf2
SLC35
SLC35
GNB1
C1orf2
GABR
PRKC
C1orf8
SKI
PLCH
−log10 GWAS p−value
0.0
0.5
1.0
1,000
1,500
2,000