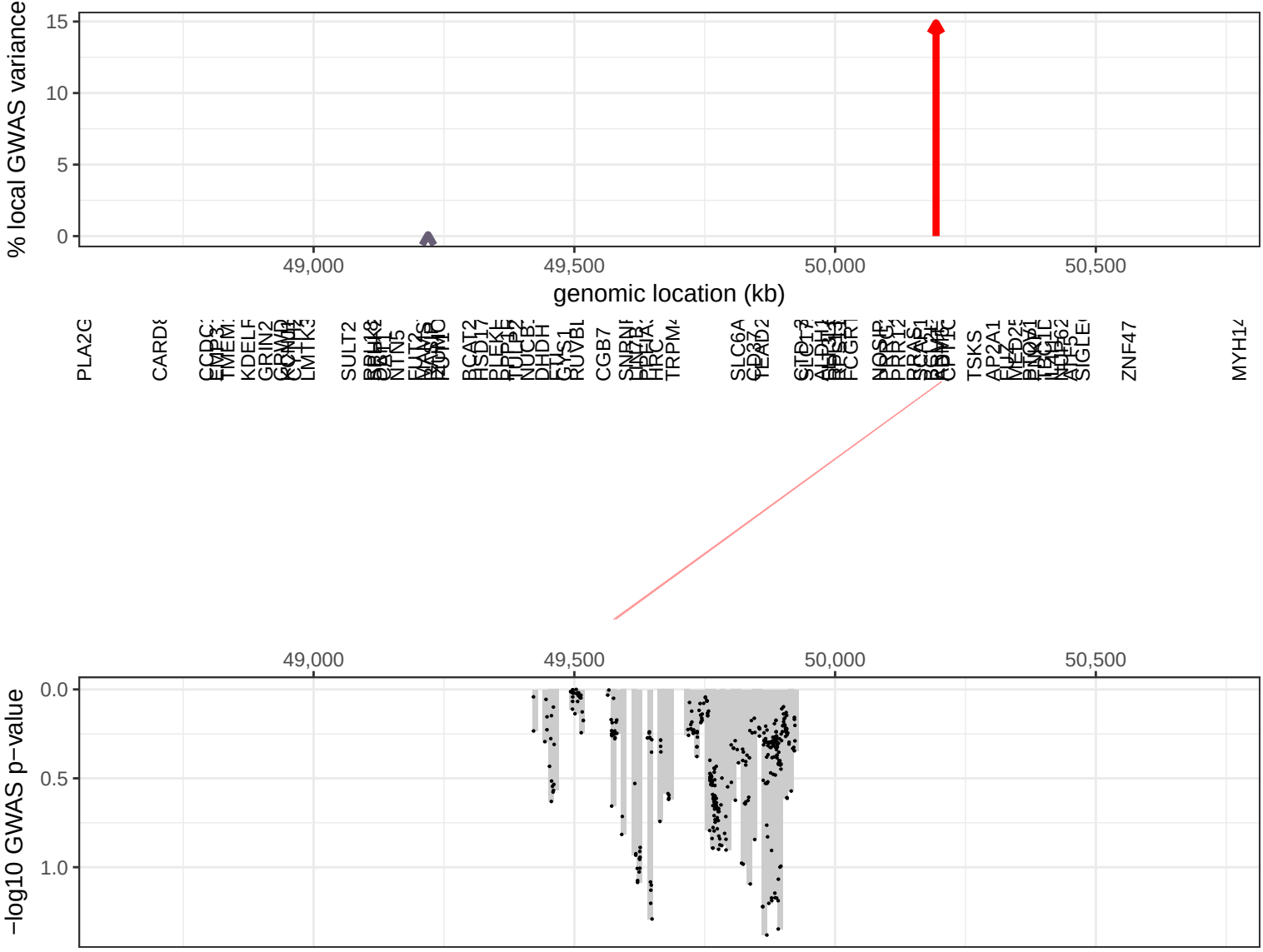
% local GWAS variance
15
10
5
0
49,000
49,500
50,000
50,500
genomic location (kb)
PLA2G
CARD
KDELF
GRIN2
LMTK
SULT2
NTN5
RSAT2
NUCB
DHDH
RUVBL
CGB7
TRPM
SLC6A
TEAD2
TSKS
AP2A1
SIGLE
ZNF47
MYH
−log10 GWAS p−value
0.0
0.5
1.0