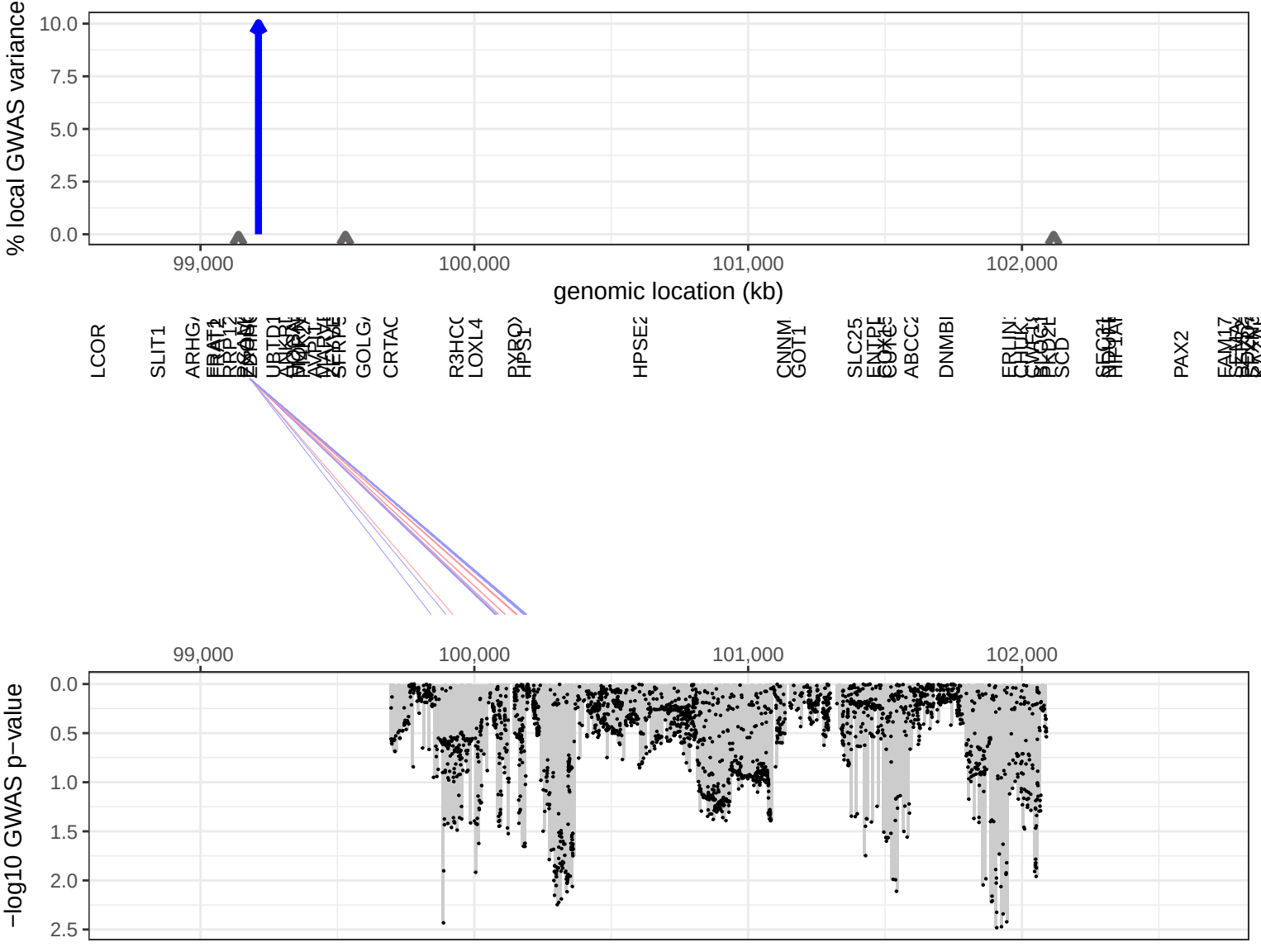
% local GWAS variance
10.0
7.5
5.0
2.5
0.0
99,000
100,000
101,000
102,000
genomic location (kb)
LCOR
SLIT1
GOLG
CRTAC
R3HCC
LOXL4
HPSE2
CNNM
GOT1
SLC25
ABCC2
DNMBI
SCD
PAX2
−log10 GWAS p−value
0.0
0.5
1.0
1.5
2.0
2.5