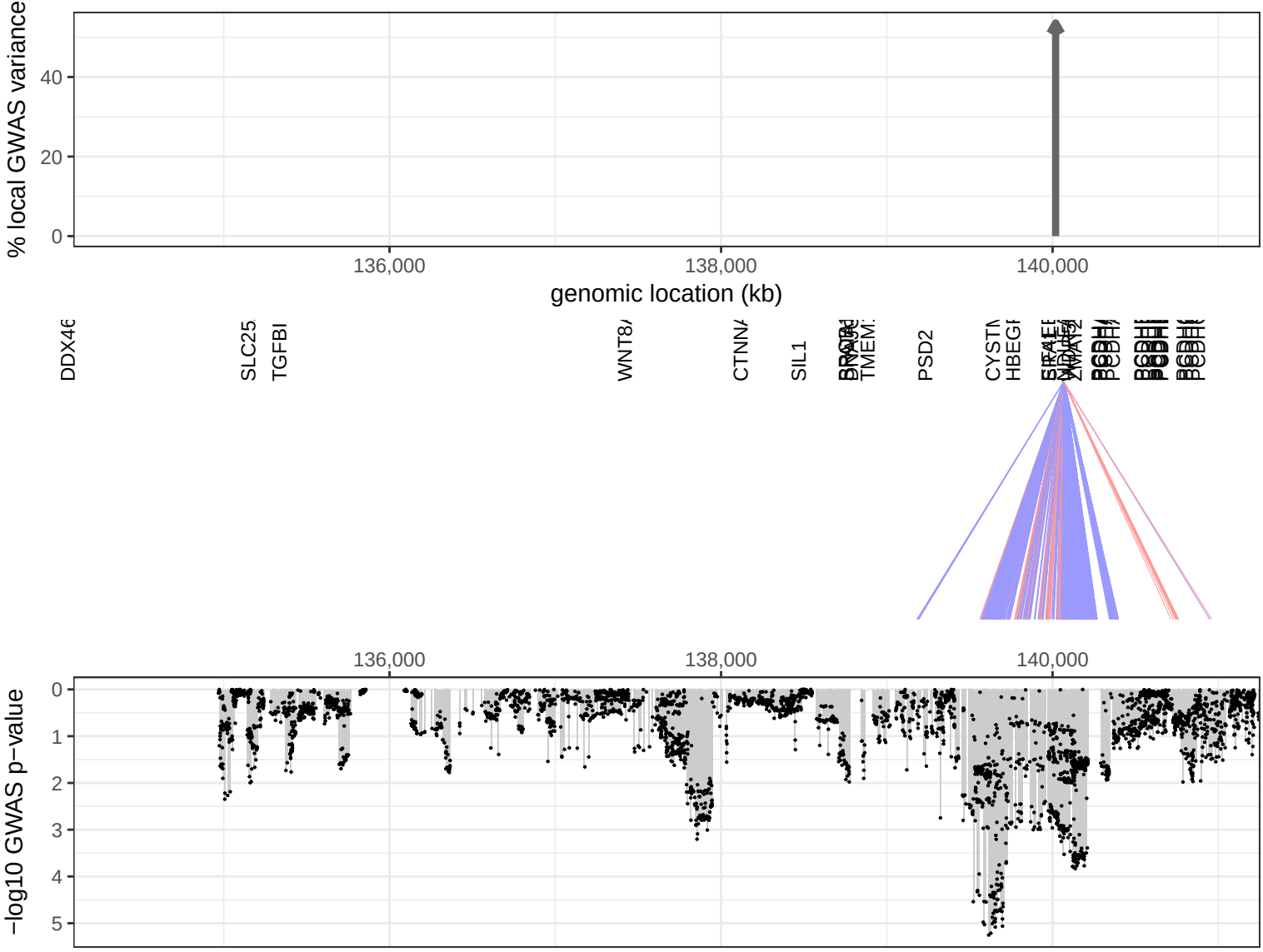
% local GWAS variance
0
20
40
136,000
138,000
140,000
genomic location (kb)
DDX46
SLC25
TGFBI
WNT8A
CTNNA
SIL1
TMEM
PSD2
CYSTM
HBEGF
−log10 GWAS p−value
0
1
2
3
4
5
136,000
138,000
140,000