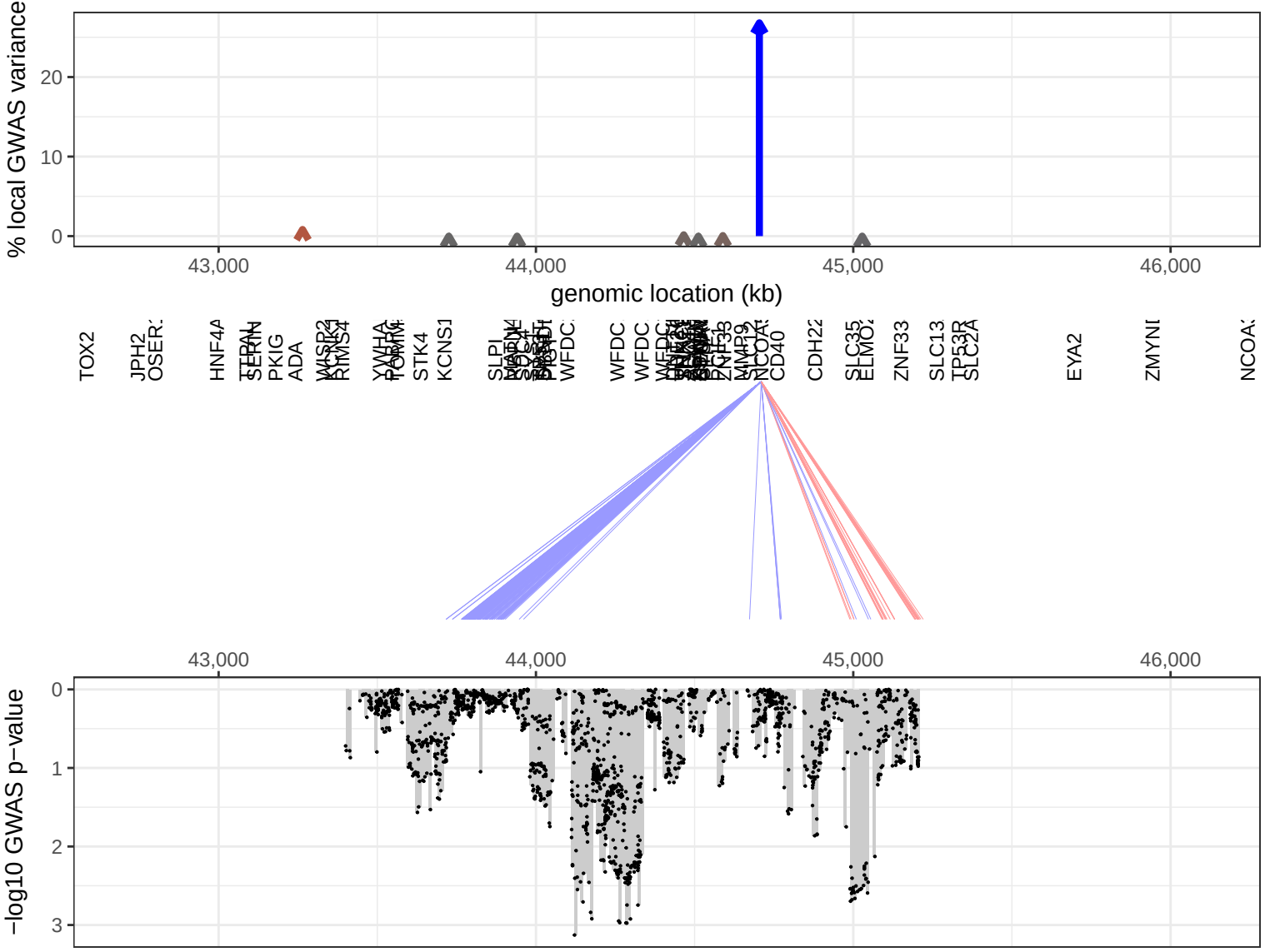
% local GWAS variance
0
10
20
43,000
44,000
45,000
46,000
genomic location (kb)
TOX2
JPH2
HNF4A
PKIG
ADA
STK4
KCNS1
WFDC
CD40
CDH22
SLC35
ELMO2
ZNF33
SLC13
EYA2
ZMYND
NCOA
−log10 GWAS p−value
0
1
2
3