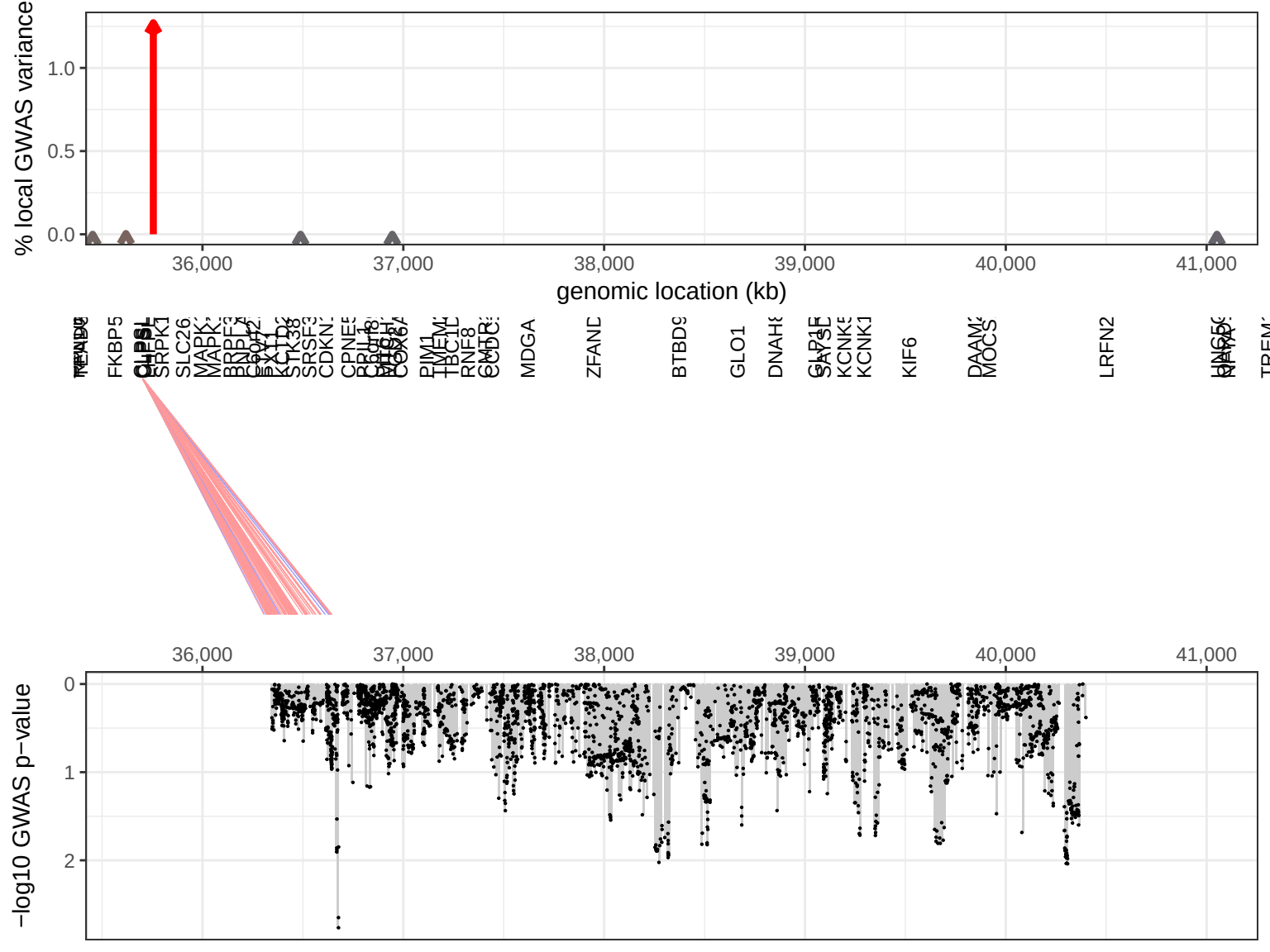
% local GWAS variance
1.0
0.5
0.0
36,000
37,000
38,000
39,000
40,000
41,000
genomic location (kb)
FKBP5
SRPK1
SLC26
STK38
SRSF
CDKN
CPNE
RNF8
MDGA
ZFAND
BTBD
GLO1
DNAH
KCNK
KCNK1
KIF6
MOCS
LRFN2
−log10 GWAS p−value
0
1
2