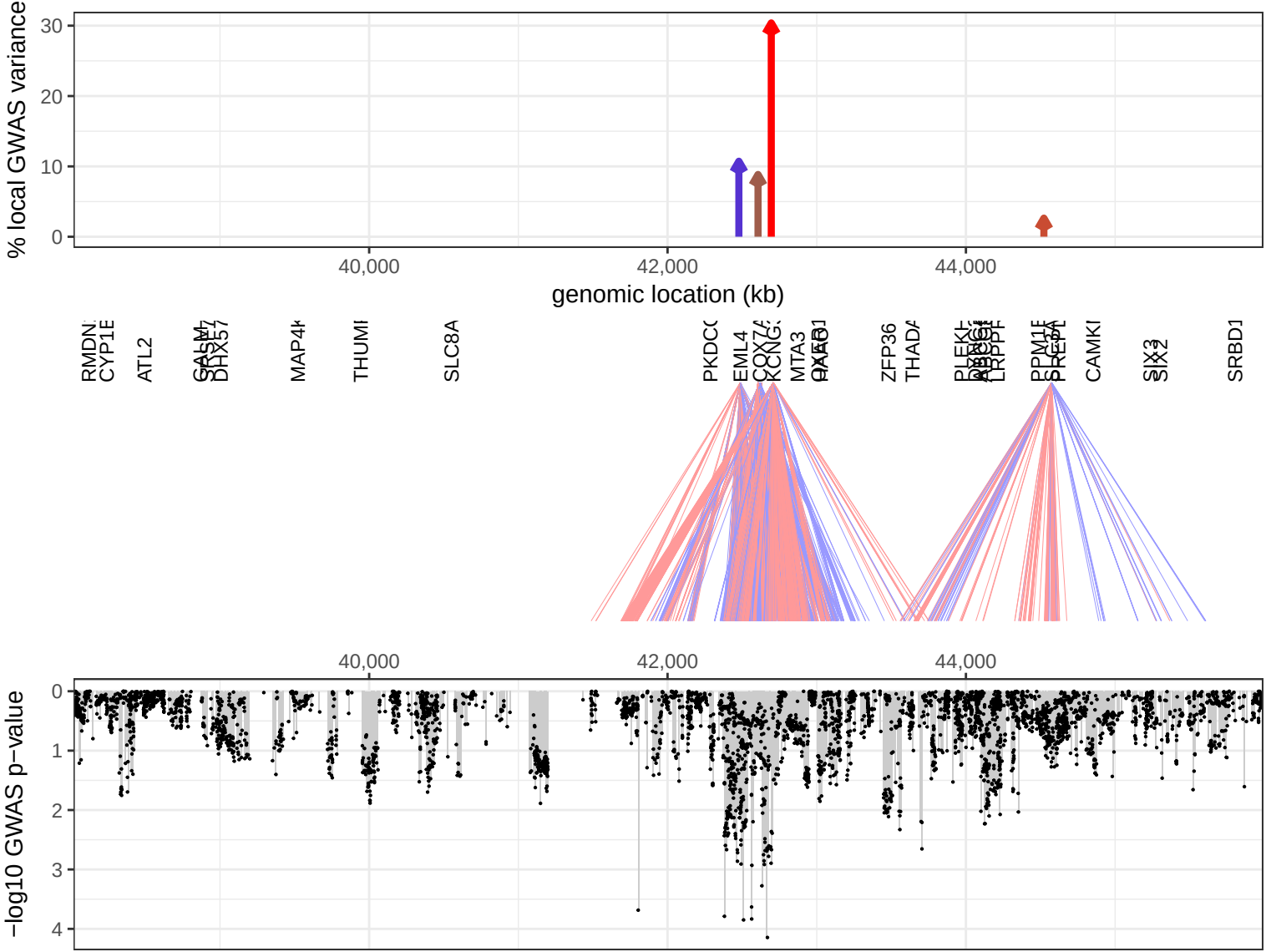
% local GWAS variance
30
20
10
0
40,000
42,000
44,000
genomic location (kb)
RMDN
CYP1E
ATL2
DHX57
MAP4K
THUMP
SLC8A
PKDCC
EML4
MTA3
ZFP36
THAD
CAMK
SIX3
SIX2
SRBD1
40,000
42,000
44,000
−log10 GWAS p−value
0
1
2
3
4