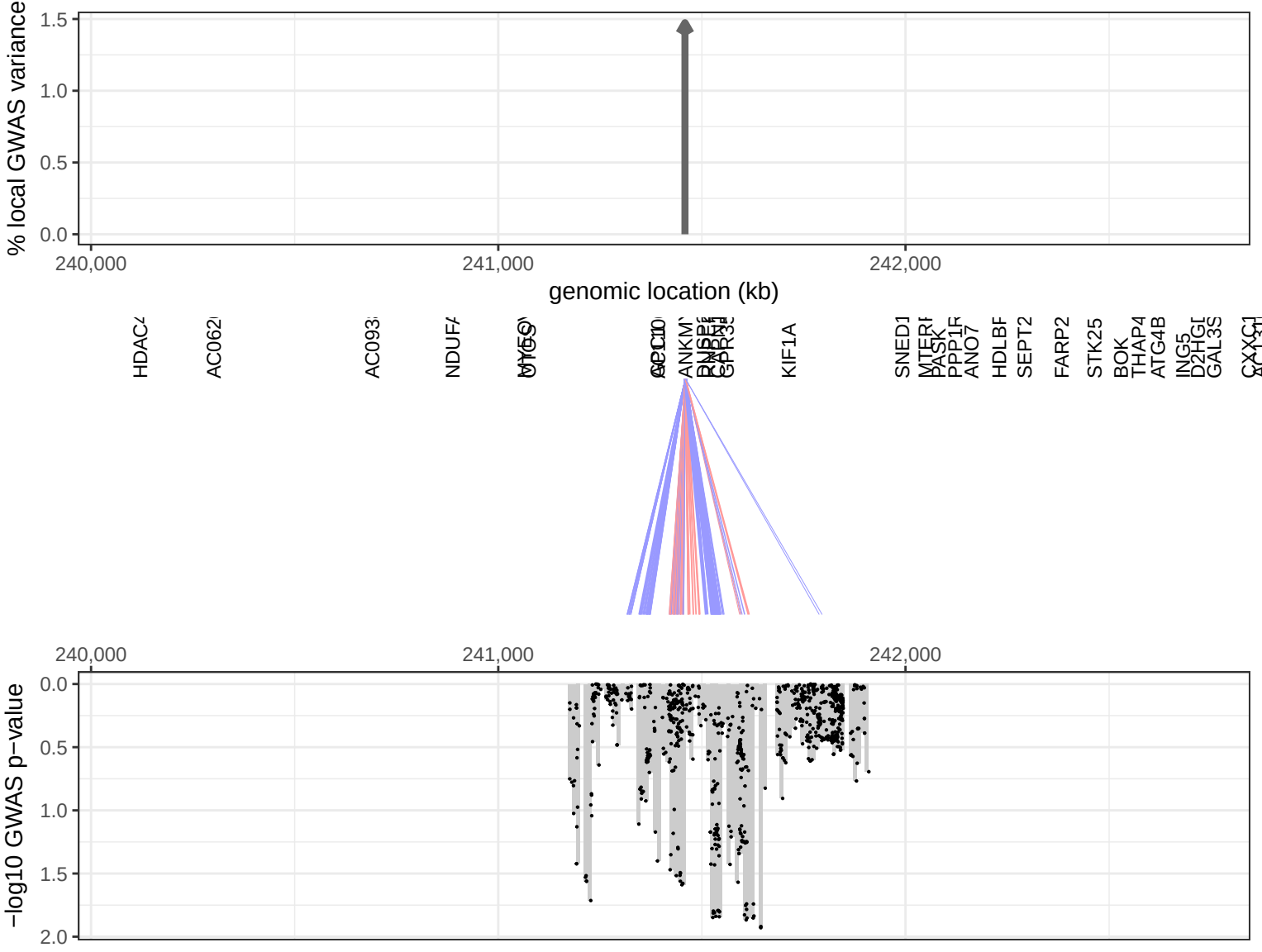
% local GWAS variance
1.5
1.0
0.5
0.0
240,000
241,000
242,000
genomic location (kb)
KIF1A
SNED1
HDLBP
SEPT2
FARP2
STK25
BOK
THAP4
ATG4B
ANO7
−log10 GWAS p−value
2.0