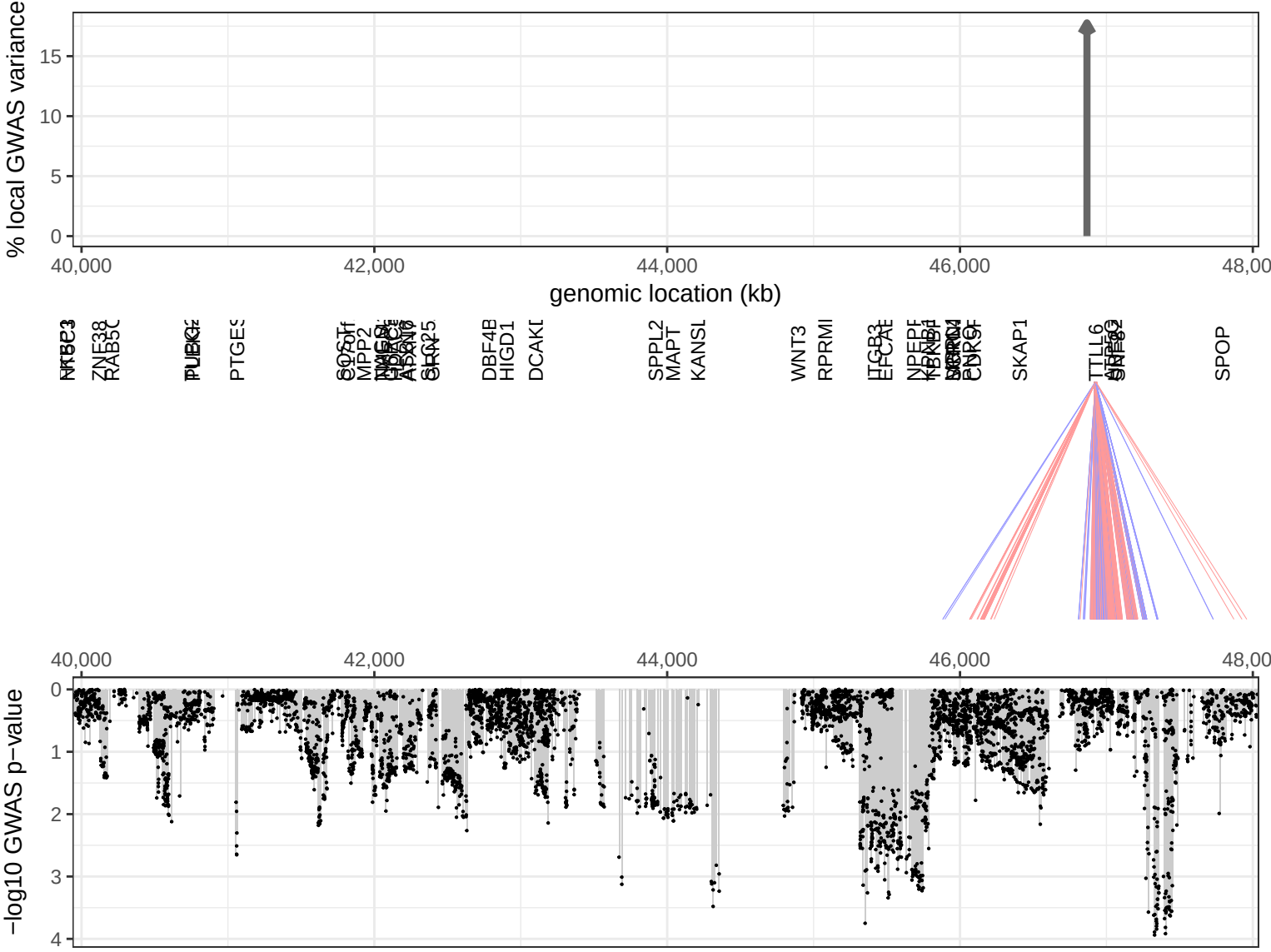
% local GWAS variance
0
5
10
15
40,000
42,000
44,000
46,000
48,00
genomic location (kb)
PTGES
DBF4B
HIGD1
DCAKD
SPPL2
MAPT
KANSL
WNT3
RPRMI
SKAP1
SPOP
−log10 GWAS p−value
0
1
2
3
4