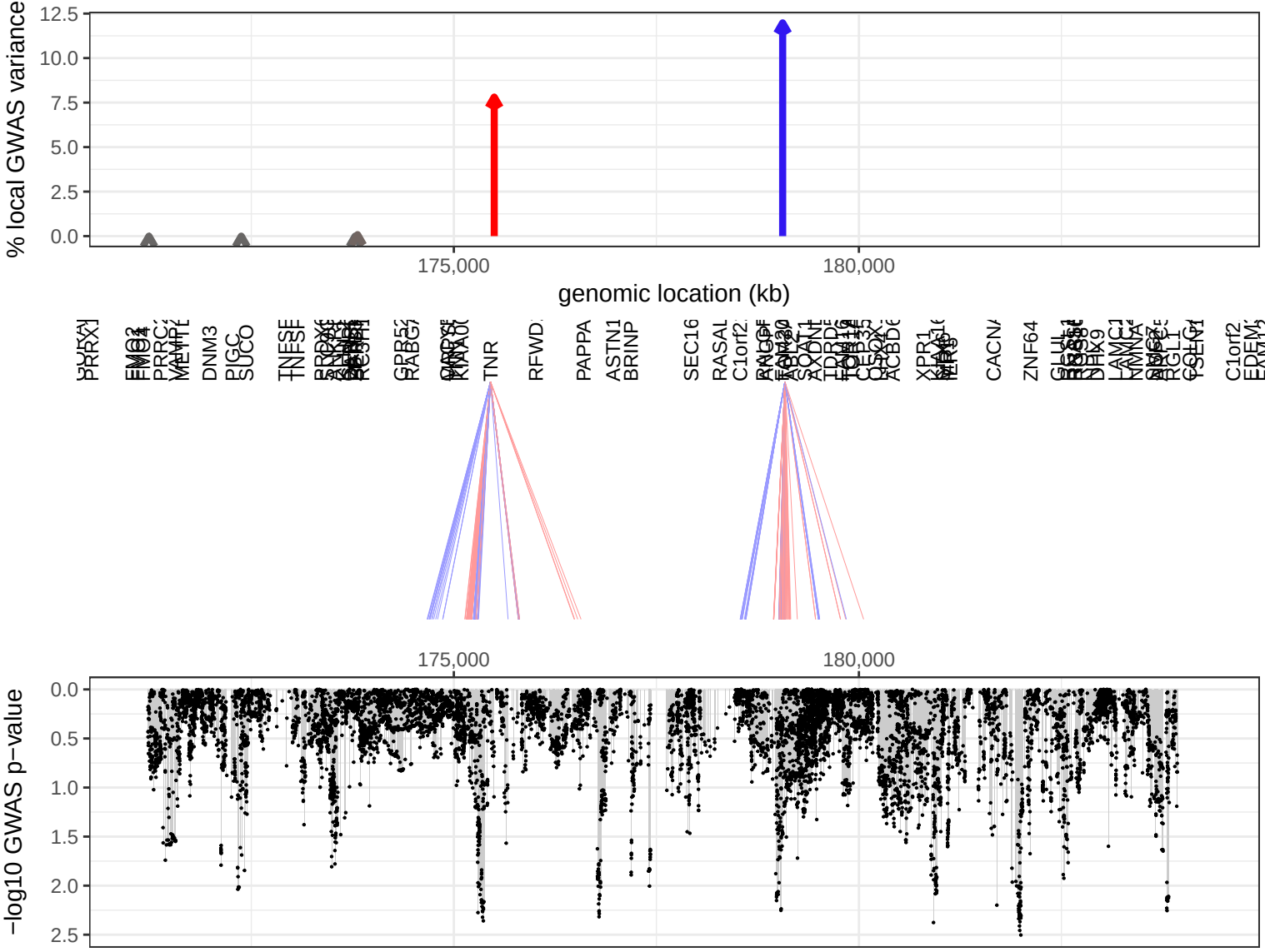
% local GWAS variance
12.5
10.0
7.5
5.0
2.5
0.0
175,000
180,000
genomic location (kb)
DNM3
PIGC
SUCO
TNESF
GPB5
RABG
TNR
RFWD
PAPPA
ASTN1
BRINP
SEC16
RASAL
C1orf2
XPR1
CACN
ZNF64
C1orf2
175,000
180,000
−log10 GWAS p−value
0.0
0.5
1.0
1.5
2.0
2.5