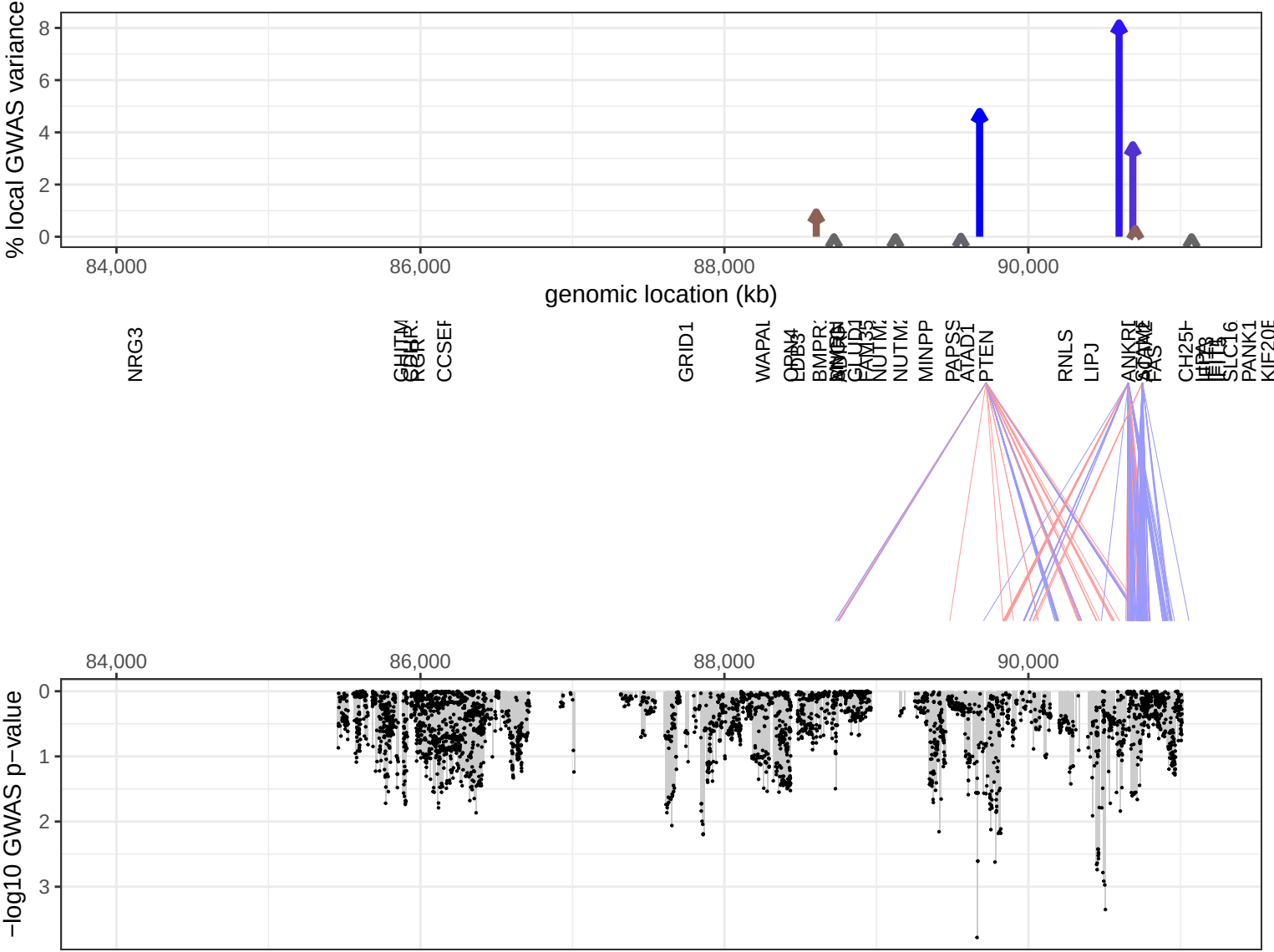
% local GWAS variance
0
2
4
6
8
84,000
86,000
88,000
90,000
genomic location (kb)
NRG3
CCSER
GRID1
WAPAL
MINPP
PAPSS
ATAD1
PTEN
RNLS
LIPJ
PAS
PANK1
−log10 GWAS p−value
0
1
2
3
84,000
86,000
88,000
90,000