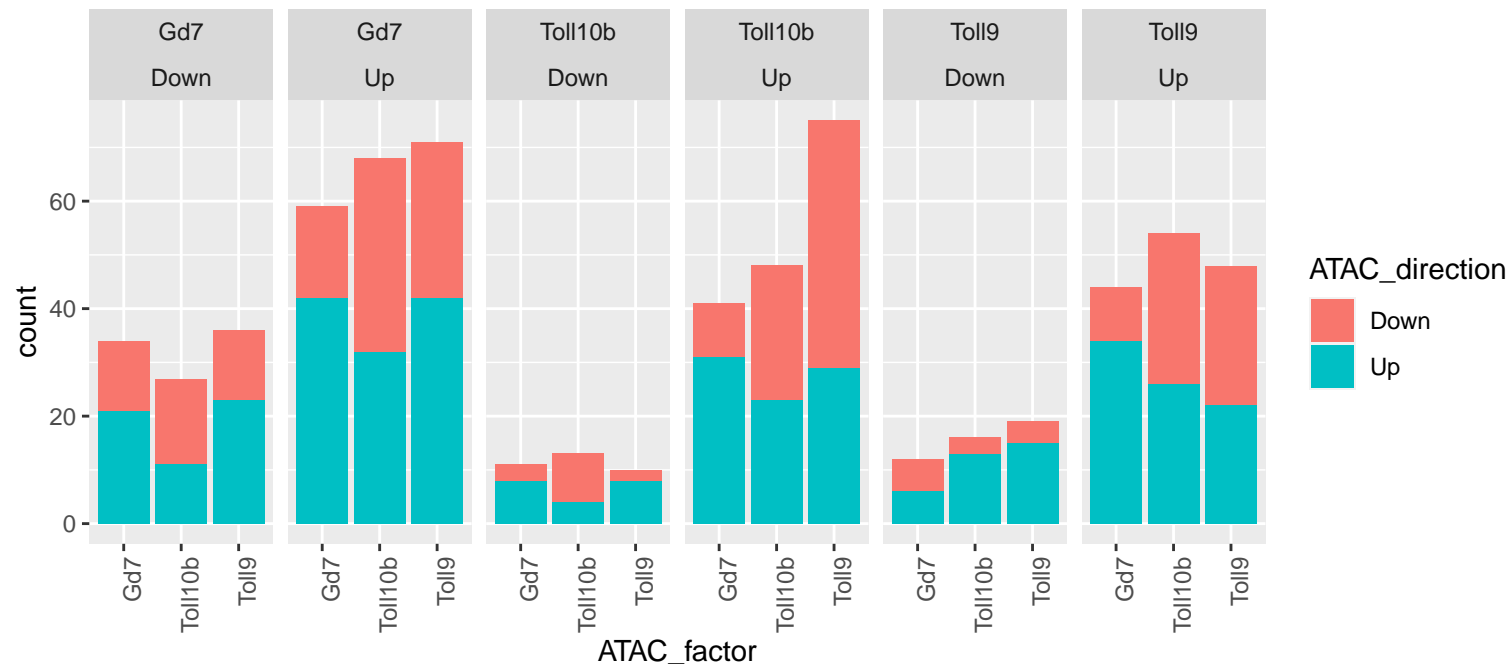


ATACseq peak and PROseq Gene correlation

100 – 1000 kb up– or downstream of gene



Gene is subset of genes that were identified using PROseq data

Genes are divided based on PROseq analysis annotation

Peak is 100 – 1000 kb up– or downstream of gene