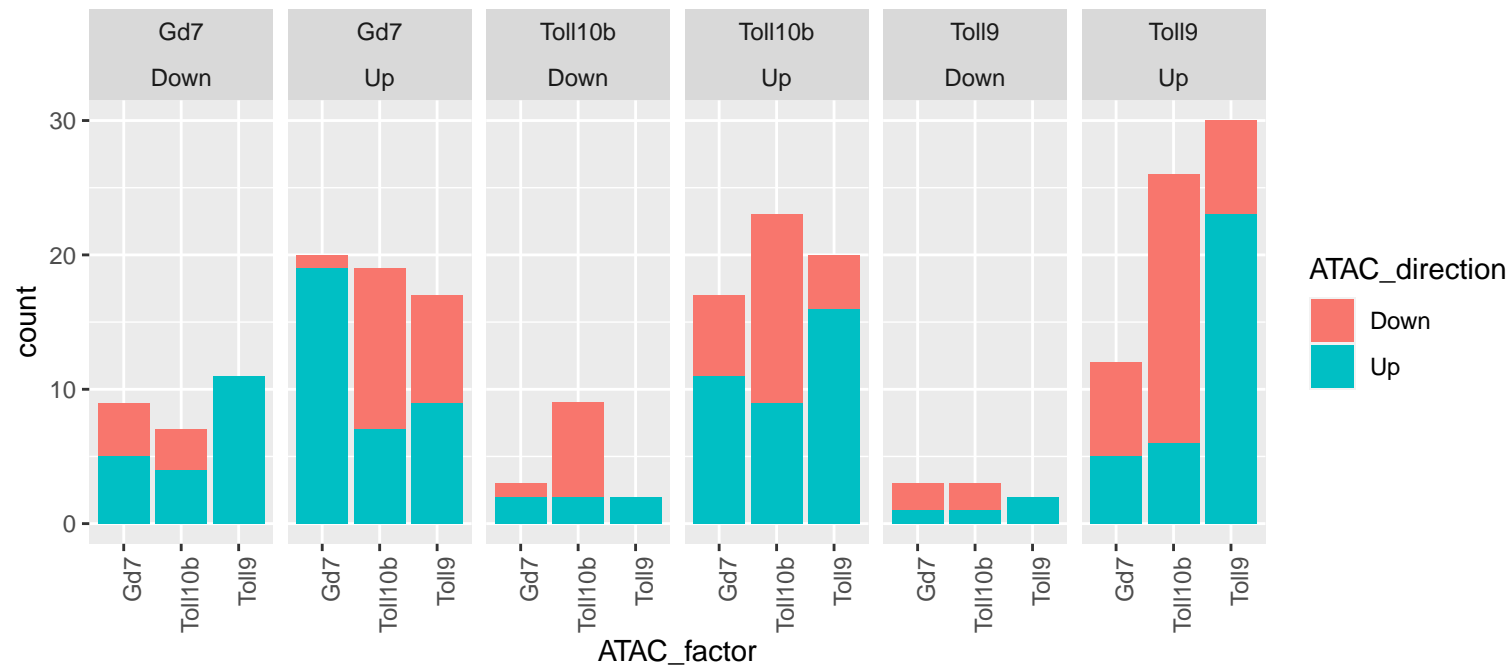


# ATACseq peak and PROseq Gene correlation

15 –100 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 15 –100 kb up– or downstream of gene