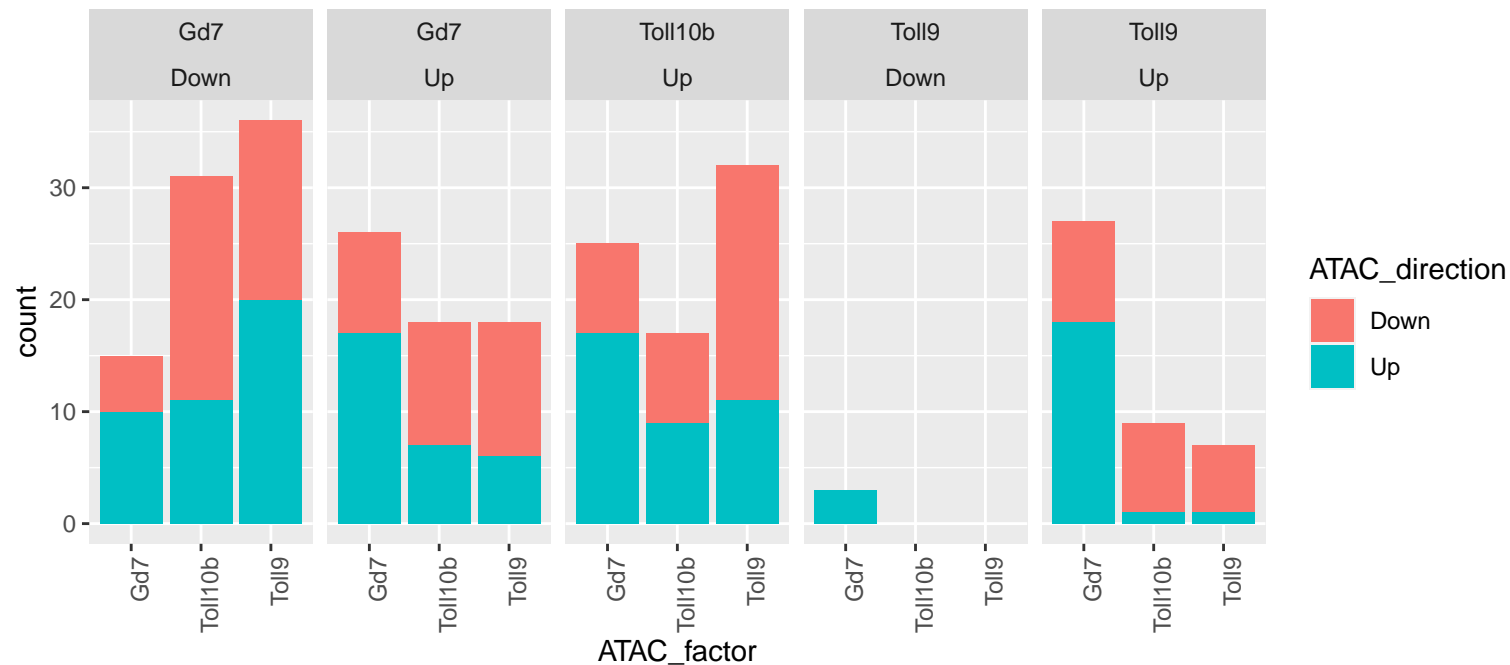


# ATACseq peak and PROseq Gene correlation

>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 >1000 kb up- or downstream of gene