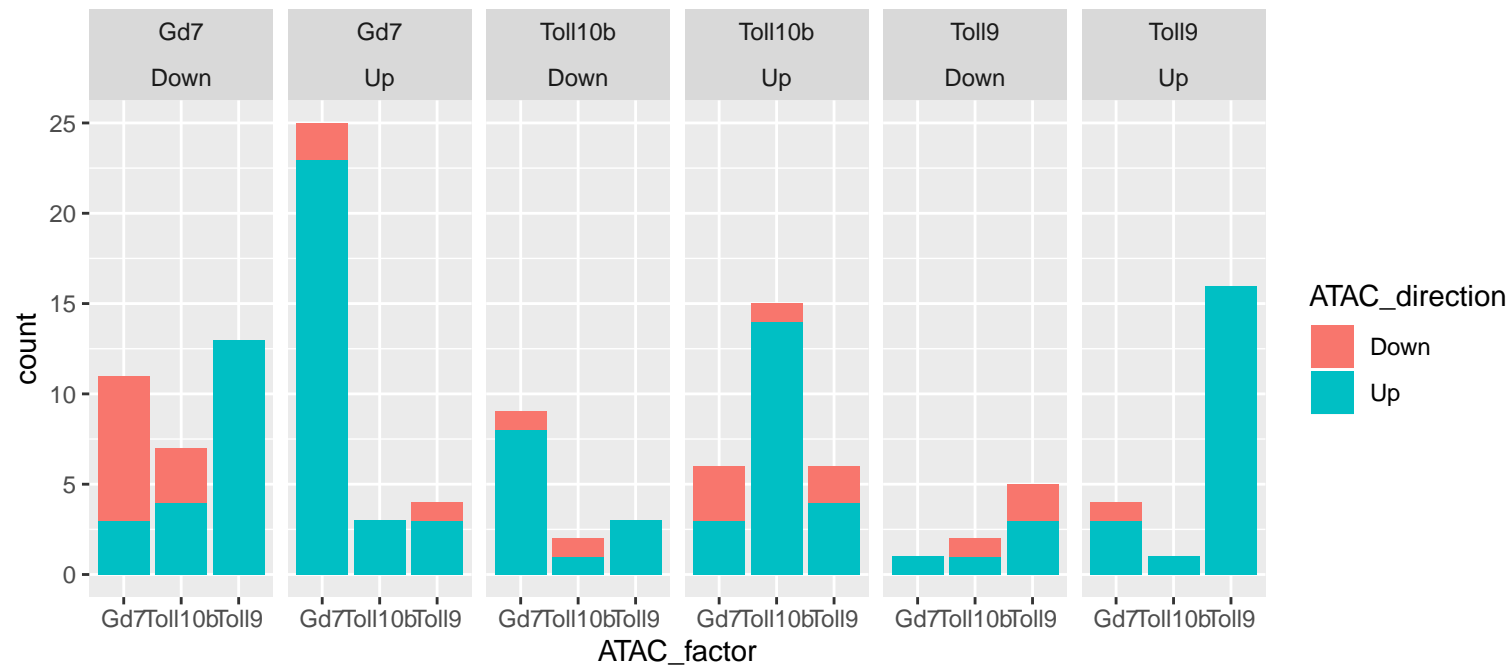


# Closest significant gene to ATAC peak

Within 15 kb



Gene is subset of genes that were identified using PROseq data  
Genes are divided based on PROseq analysis annotation  
Peak is located between 0 and 15 kb from gene boundaries