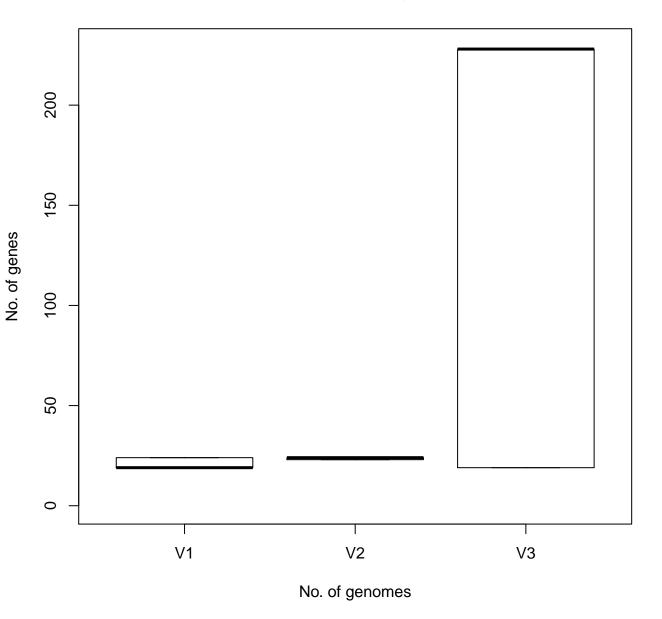
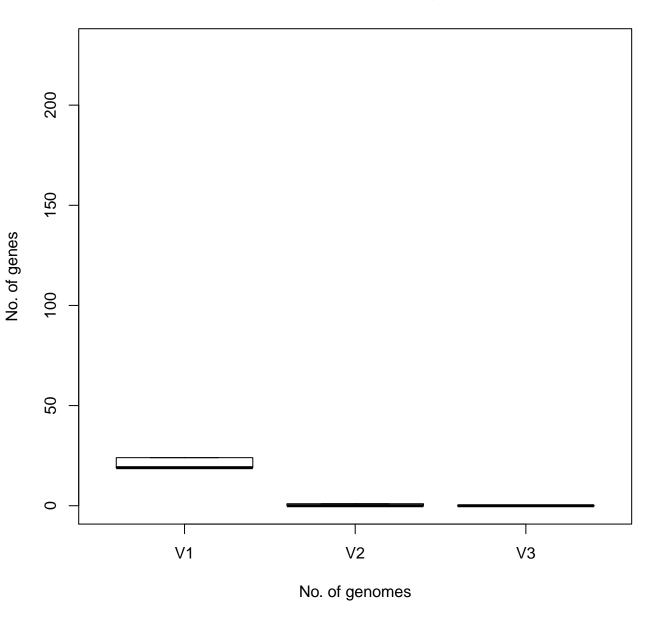
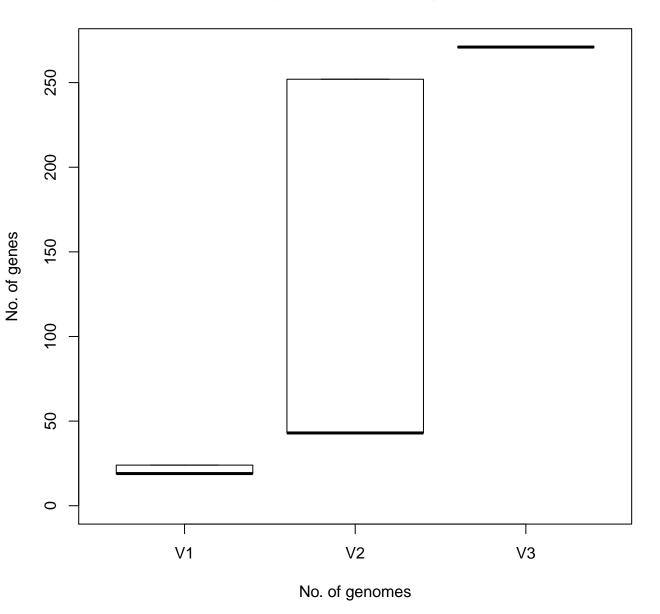
### Number of new genes



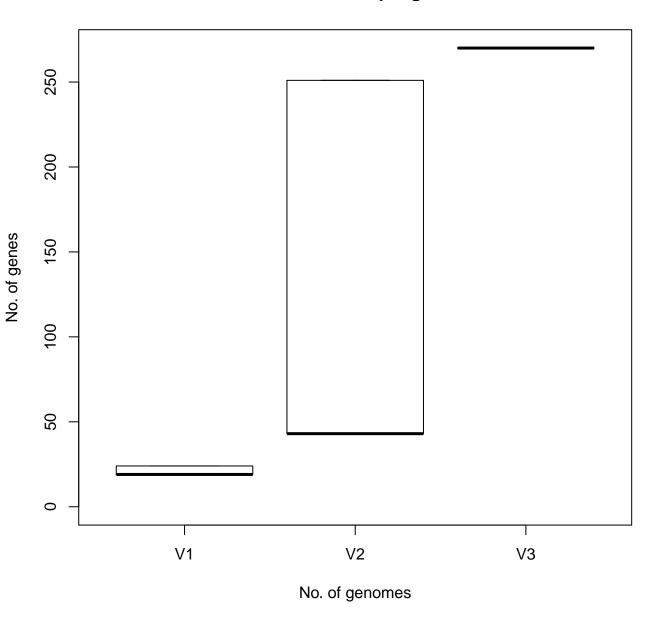
### Number of conserved genes



## No. of genes in the pan-genome



# Number of unique genes



#### Number of blastp hits with different percentage identity

