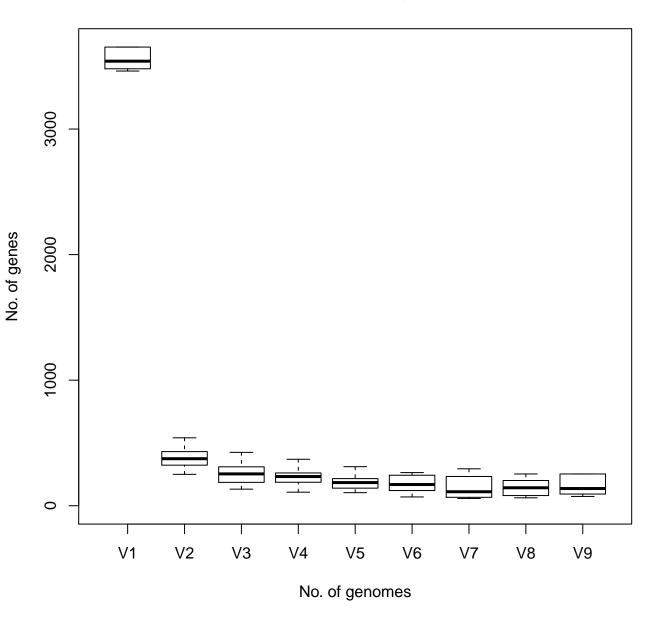
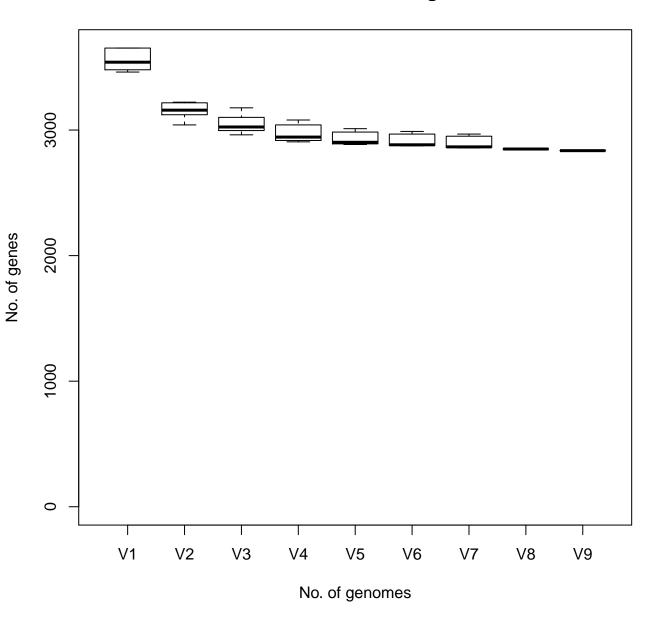
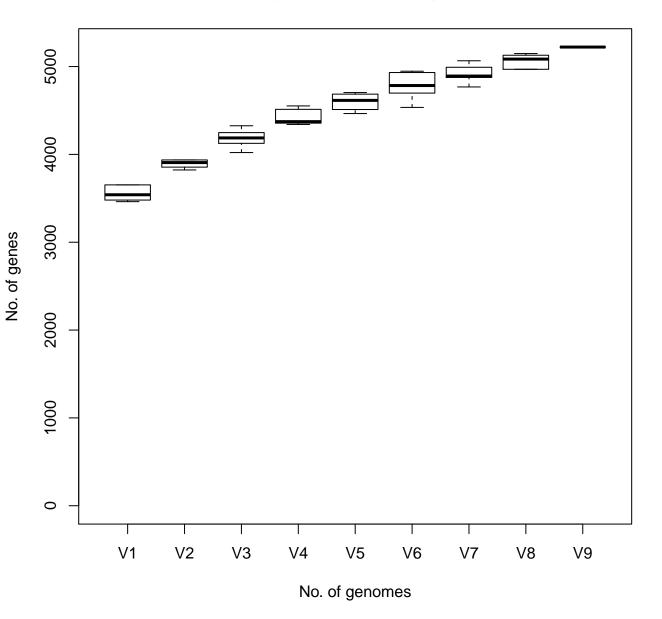
#### Number of new genes



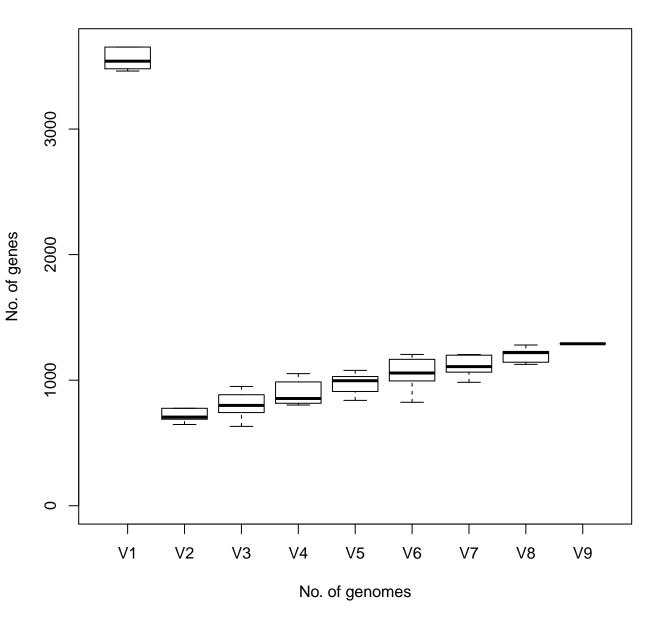
### Number of conserved genes



# No. of genes in the pan-genome



# Number of unique genes



### Number of blastp hits with different percentage identity

