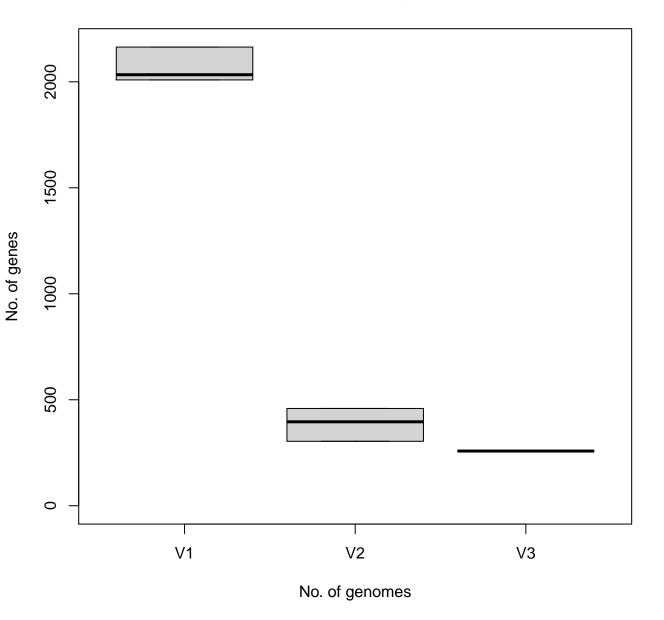
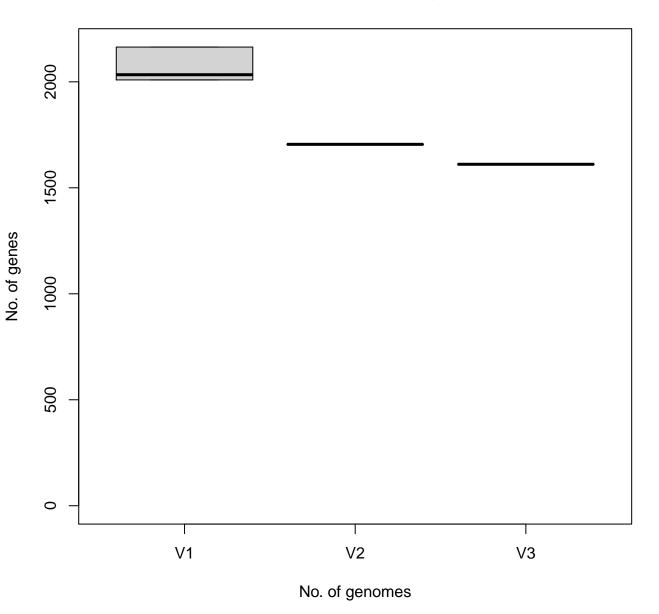
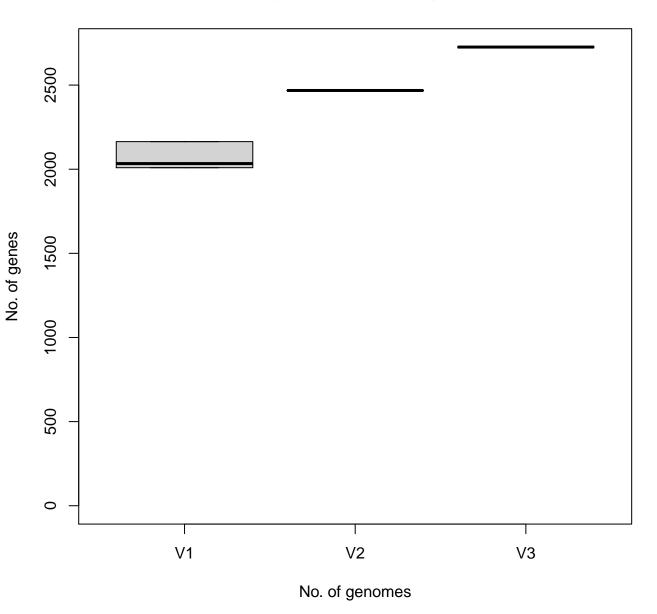
# Number of new genes



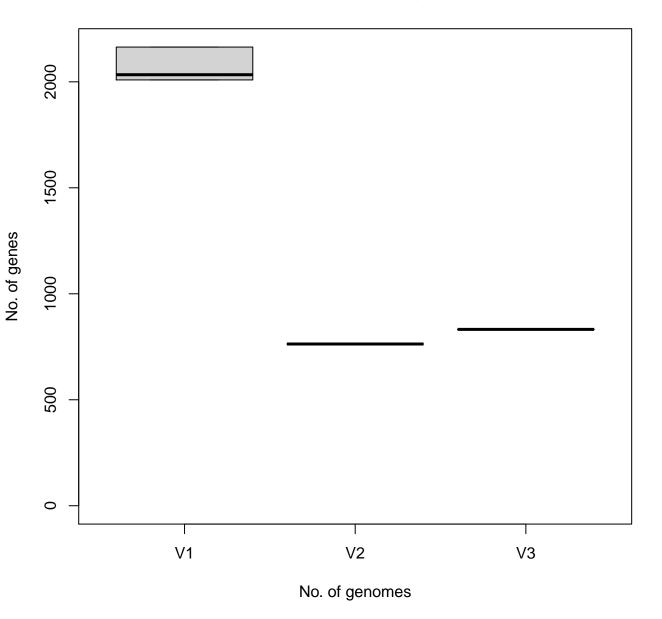
#### Number of conserved genes



# No. of genes in the pan-genome



### Number of unique genes



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

