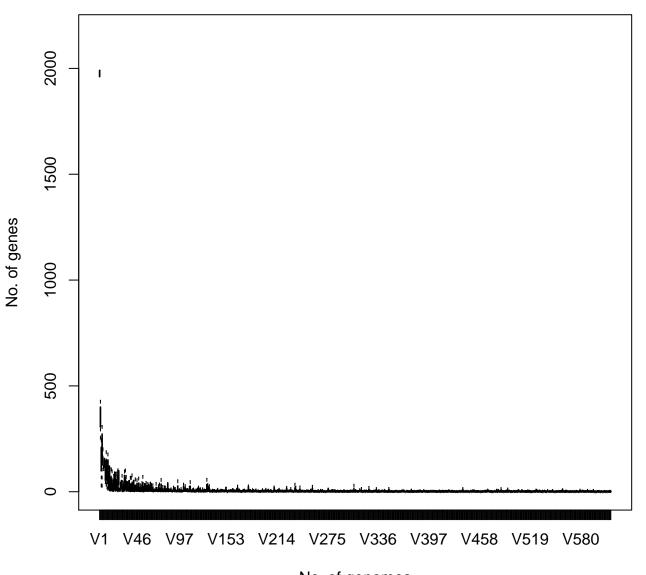
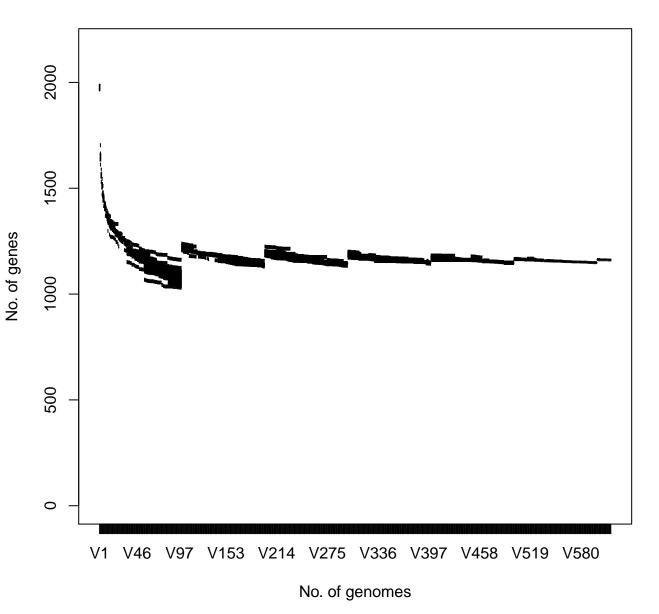
## Number of new genes

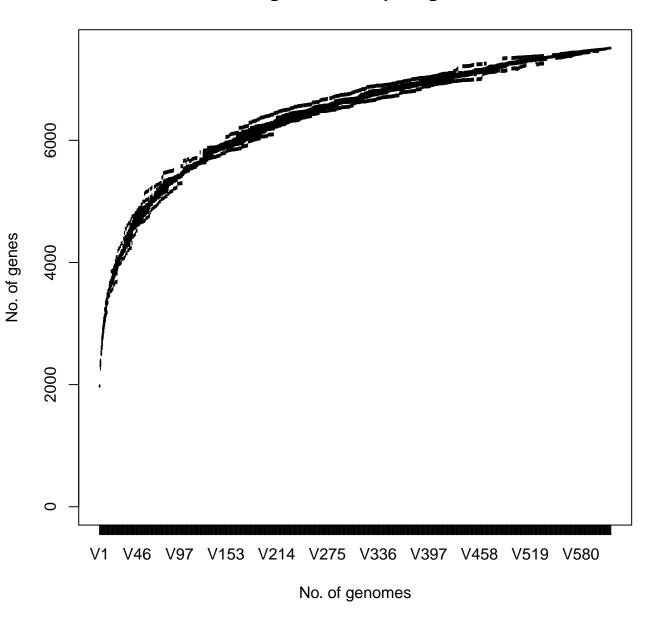


No. of genomes

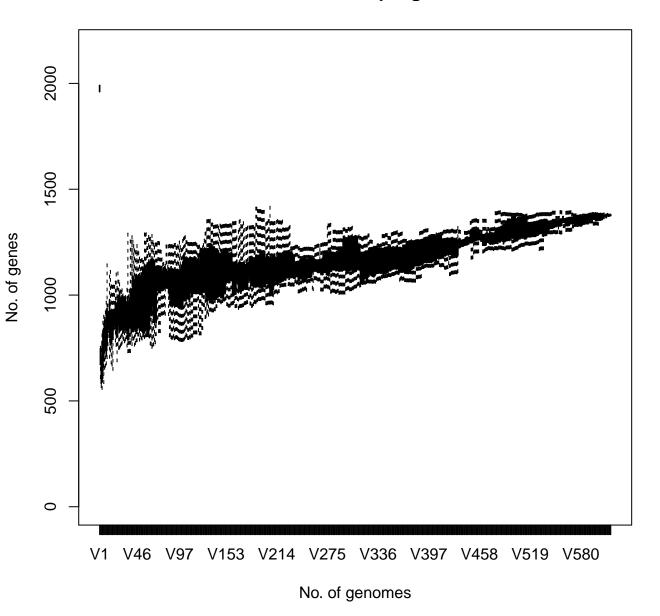
#### Number of conserved genes



#### No. of genes in the pan-genome



# Number of unique genes



### Number of blastp hits with different percentage identity

