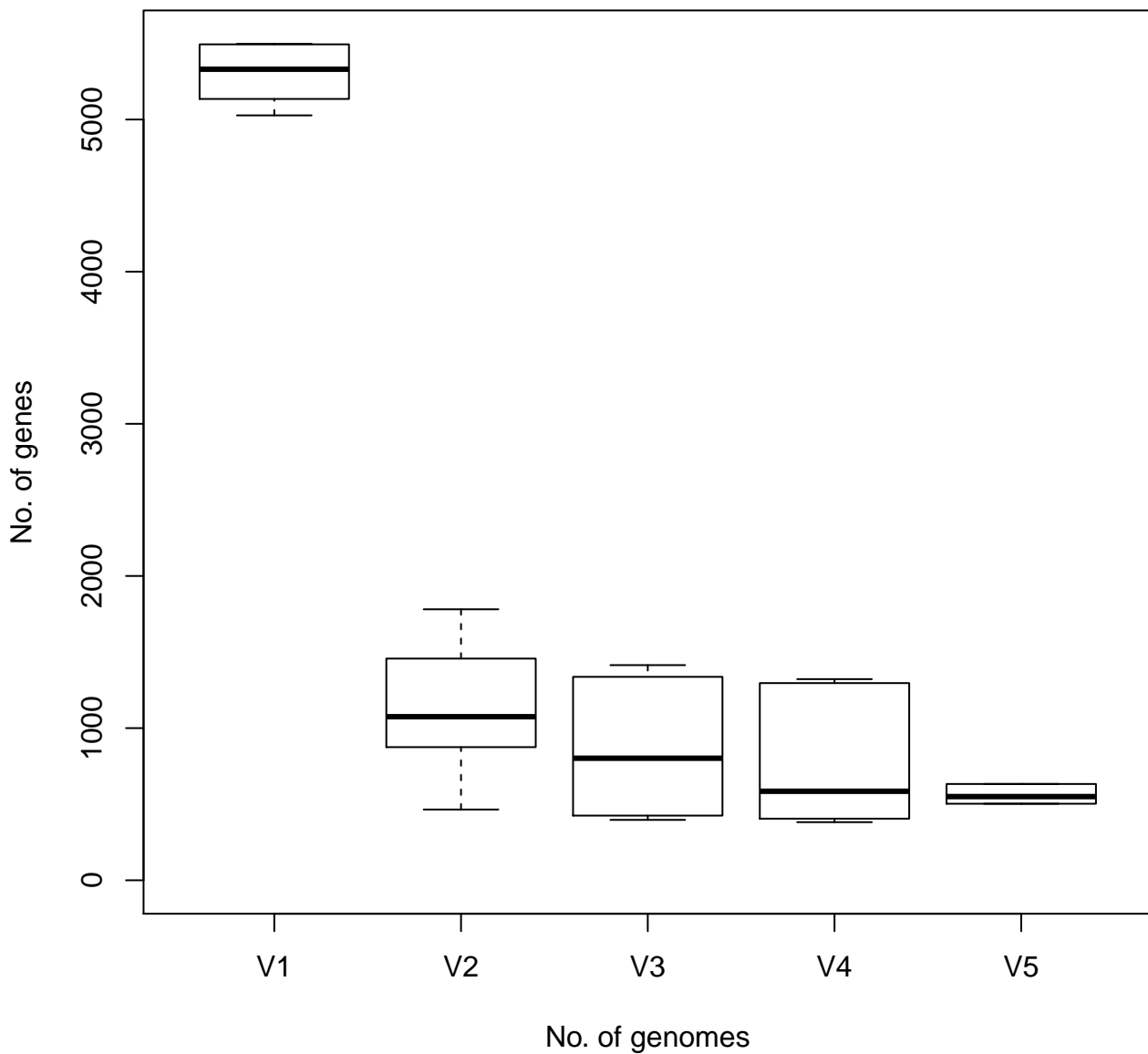
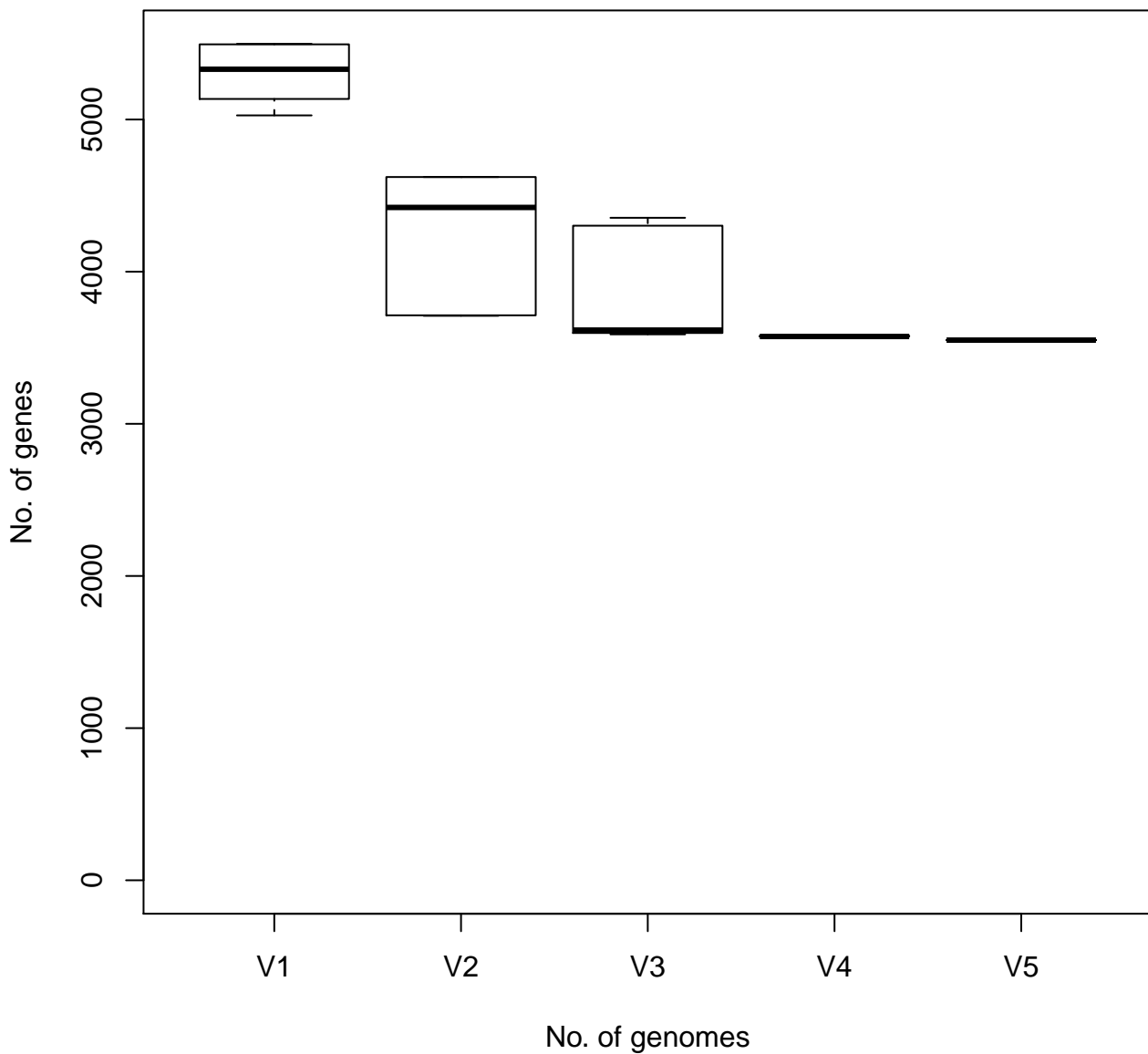


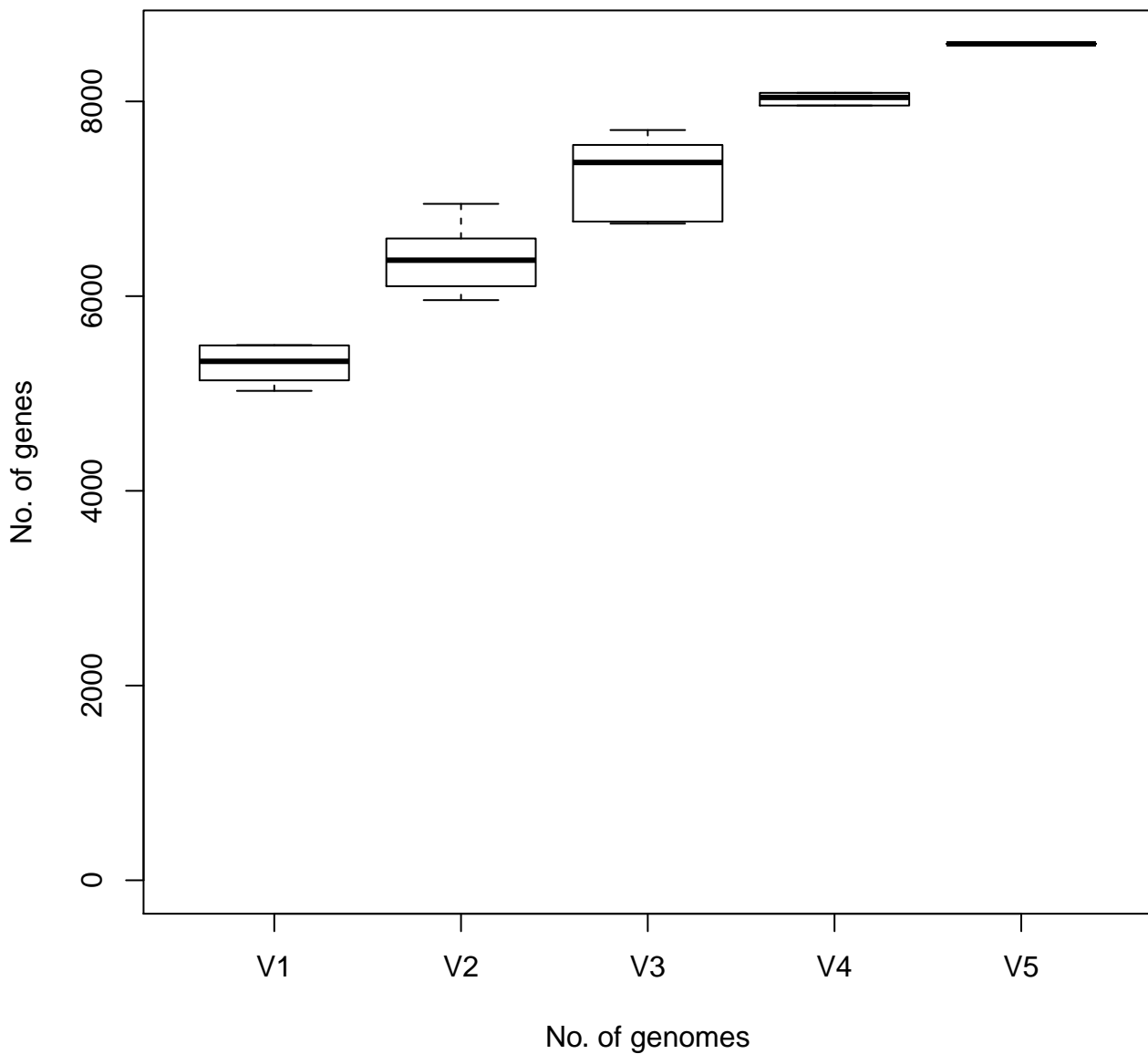
## Number of new genes



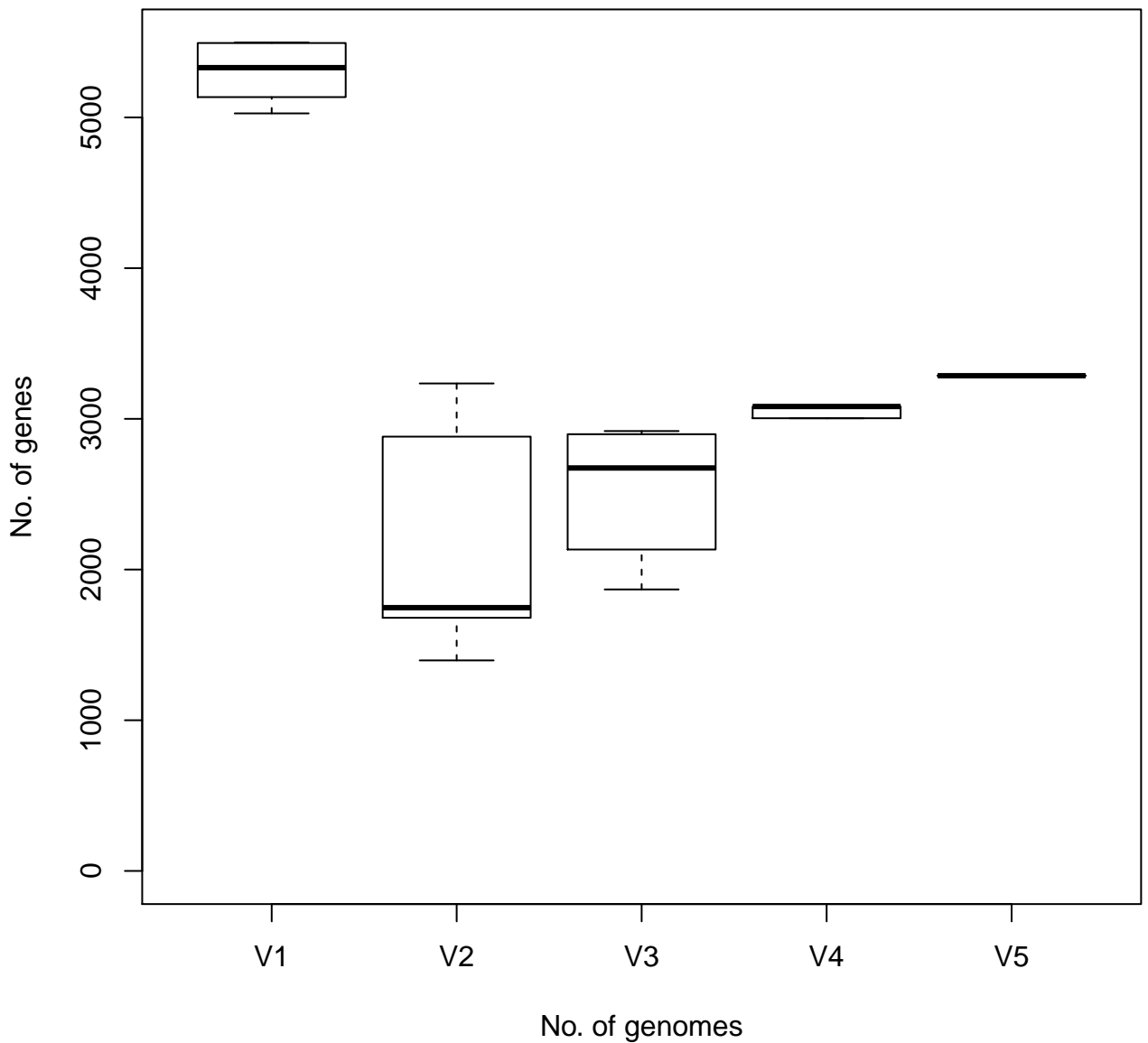
## Number of conserved genes



## No. of genes in the pan-genome



# Number of unique genes



# Number of blastp hits with different percentage identity

