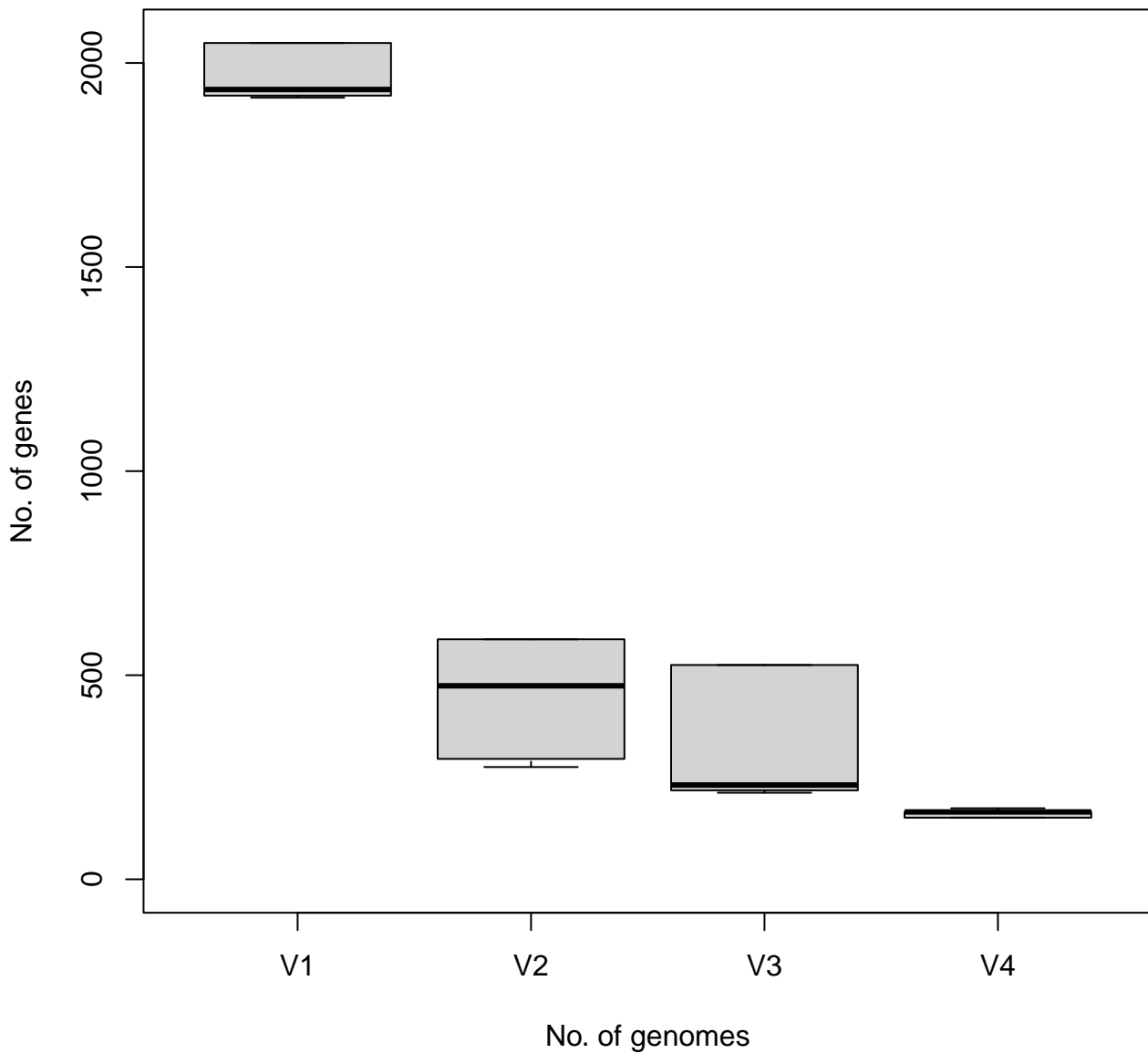
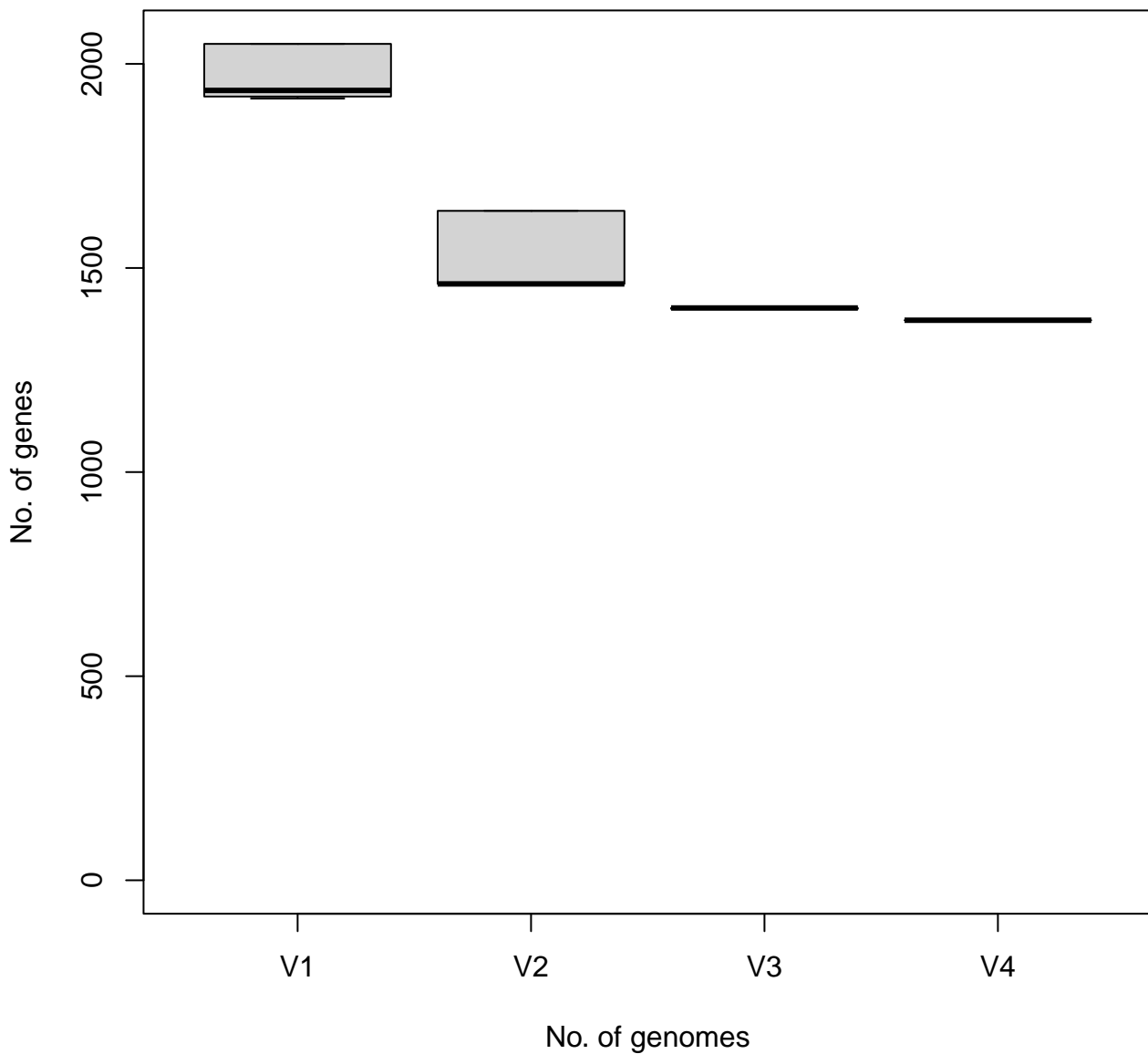


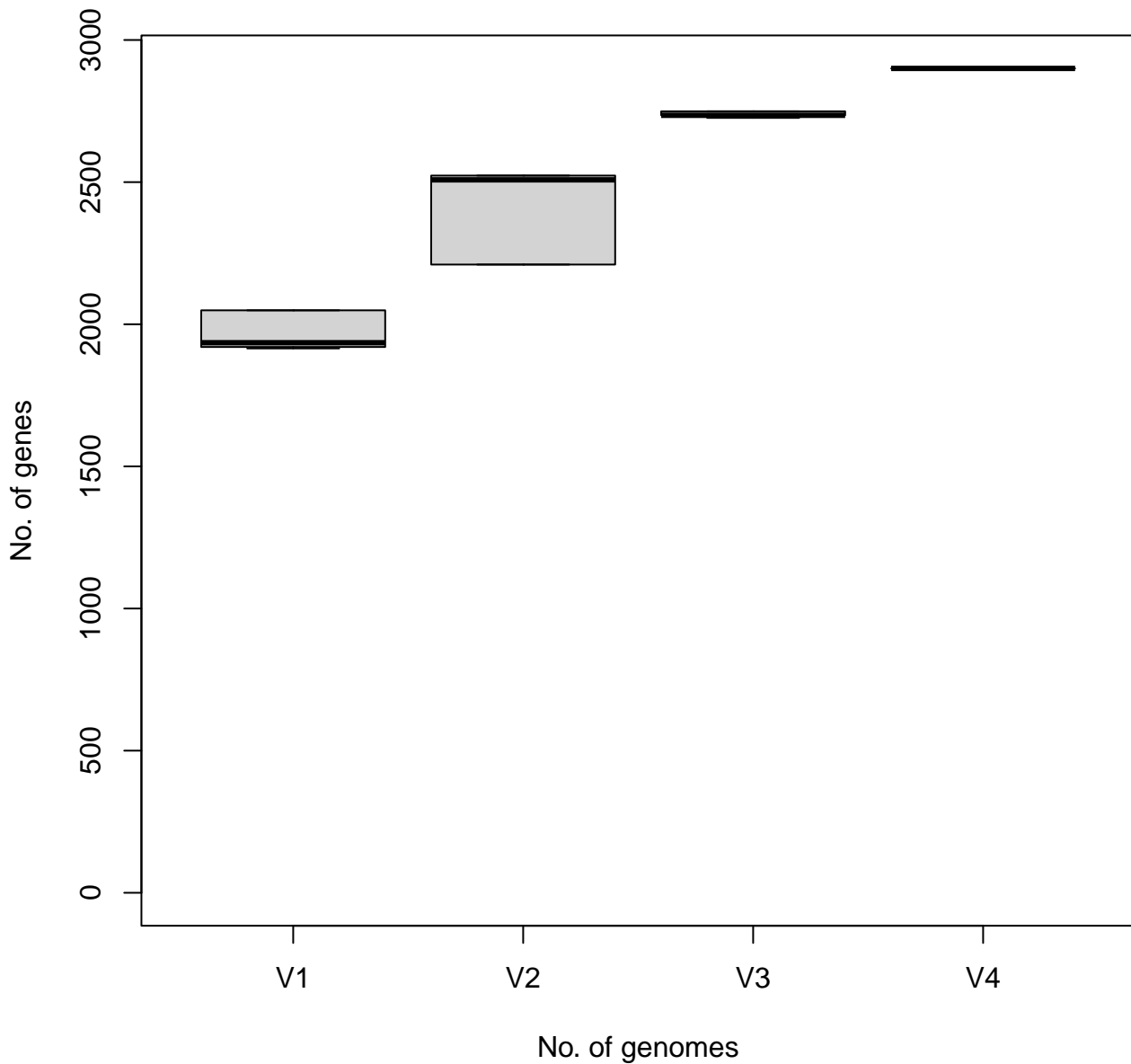
## Number of new genes



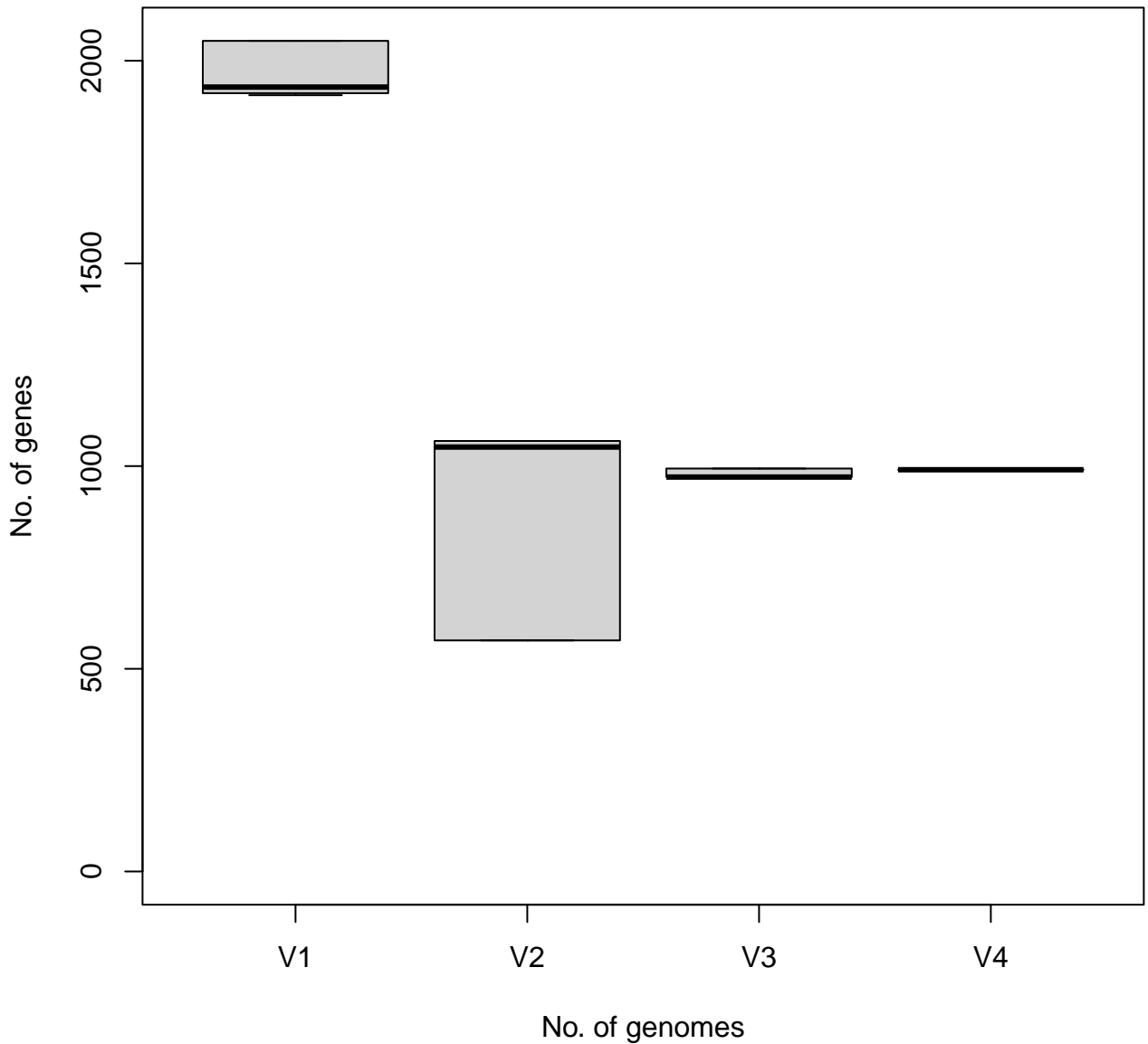
# Number of conserved genes



## No. of genes in the pan-genome



# Number of unique genes



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

