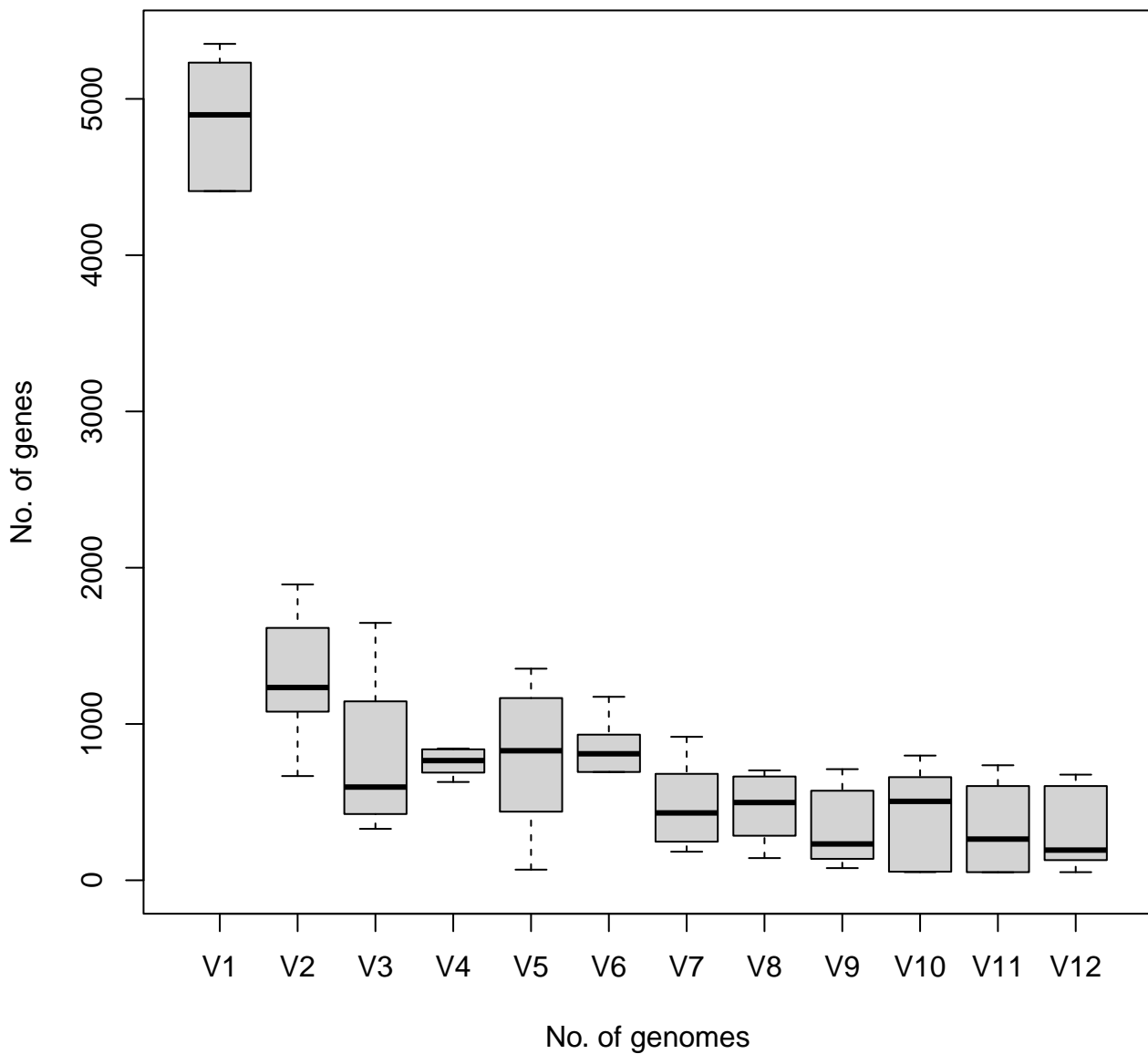
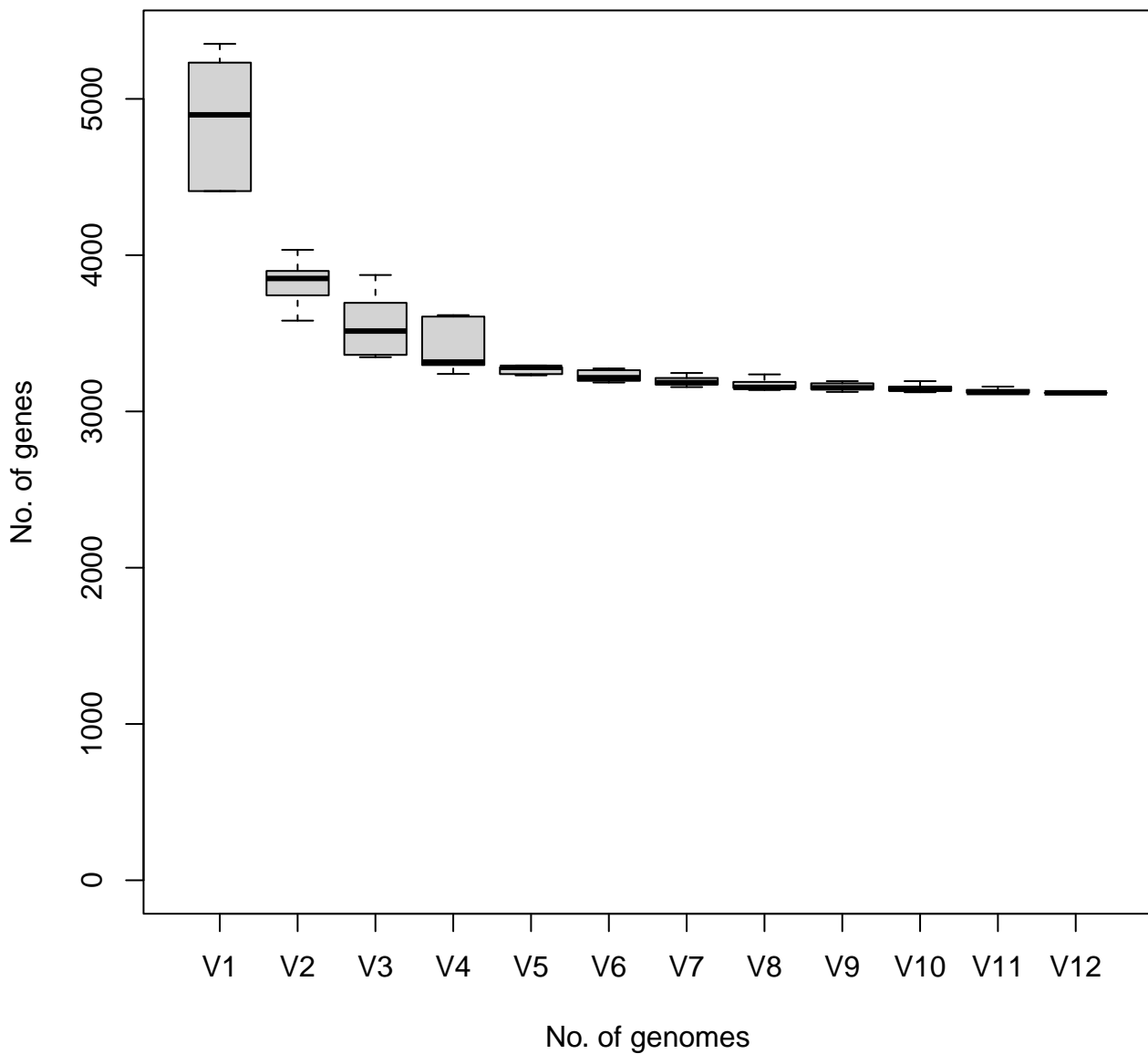


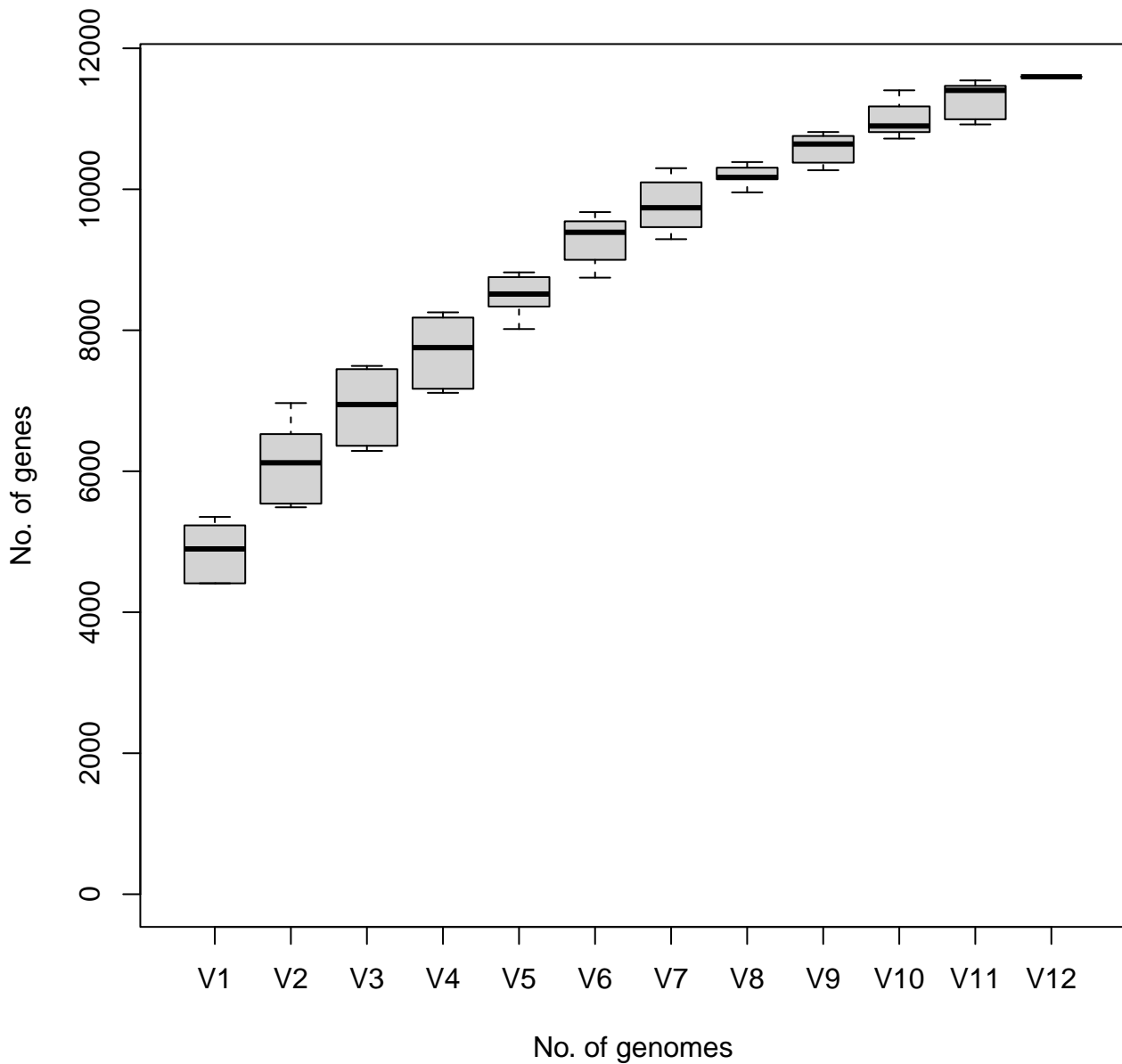
## Number of new genes



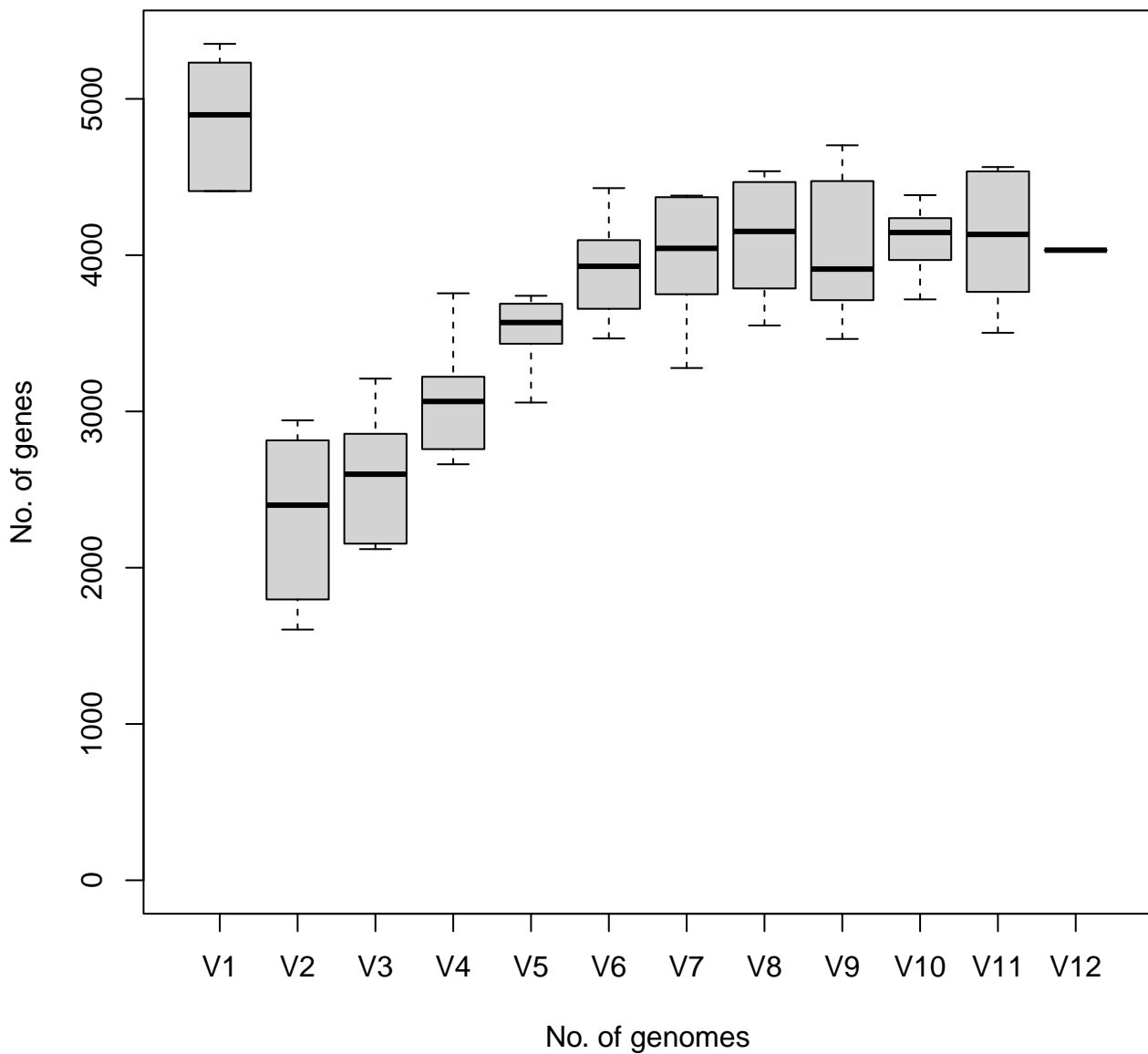
# Number of conserved genes



## No. of genes in the pan-genome



## Number of unique genes



# Number of blastp hits with different percentage identity

