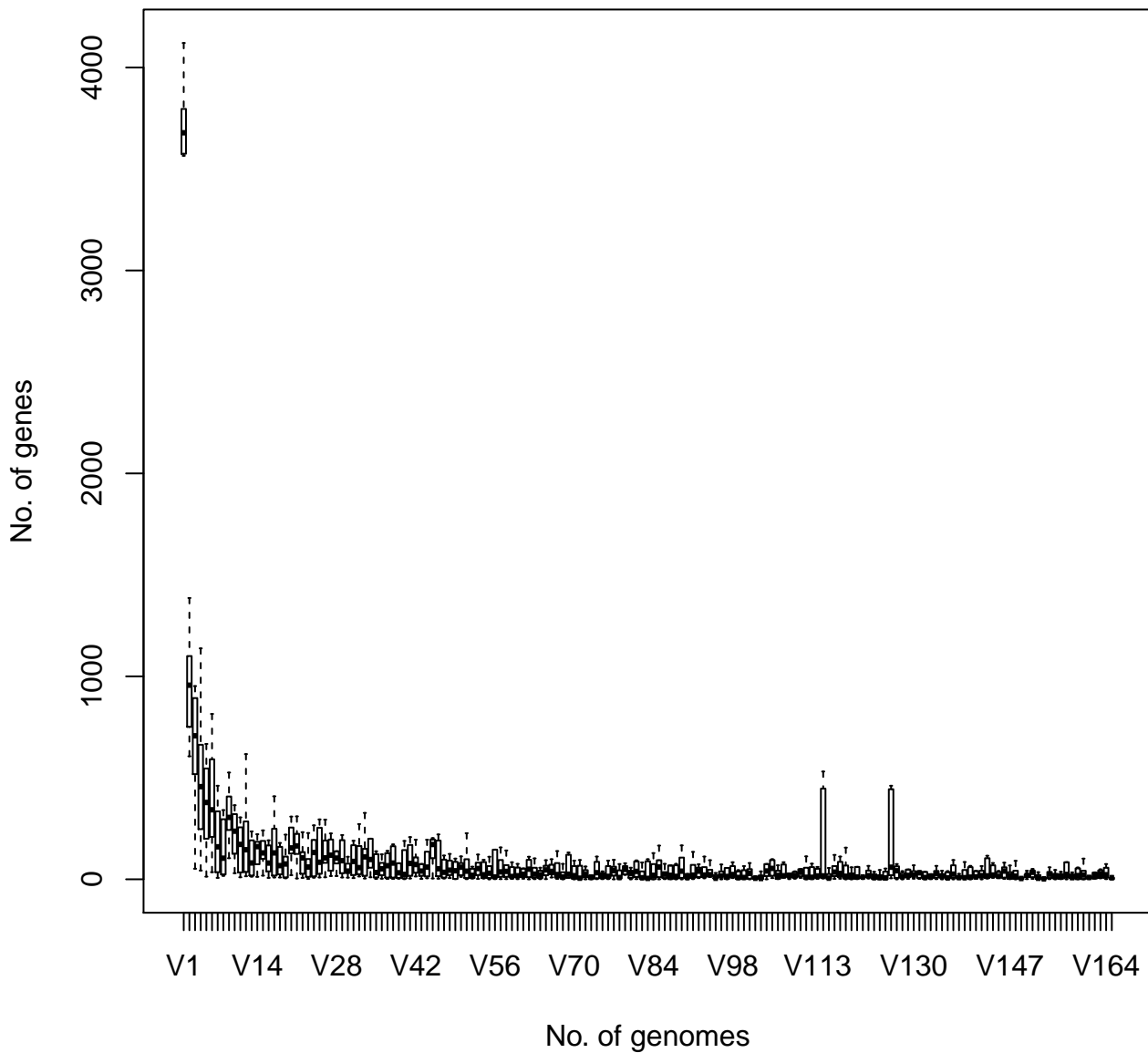
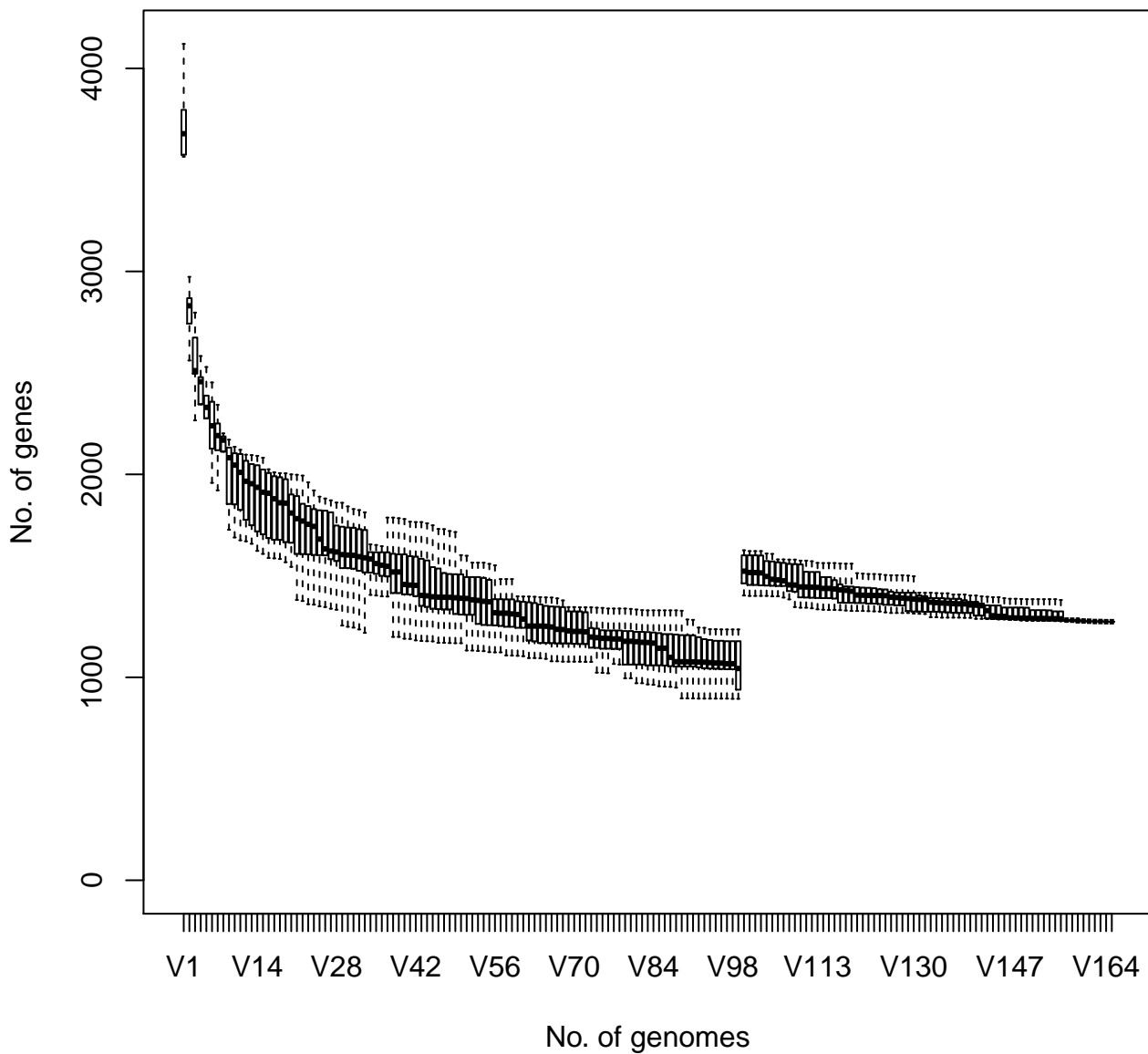


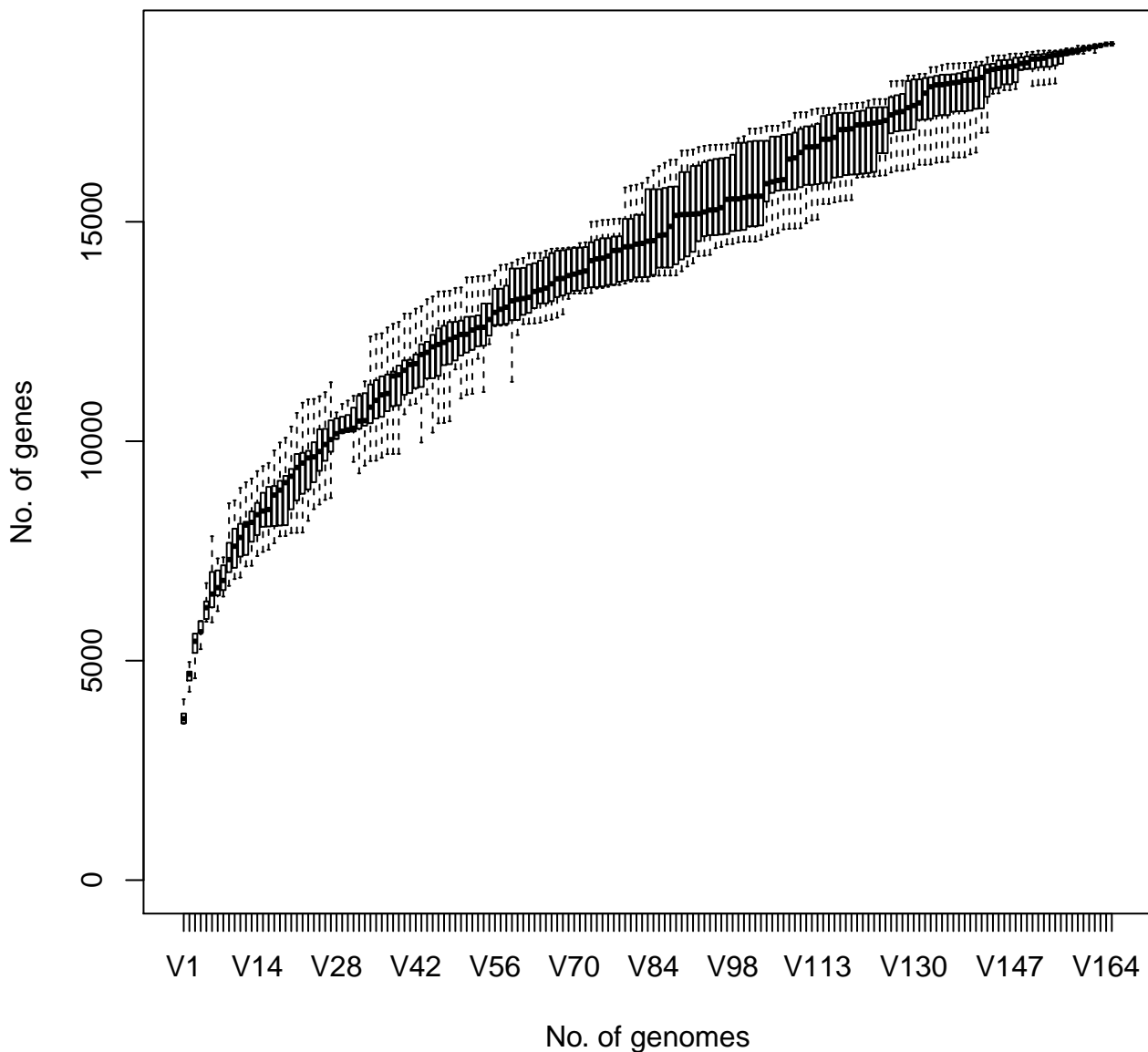
# Number of new genes



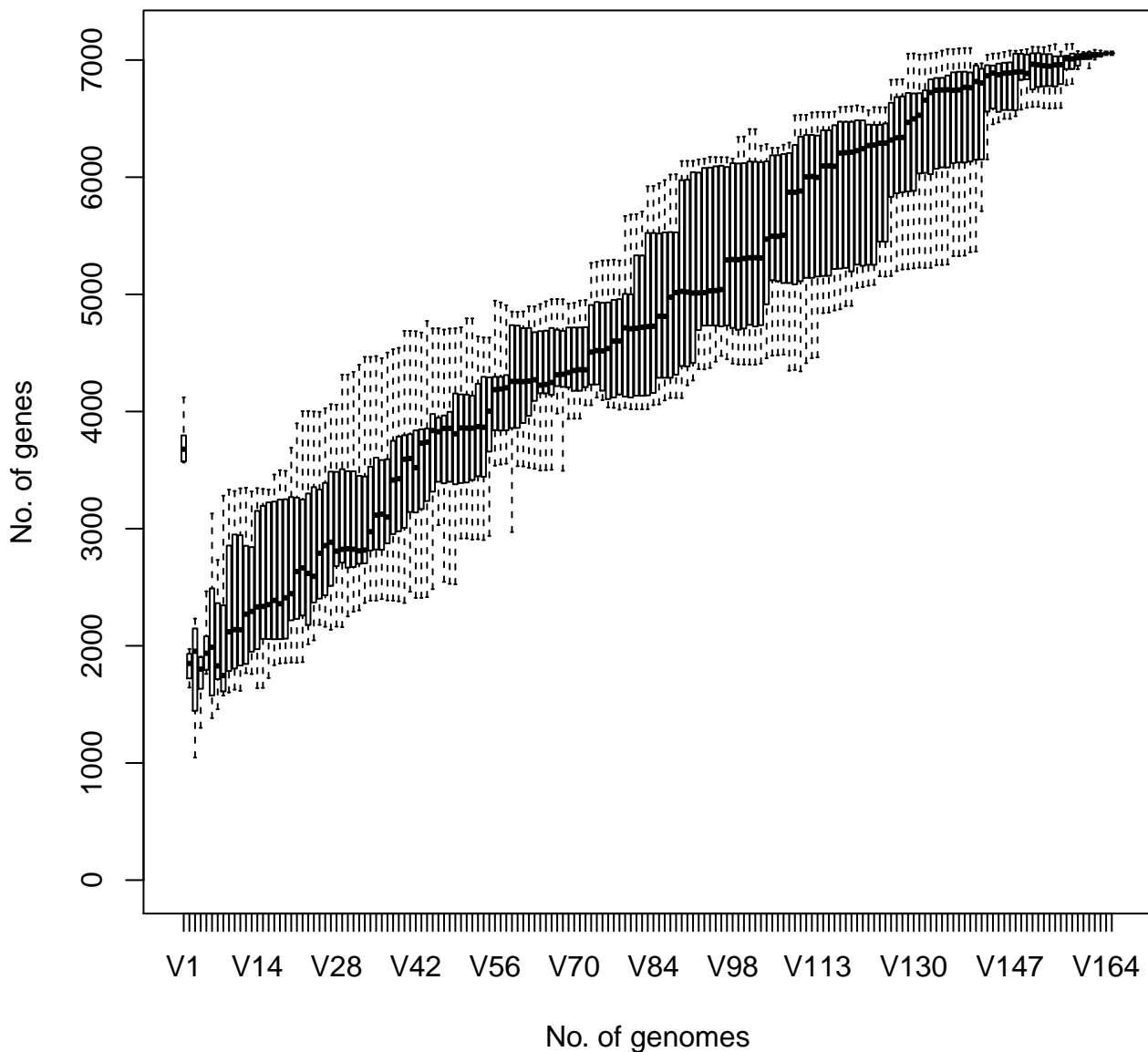
# Number of conserved genes



# No. of genes in the pan-genome



# Number of unique genes



# Number of blastp hits with different percentage identity

