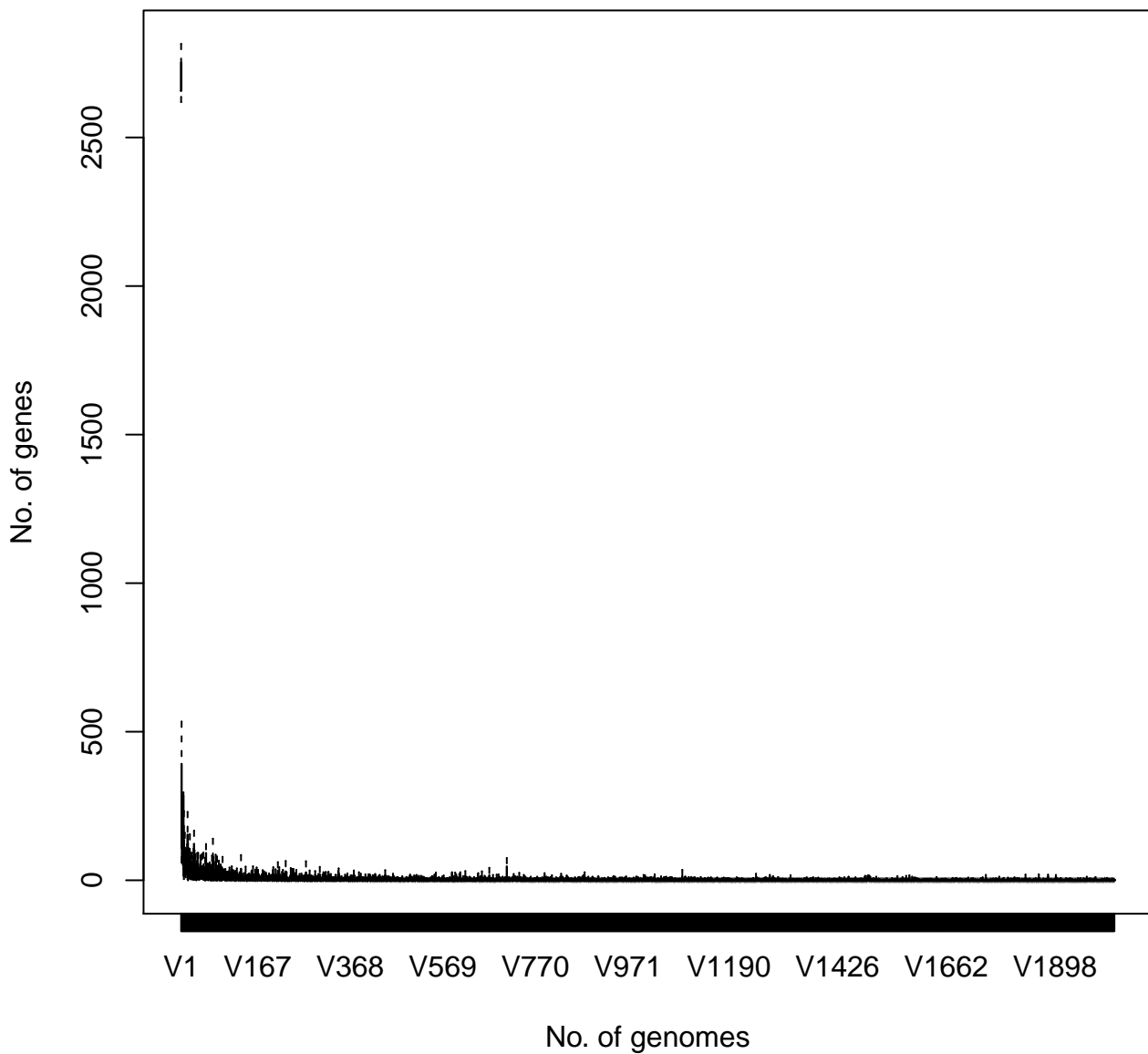
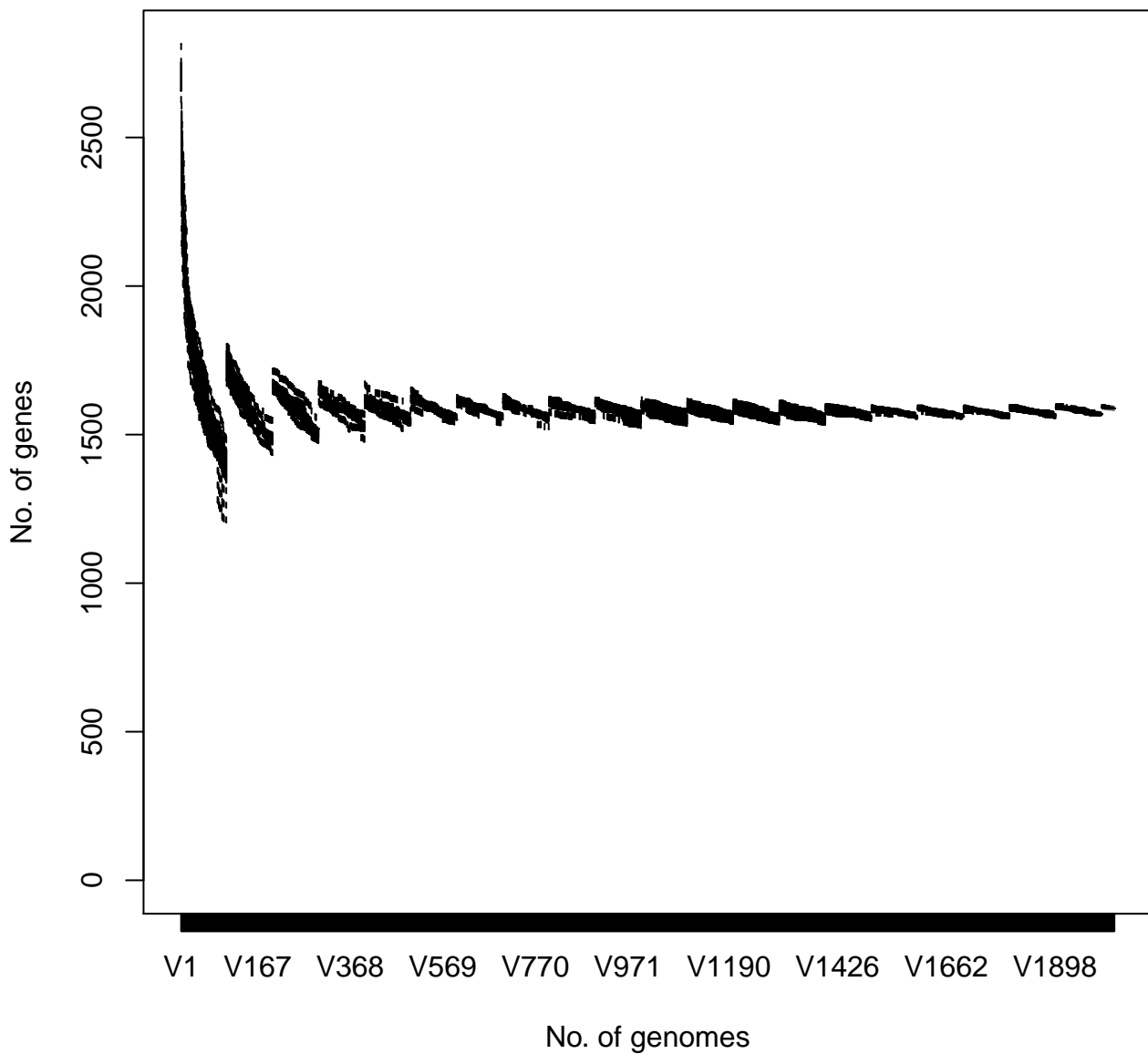


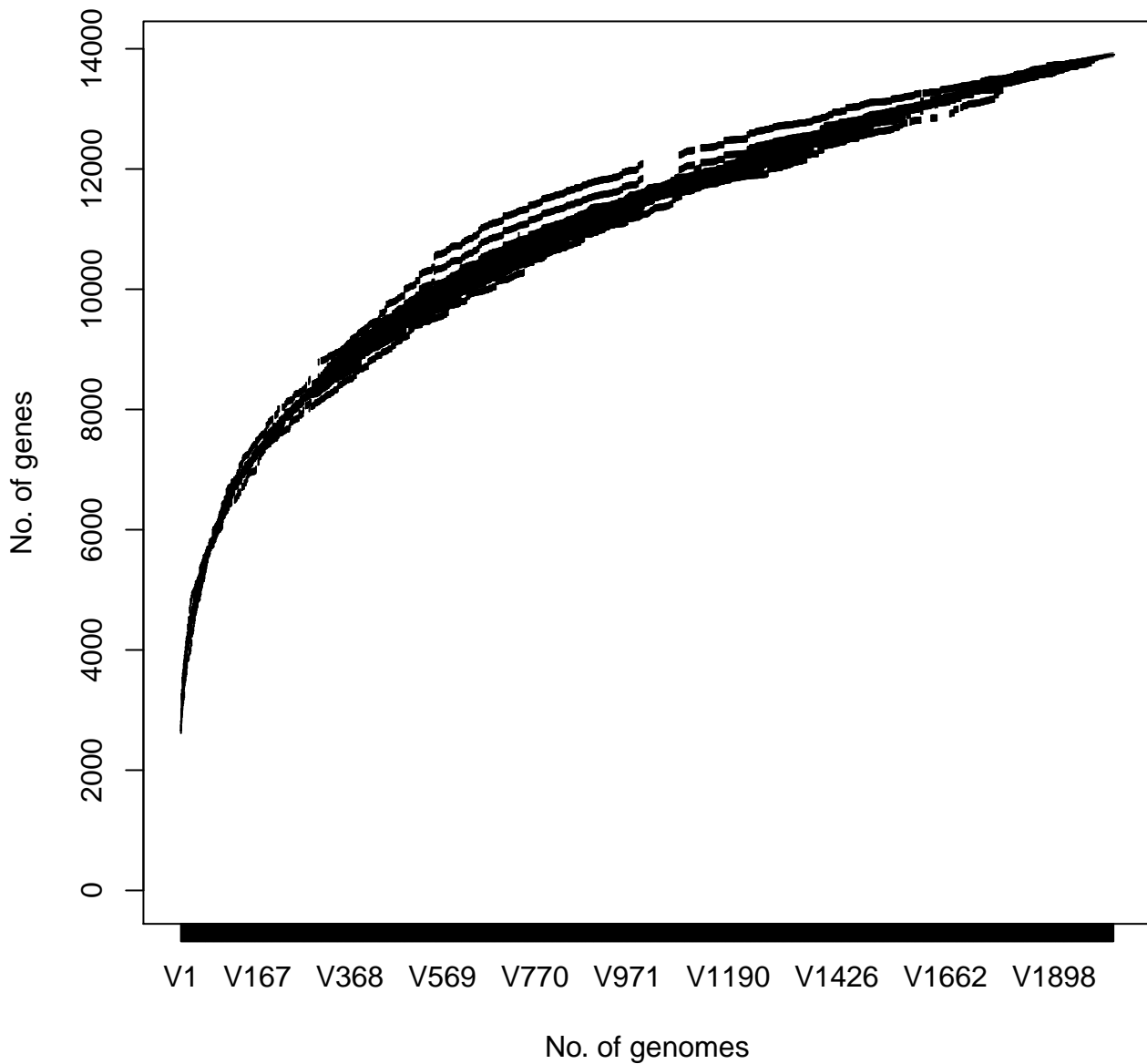
# Number of new genes



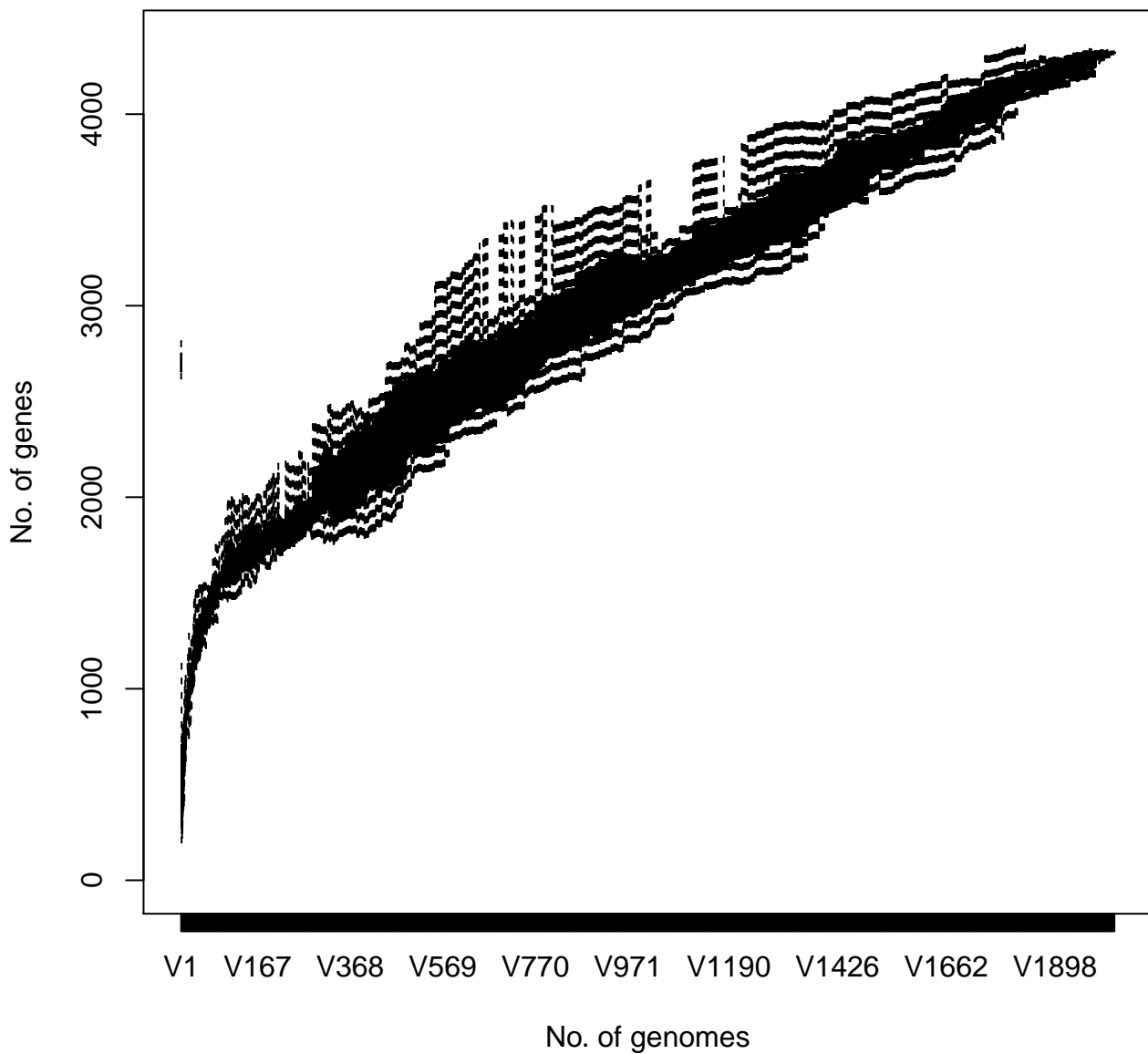
# Number of conserved genes



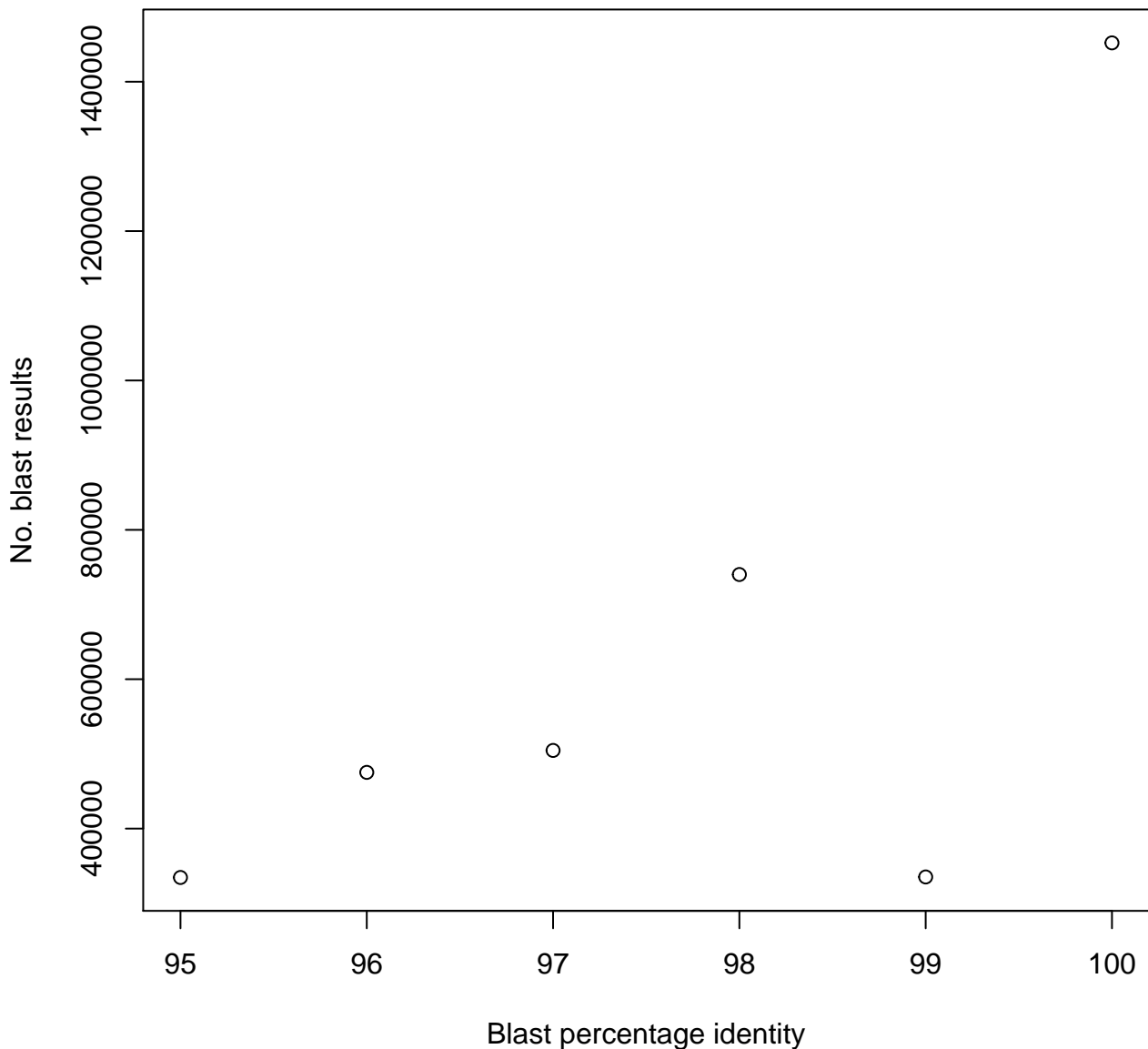
## No. of genes in the pan-genome



# Number of unique genes



# Number of blastp hits with different percentage identity



# Key

- Conserved genes
- .... Total genes

