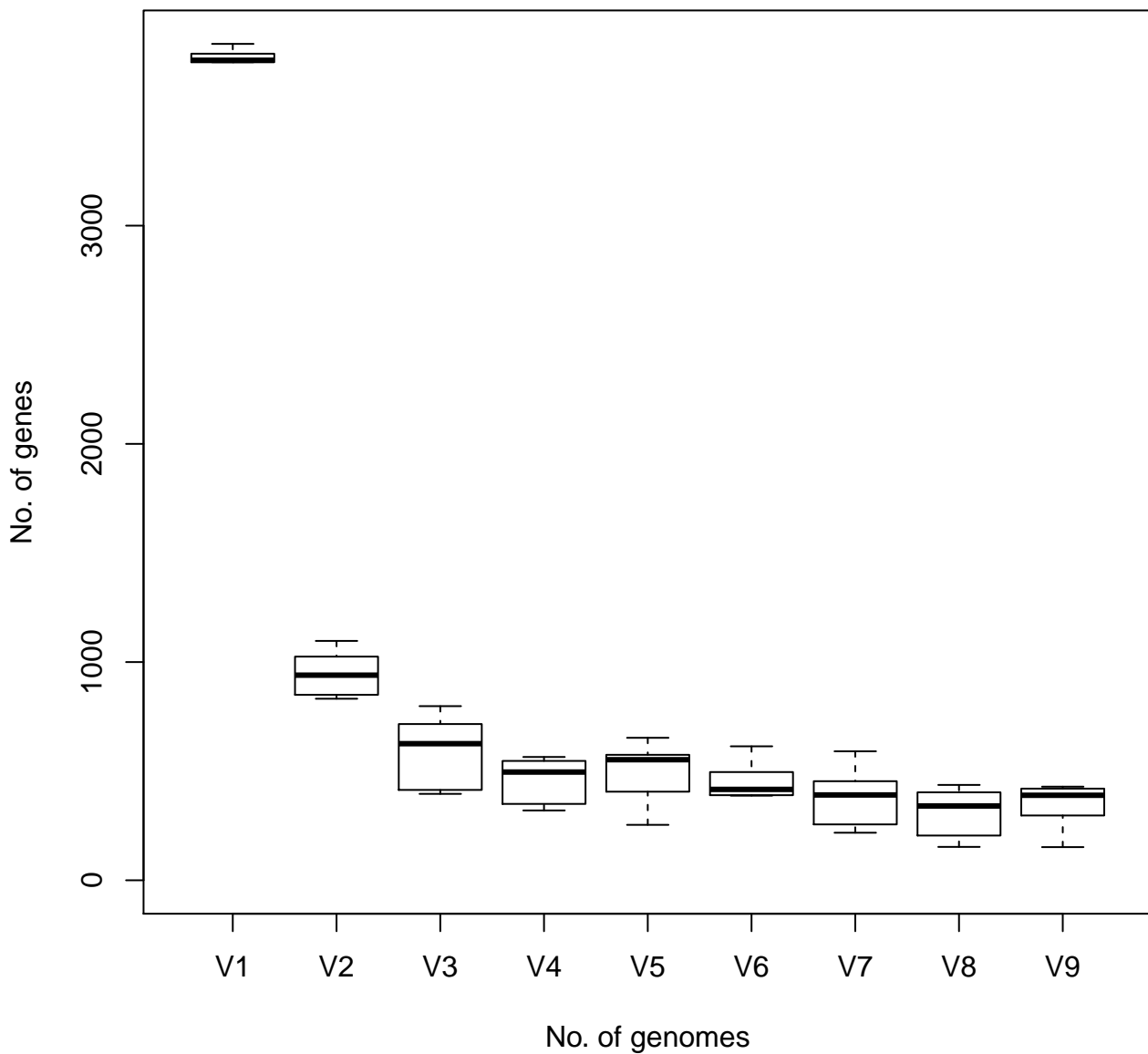
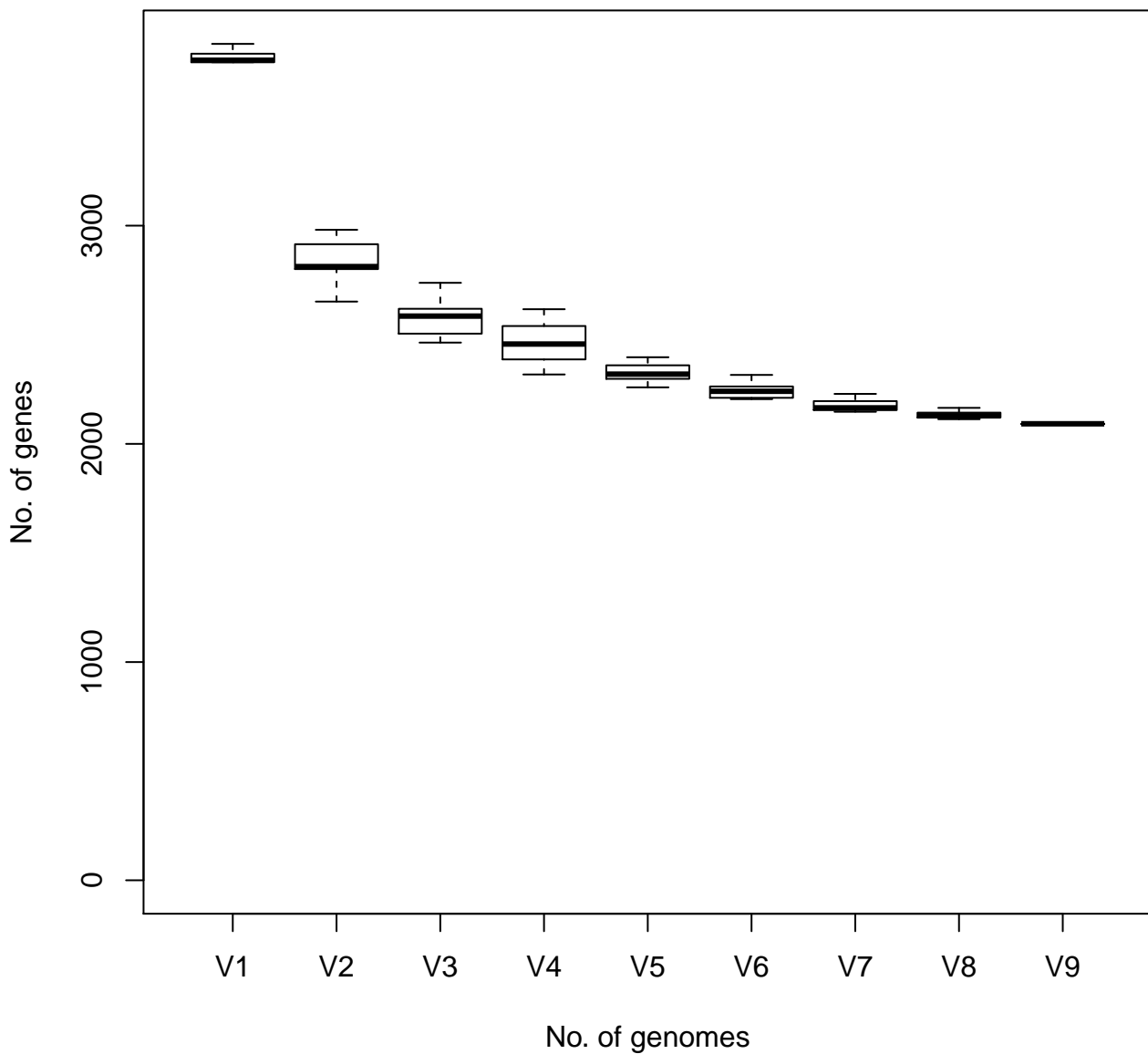


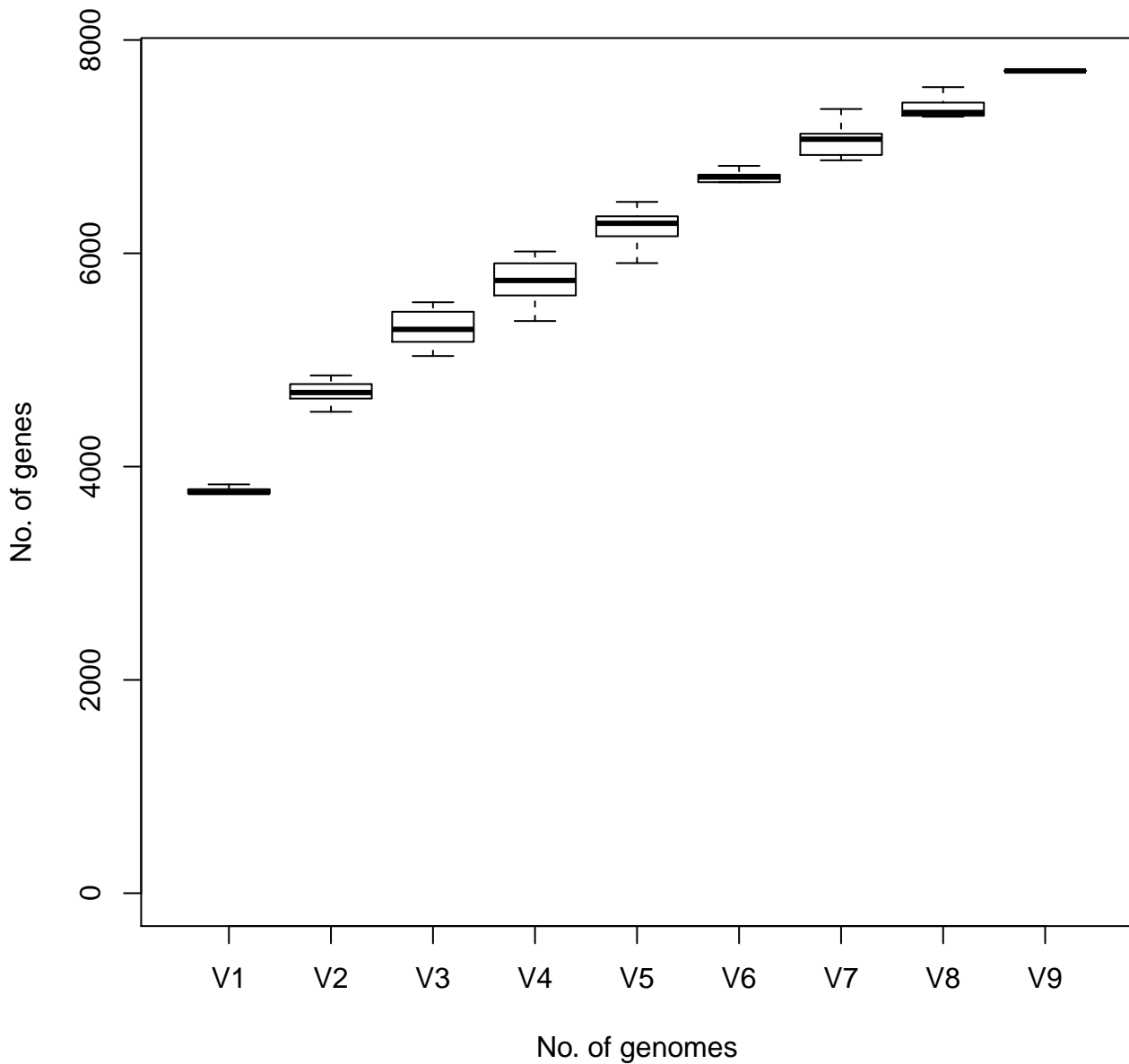
Number of new genes



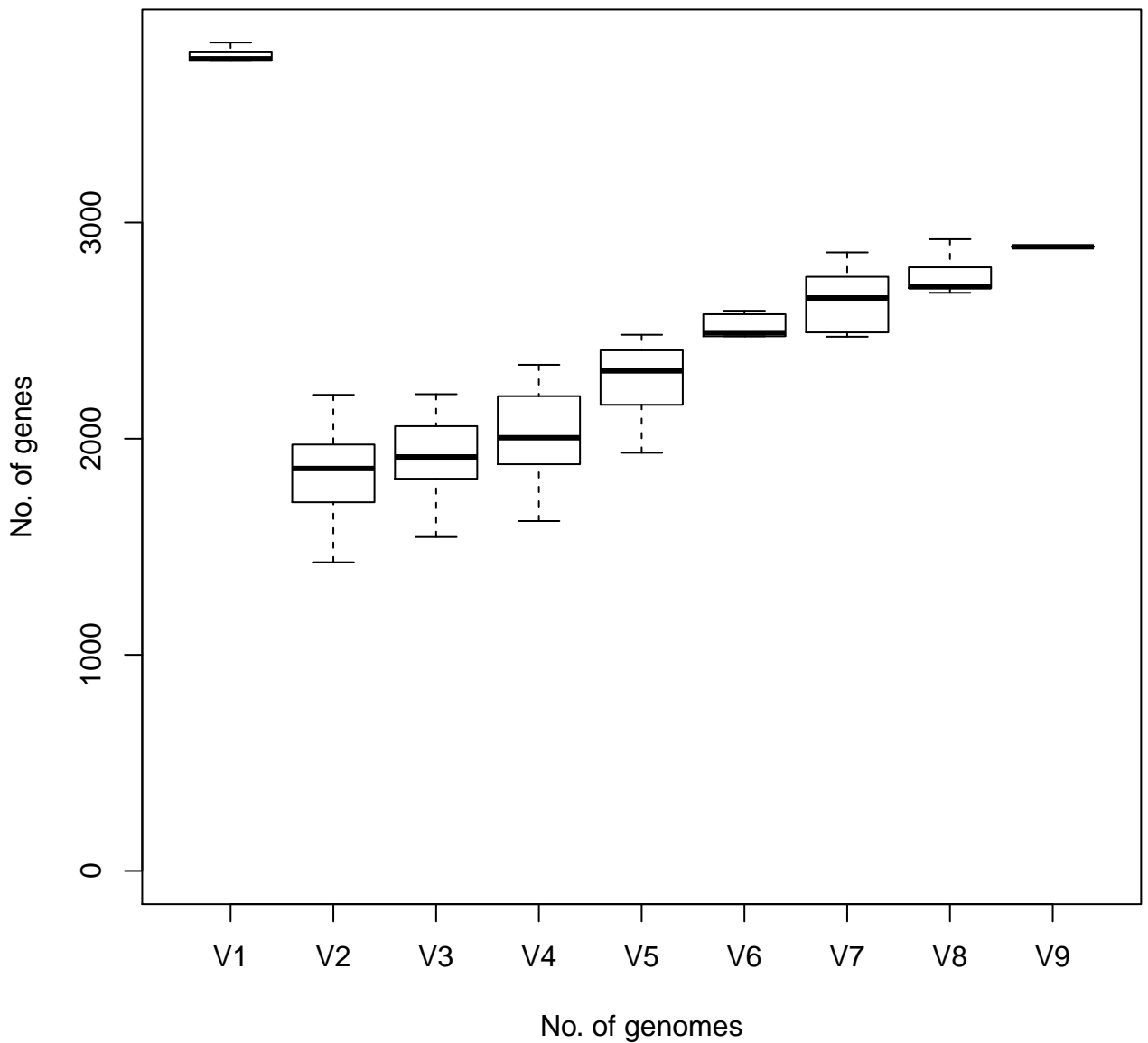
Number of conserved genes



No. of genes in the pan-genome



Number of unique genes



Number of blastp hits with different percentage identity

