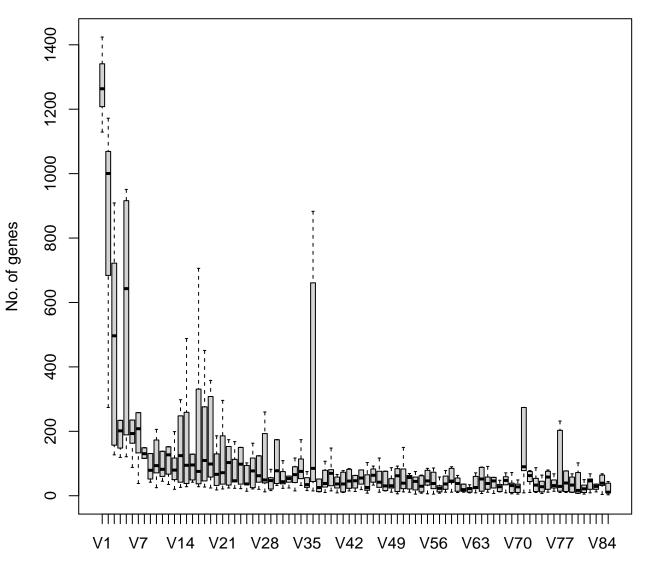
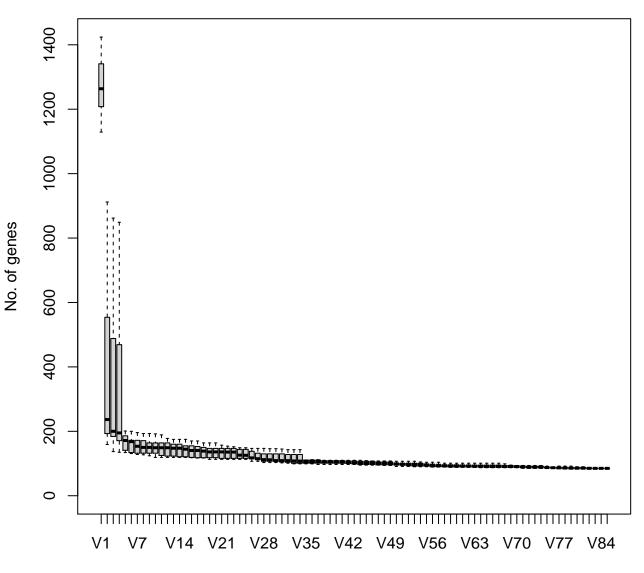
## Number of new genes



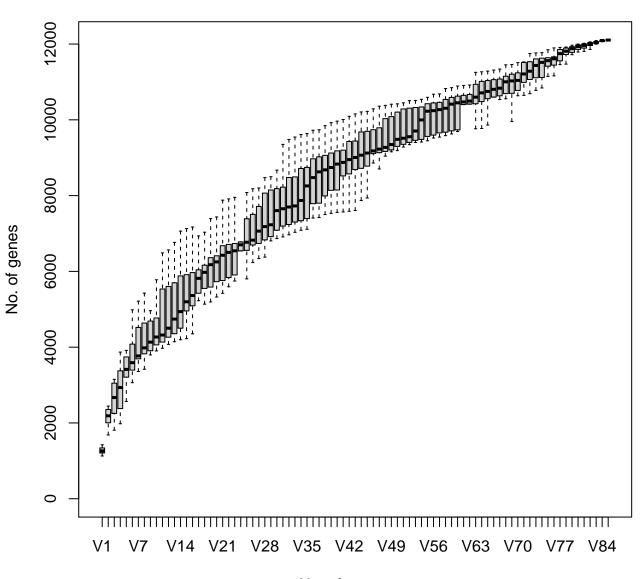
No. of genomes

### Number of conserved genes



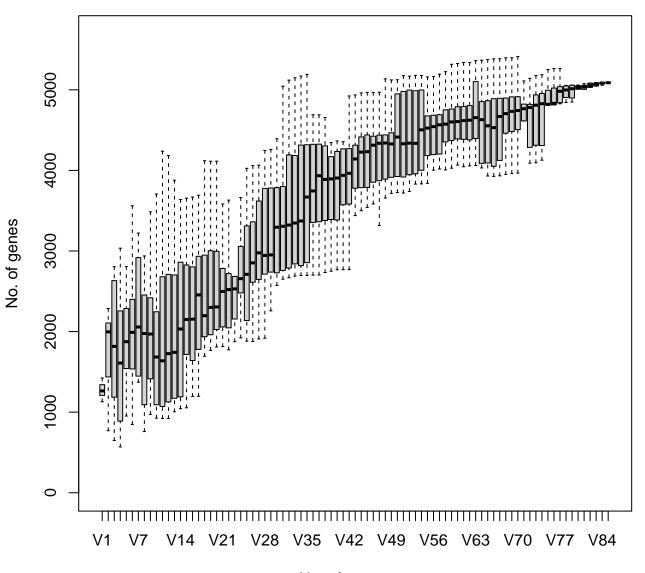
No. of genomes

### No. of genes in the pan-genome



No. of genomes

### Number of unique genes



No. of genomes

# Number of blastp hits with different percentage identity

