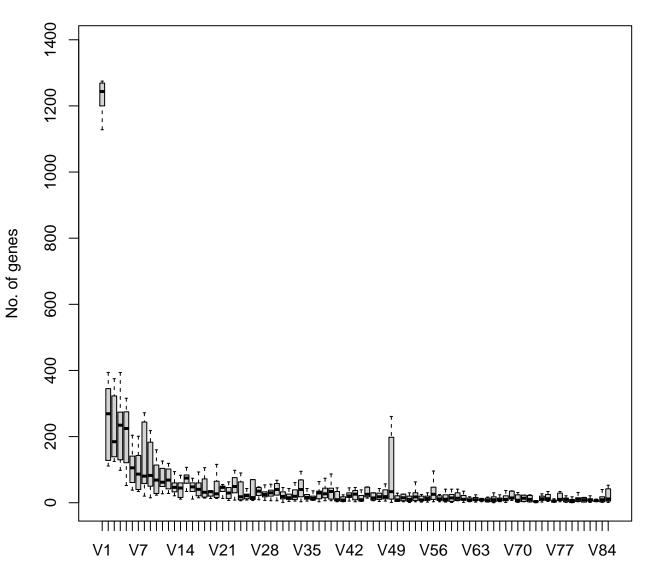
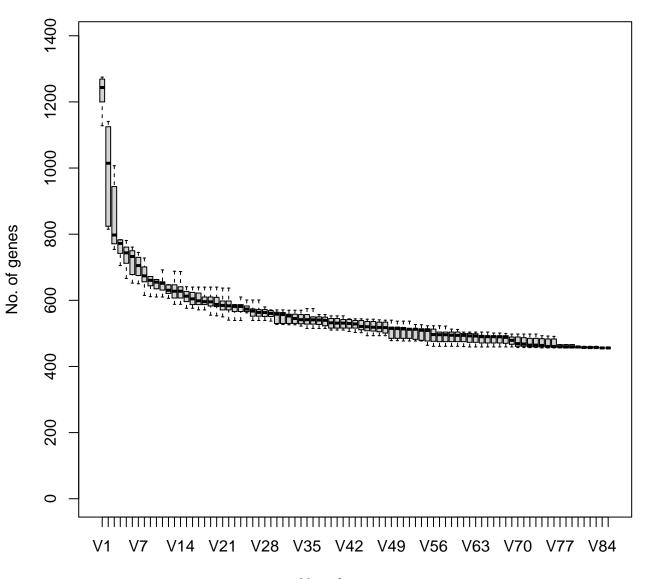
#### Number of new genes



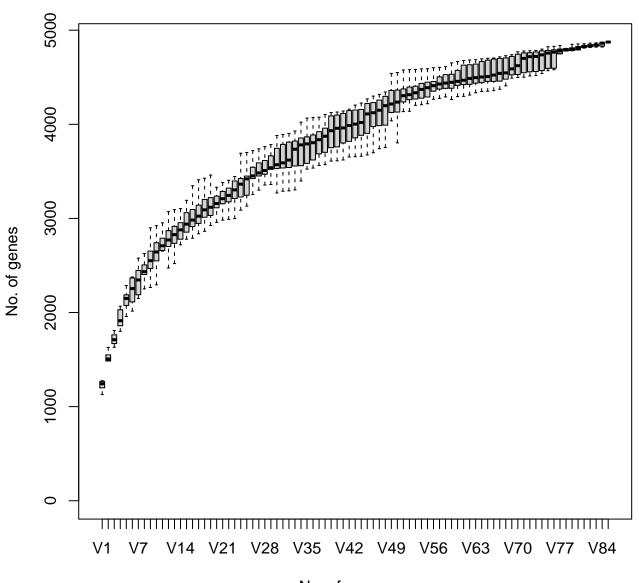
No. of genomes

### Number of conserved genes



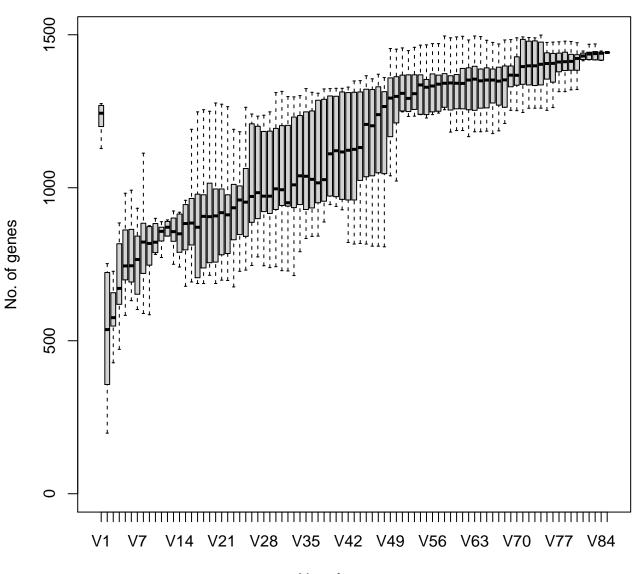
No. of genomes

# 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

