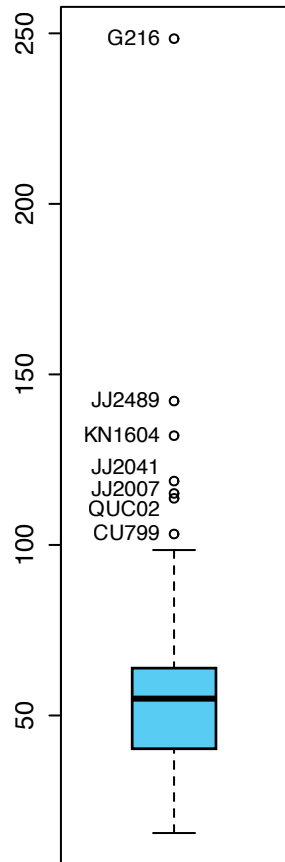
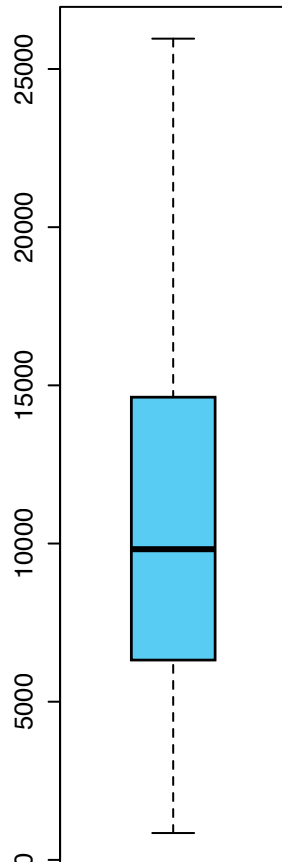
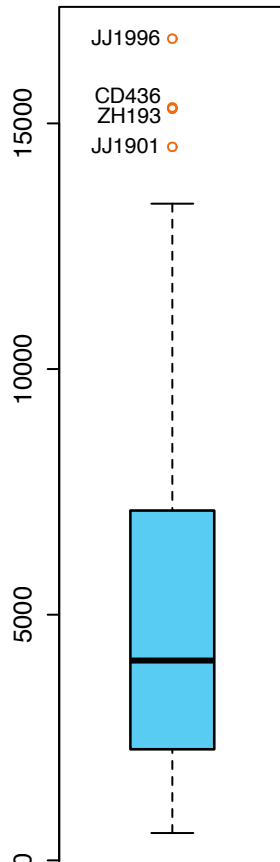
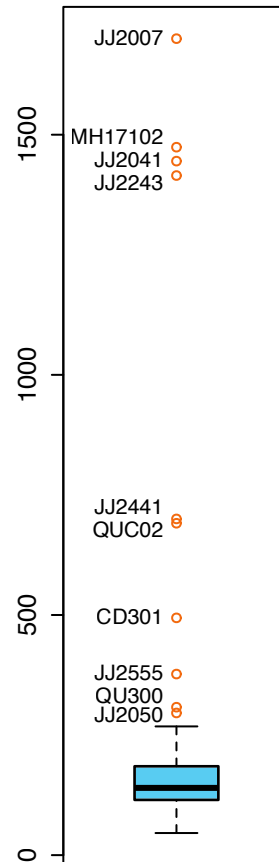


Coverage**No. of unmapped bases****No. of uncalled bases****No. of scaffolds****Genome size**