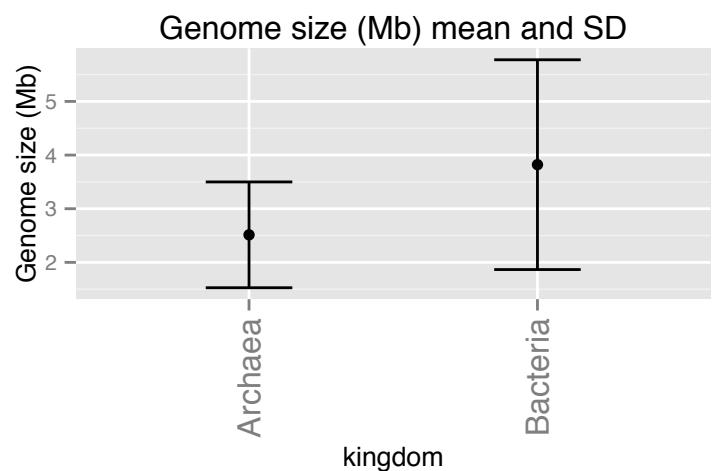
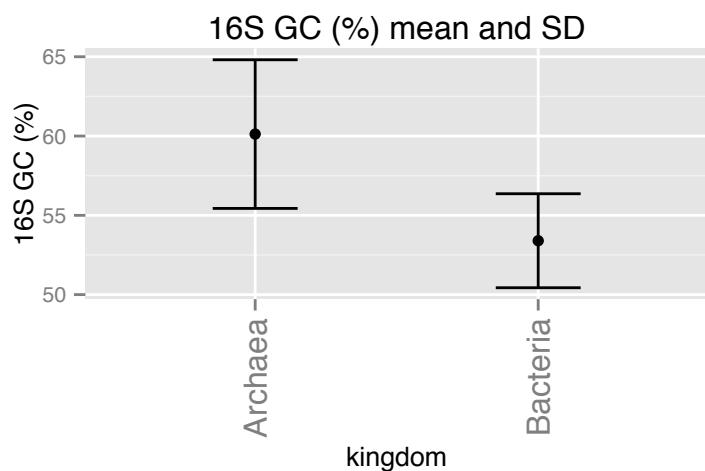
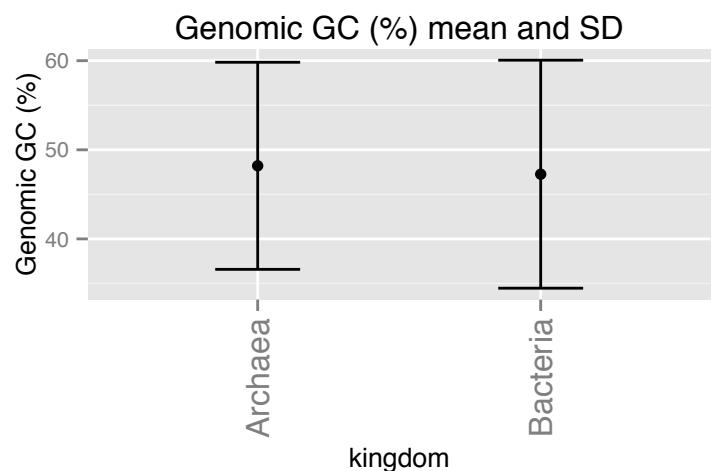
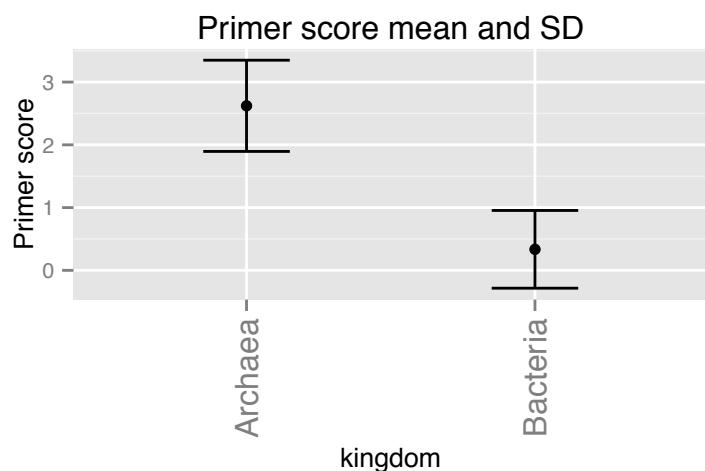
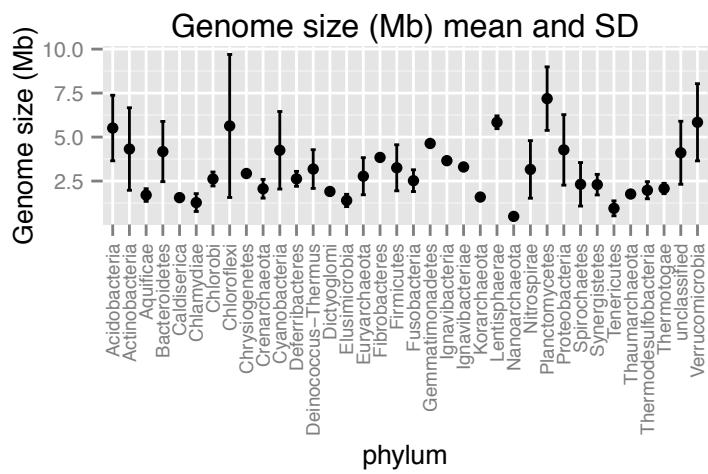
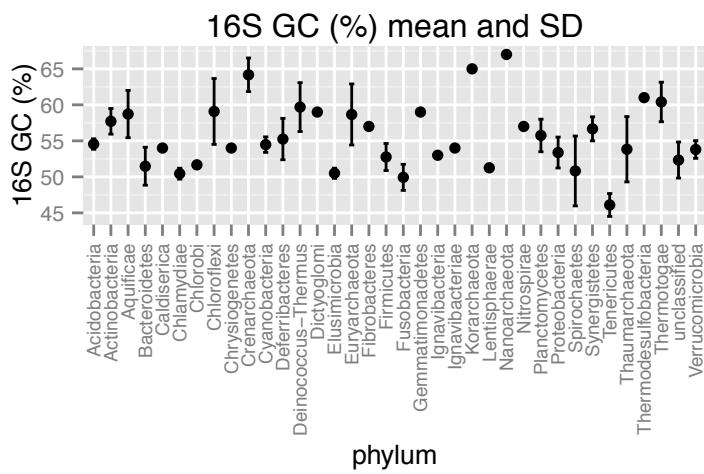
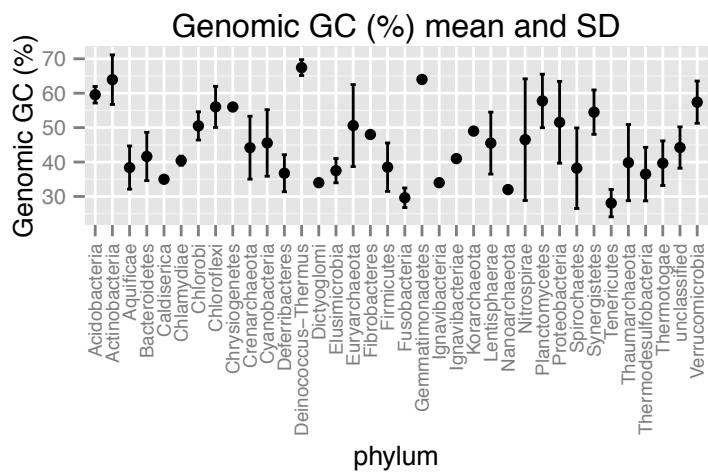
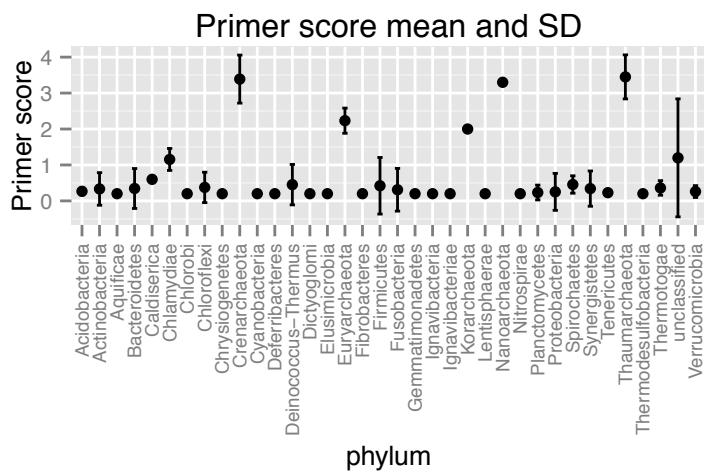


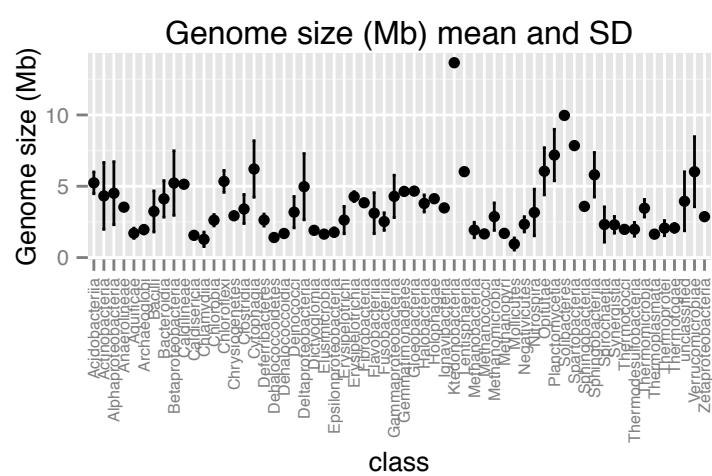
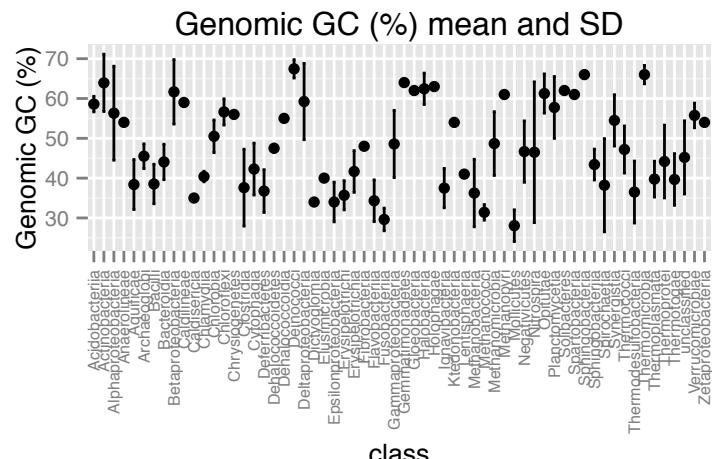
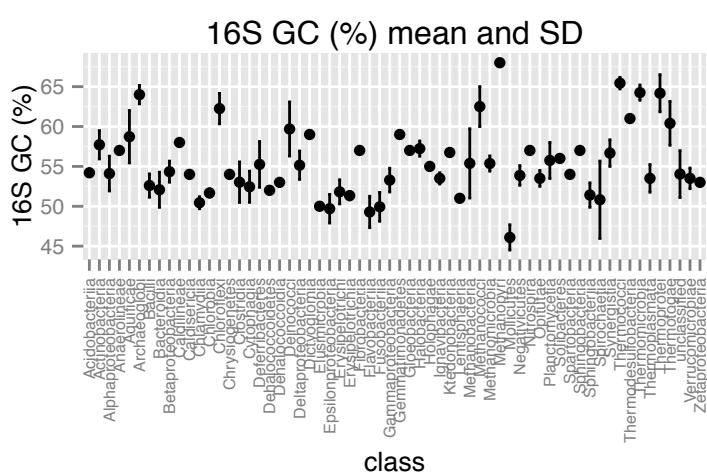
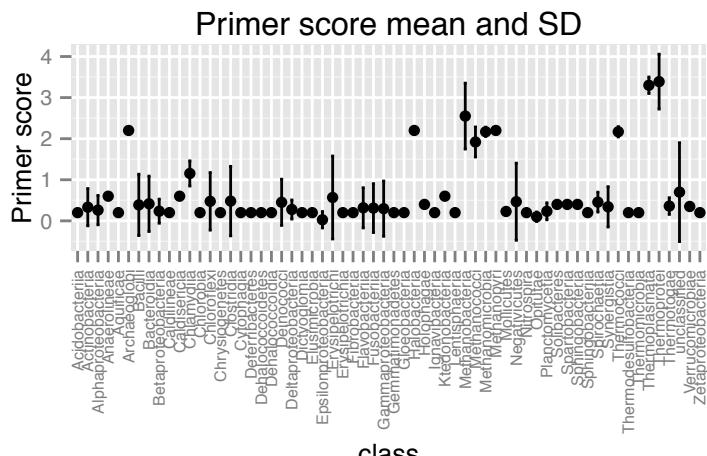
Distributions for kingdom level



Distributions for phylum level

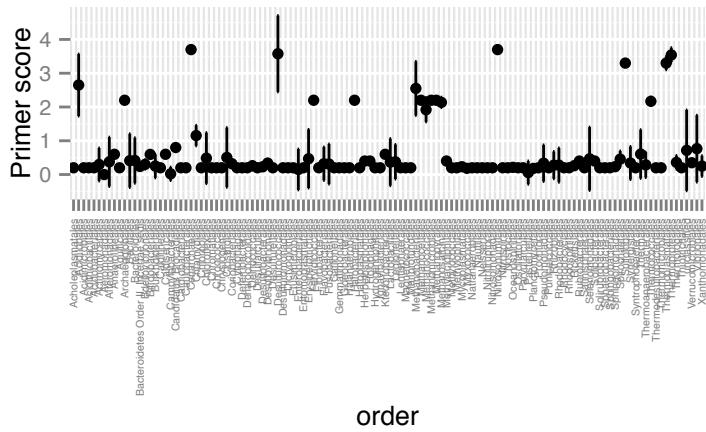


Distributions for class level

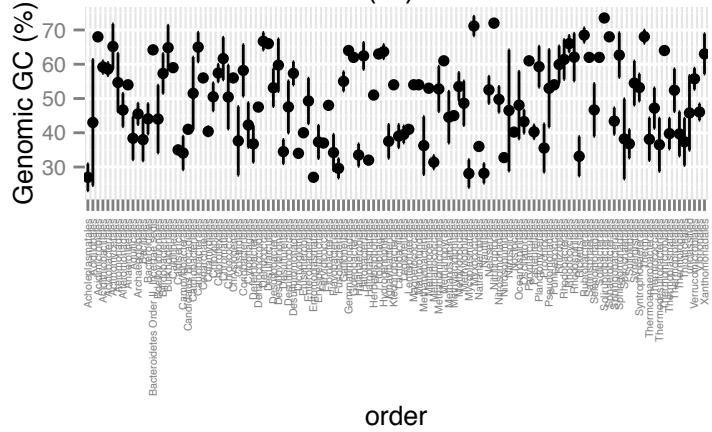


Distributions for order level

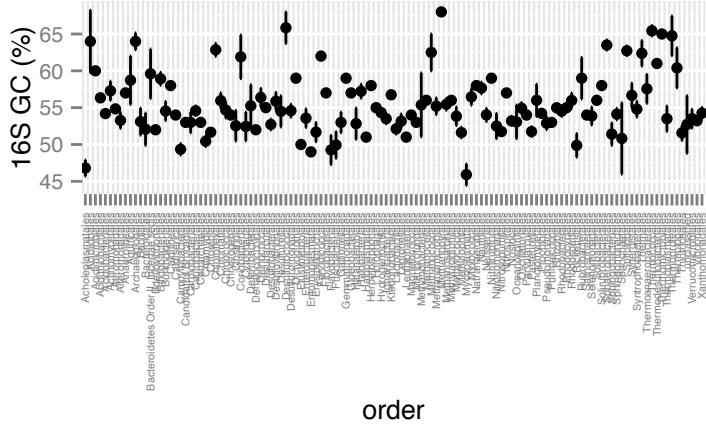
Primer score mean and SD



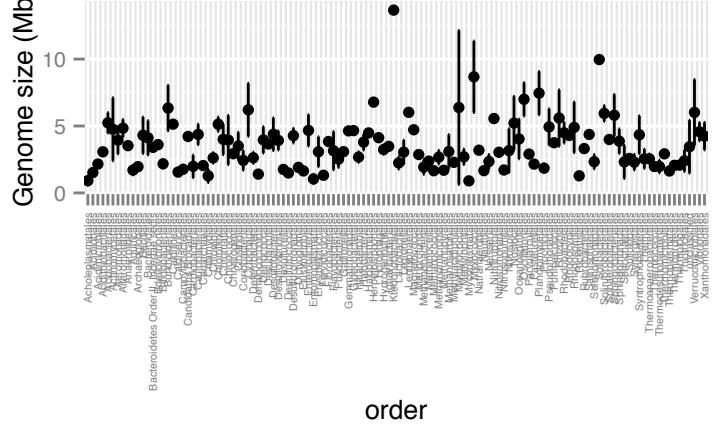
Genomic GC (%) mean and SD



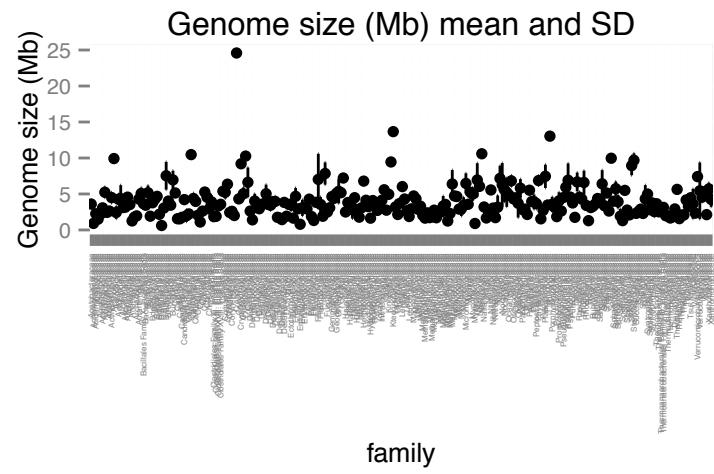
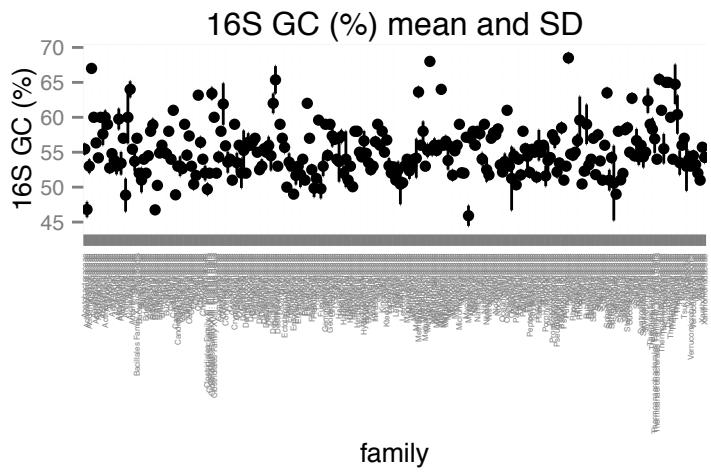
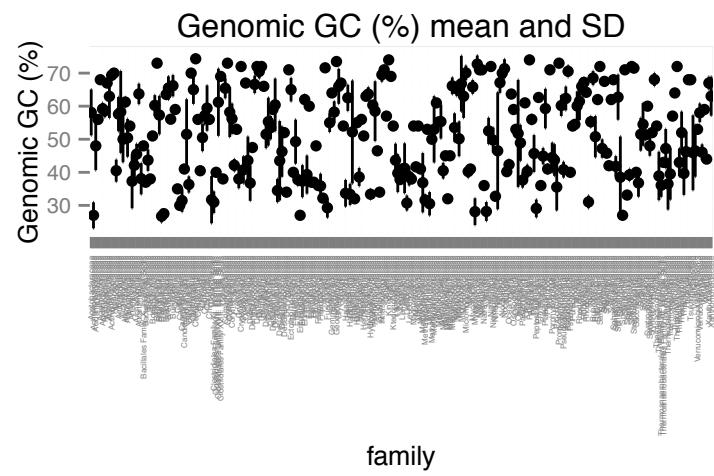
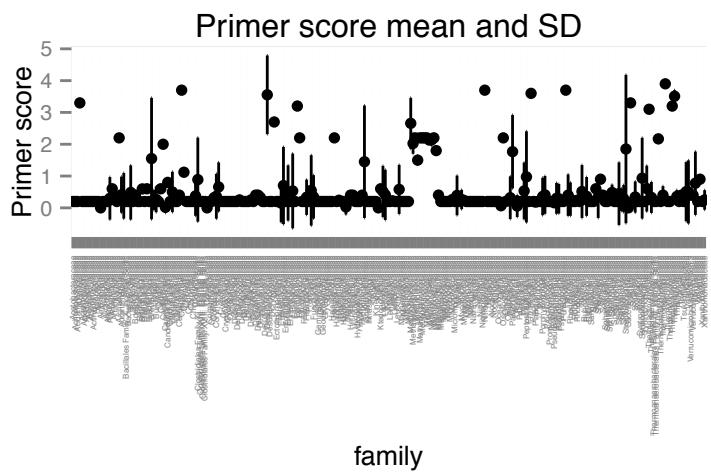
16S GC (%) mean and SD



Genome size (Mb) mean and SD

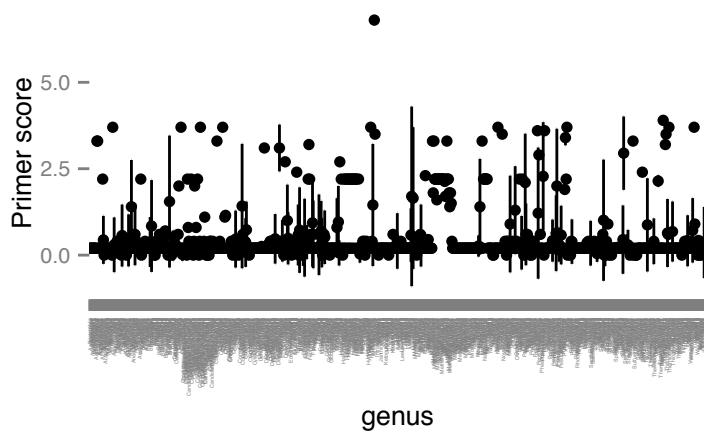


Distributions for family level

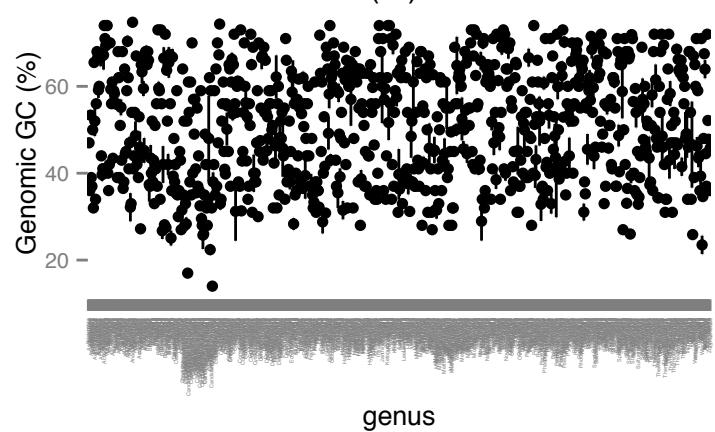


Distributions for genus level

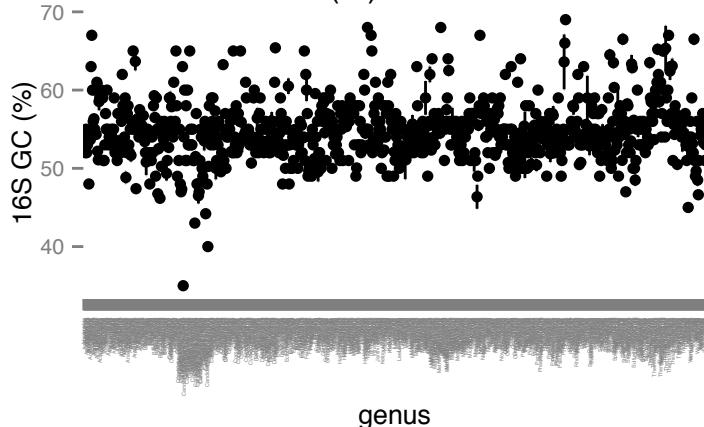
Primer score mean and SD



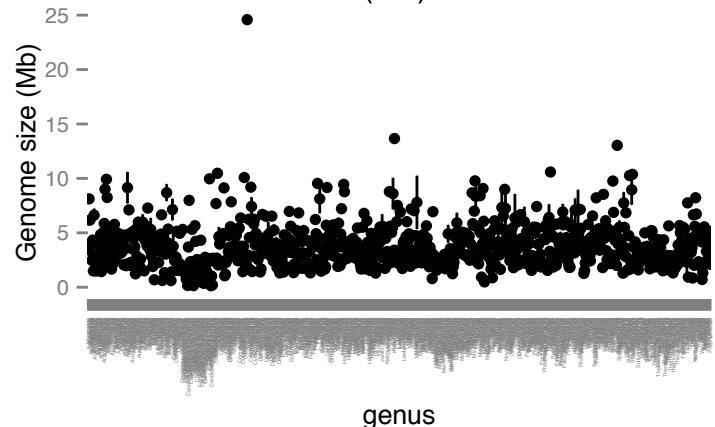
Genomic GC (%) mean and SD



16S GC (%) mean and SD

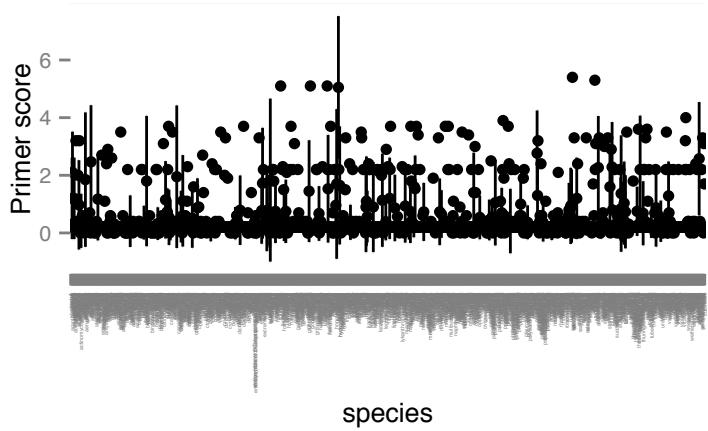


Genome size (Mb) mean and SD

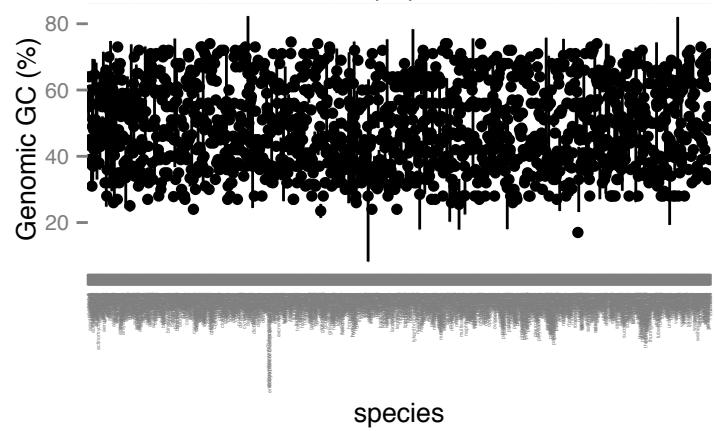


Distributions for species level

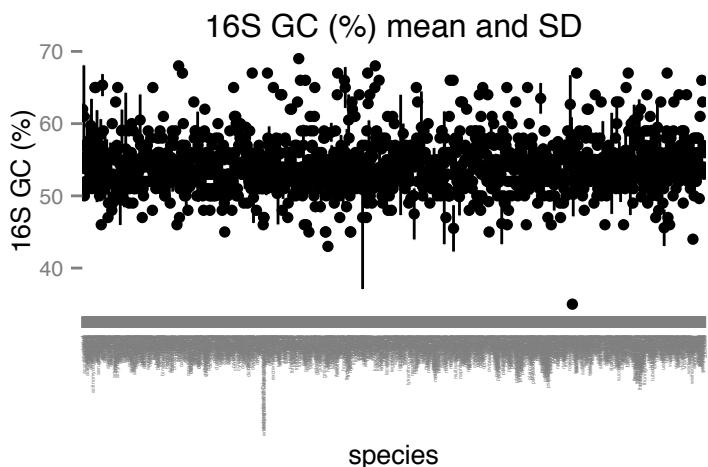
Primer score mean and SD



Genomic GC (%) mean and SD



16S GC (%) mean and SD



Genome size (Mb) mean and SD

