

Illumina GC bias

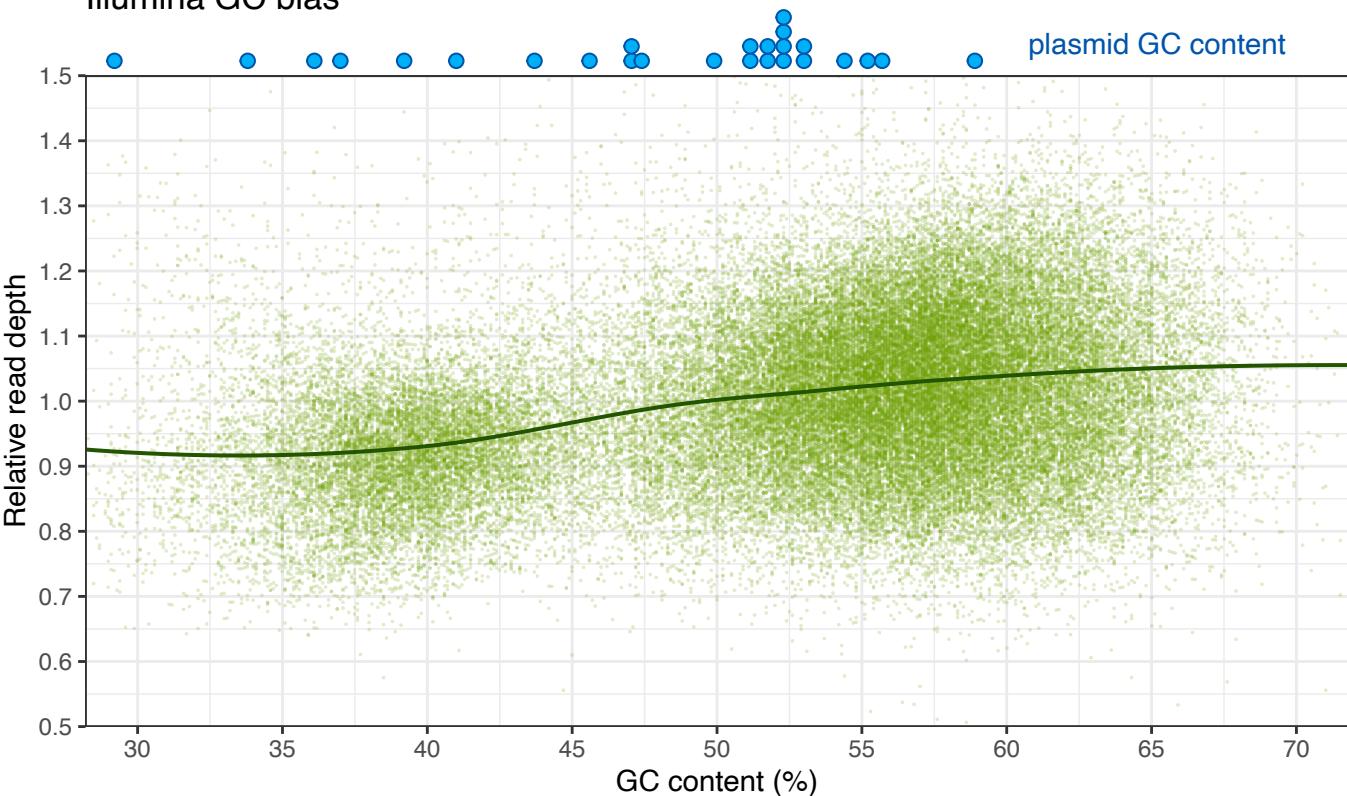


Figure S3: the effect of GC content on Illumina read depth. Each of the green dots represents a 1 kbp sequence window from the chromosomes of the genomes in this study, where read depth was normalised using the average depth for 50% GC windows. The dark green line is a LOESS regression curve fitted to the points. The blue dots on the top of the plot show the average GC content of each plasmid in the study's genomes.