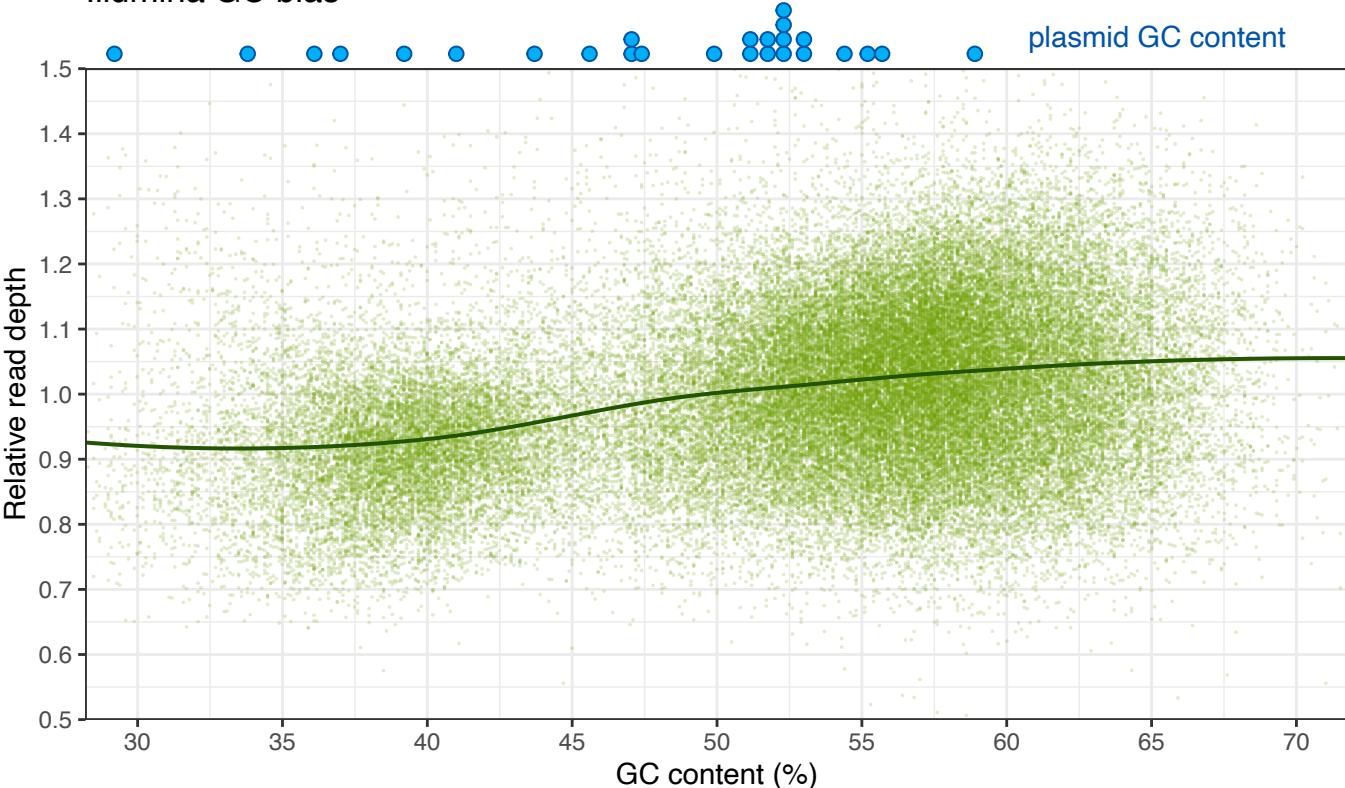


### Illumina GC bias



**Figure S3:** the association between GC content on Illumina read depth. Each green point represents a 1 kbp sequence window from the chromosome of a genome in this study, where read depth was normalised using the mean depth for 49–51% GC windows. The dark green line is a LOESS regression curve fitted to the points. The blue points on the top of the plot show the mean GC content of each plasmid in the study genomes.