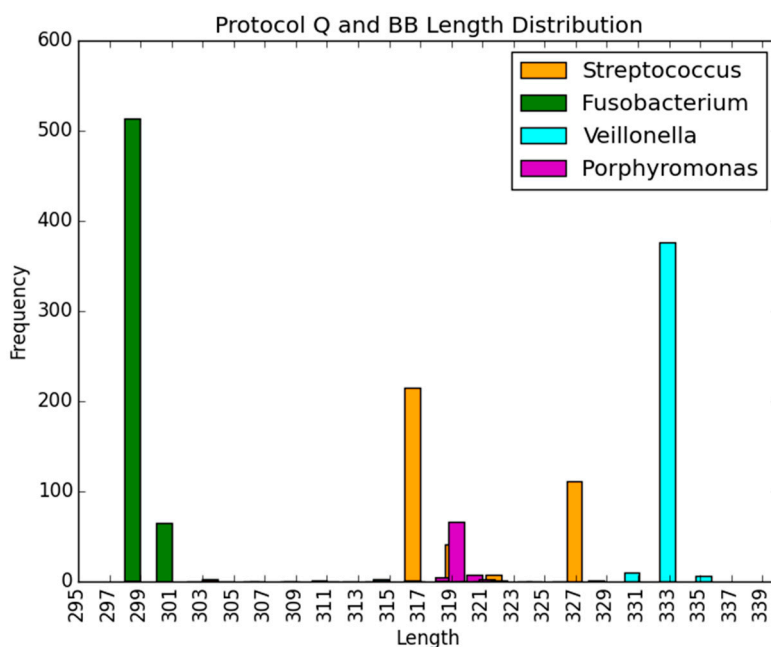


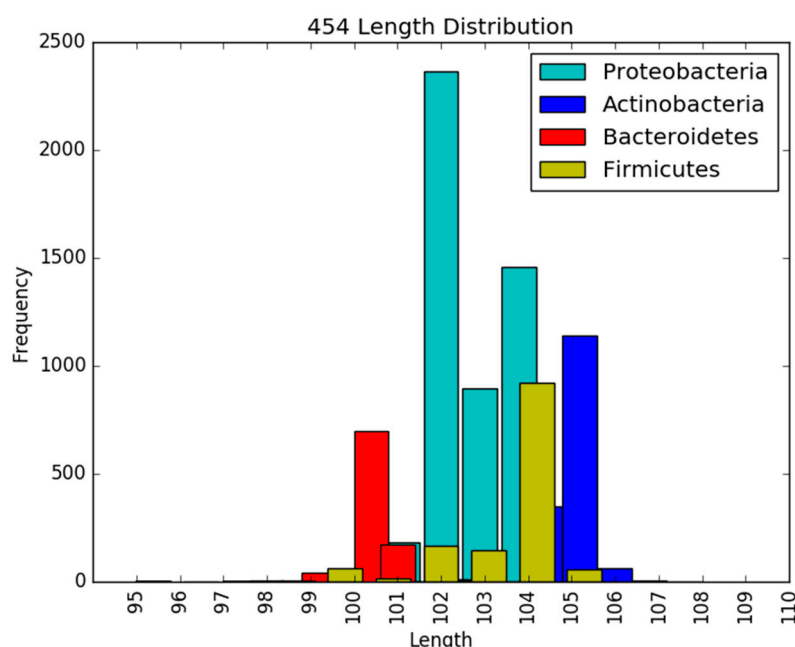


# Supplementary Materials: Phylogenetic Heatmaps Highlight Composition Biases in Sequenced Reads

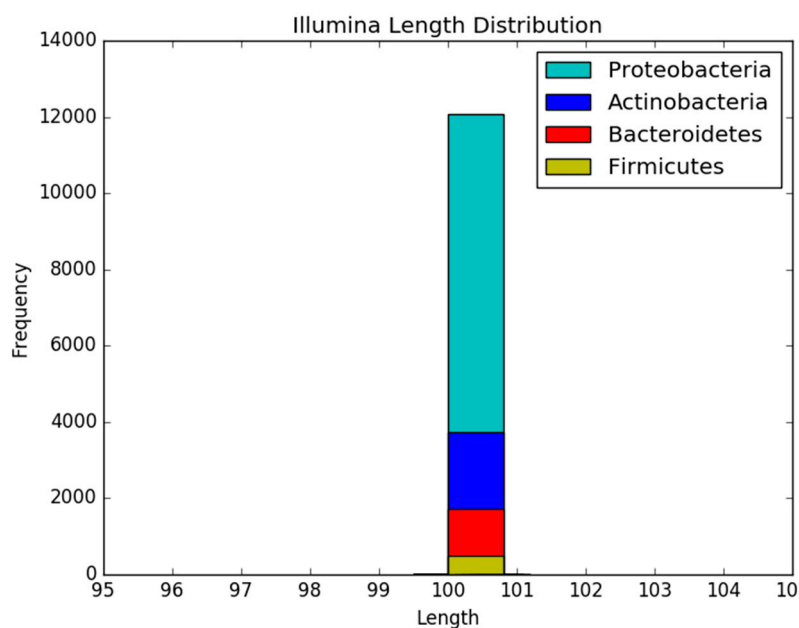
Sulbha Choudhari and Andrey Grigoriev



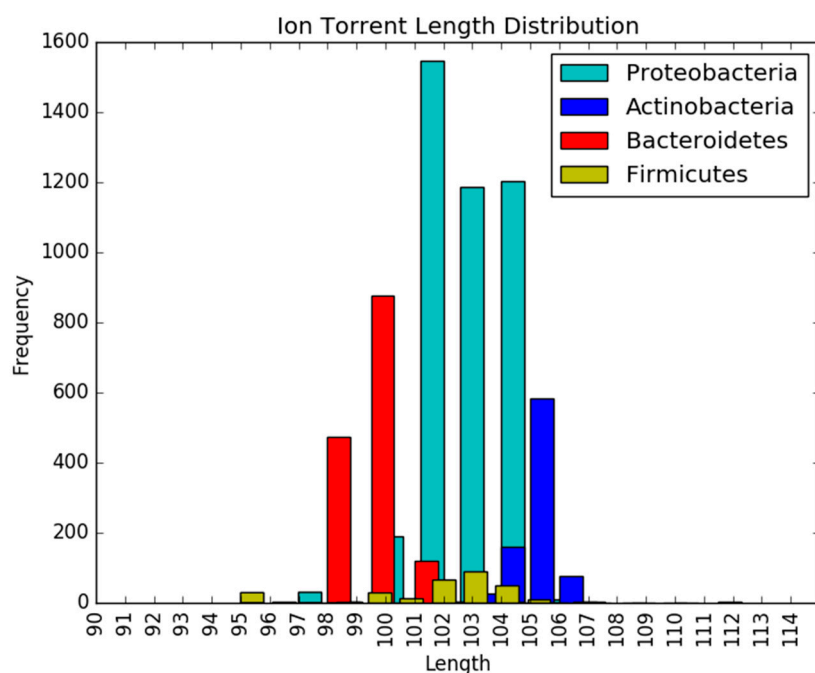
**Figure S1.** Length distribution for the samples in Figure 1 (colors as in Figure 1). Two peaks for *Streptococcus* correspond to the two species.



**Figure S2.** Length distribution for the 454 sample in Figure 6 (colors as in Figure 6B).



**Figure S3.** Length distribution for the Illumina sample in Figure 5 (colors as in Figure 5B).



**Figure S4.** Length distribution for the Ion Torrent sample in Figures 5 and 6 (colors as in Figures 5B and 6B).



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