



Corrigendum: Genomic GC-Content Affects the Accuracy of 16S rRNA Gene Sequencing Based Microbial Profiling due to PCR Bias

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In the original article, there was a mistake in the legend and image for **Figure 2** as published. The legend incorrectly states Log2 ratio and the x-axis label in the image incorrectly states "Log2(Measured/expected relative abundance)". The corrected legend and image for **Figure 2** appears below.

The authors apologize for this error and state that this does not change the scientific conclusions of the article in any way. The original article has been updated.

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FIGURE 2 | Accuracy of the abundance estimates across all five sequencing runs for each bacterial species, expressed as the Log₁₀ ratio of measured relative abundance to the expected relative abundance. Boxplot show the median with 25 and 75 percentiles within the box and whiskers show the range. Dashed line indicates the expected relative abundance of 5%.