Corrigendum: A Metataxonomic Tool to Investigate the Diversity of Treponema

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In the original article, there was a mistake in Figure 1 as published. The figure contains an error in the number of cycles for the “add indices and adapters” step. The step should read 8 cycles instead of 20 cycles. The corrected Figure 1 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 1 | Study design of the metataxonomic assay targeting the V4-region of the 16S rRNA gene. The gray boxes show the tested cycle (c) conditions for each step. The blue shading indicates the modular two-step library preparation.