



## CORRECTION OPEN



# Correction to: Global distribution of a chlorophyll *f* cyanobacterial marker

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Following the publication of this article, the authors noted errors regarding the sequences and lengths of the primers *f\_apcE2t\** and *f\_apcE2M\** listed in Table 1 and Table S2. In addition, the colors of the highlighted bases were missing in Table 1.

The original article and the Supplementary Information have been corrected.

**Table 1.** Main sets of primers used in this study.

<i>f_apcE2M*</i>	CAGAGGATTGCGCGGTGT <span style="background-color: #e0f0ff;">H</span> AT <span style="background-color: #e0ffe0;">Y</span> CC <span style="background-color: #e0f0ff;">H</span> GAAGAT <span style="background-color: #ffe0e0;">R</span> T <span style="background-color: #ffe0e0;">N</span> AC	144	36	<i>apcE2</i>	This study
<i>f_apcE2t*</i>	CAGAGGATTGCGCGGTG	0	17	<i>apcE2</i>	This study
<i>r_apcE2M</i>	GGAGCTCTGCAGATATCGCCCGATG <span style="background-color: #e0ffe0;">R</span> AT <span style="background-color: #e0f0ff;">R</span> TATTC <span style="background-color: #e0ffe0;">D</span> RY	48	37	<i>apcE2</i>	This study
<i>r_apcE2t</i>	GGAGCTCTGCAGATATCGCC	0	20	<i>apcE2</i>	This study
<i>CYA359F</i>	GGGGAAT <span style="background-color: #e0ffe0;">Y</span> TTCCGCAATGGG	2	20	16S <i>rRNA</i>	[80]
<i>C_uni_16S</i>	ACGGGCGGTGTGTAC	0	16	16S <i>rRNA</i>	[81]
<p><sup>a</sup><i>f</i> forward, <i>r</i> reverse. <i>M</i> refers to medium degeneracy in comparison with other primers tested in this study (Table S2). <i>t</i> represents tags. Asterisks mark versions of the same primer with an alternate tag.  <sup>b</sup>All degenerate primers had 5' tags. Red boxes mark any base (<i>N</i>); blue boxes mark bases with a degeneracy of 3 (H = not G; D = not C; V = not T); green boxes mark bases with a low degeneracy (Y = C/T; R = A/G).</p>					



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