Erratum to: Distribution and phylogenetic diversity of *cbbM* genes encoding RubisCO form II in a deep-sea hydrothermal field revealed by newly designed PCR primers

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In the original publication of this paper, in the sections “Primer design for *cbbM* genes” and “PCR clone library analysis”, the reverse primer name is incorrectly given as cbbM1226R which should be cbbM1126R.

Also in the section “Primer design for *cbbM* genes”, the expected amplicon size is incorrectly given as 800–900 bp which should be 700–800 bp.

The online version of the original article can be found under
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