

CORRECTION

Correction: Application of Stochastic Labeling with Random-Sequence Barcodes for Simultaneous Quantification and Sequencing of Environmental 16S rRNA Genes

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There are errors in the fourth and fifth sentences of the Quantification by qSeq subsection of the Materials and Methods section. The correct sentences are: The SPE conditions consisted of initial denaturation at 98°C for 2 min, cooling to 55°C at 0.3°C/s, and elongation at 68°C for 10 min. Subsequently, excess primers were digested by addition of 4 µl of Exonuclease I (5 U/µl, TaKaRa Bio) and incubation at 37°C for 120 min, followed by inactivation of the enzyme by incubation at 80°C for 30 min.

Reference

1. Hoshino T, Inagaki F (2017) Application of Stochastic Labeling with Random-Sequence Barcodes for Simultaneous Quantification and Sequencing of Environmental 16S rRNA Genes. PLoS ONE 12(1): e0169431. doi:[10.1371/journal.pone.0169431](https://doi.org/10.1371/journal.pone.0169431) PMID: [28052139](https://pubmed.ncbi.nlm.nih.gov/28052139/)



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