

CORRECTION

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Correction to: More than 2500 years of oil exposure shape sediment microbiomes with the potential for syntrophic degradation of hydrocarbons linked to methanogenesis

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Correction

Following publication of the original article [1], the authors requested two minor changes. The first is to the name of one of the genes of the TCA cycle (*fumABC*) in Fig. 3. The correct figure with the name of the gene added is included below.

The second is to a sentence in the Authors' Contribution section which should be changed from:

KT, RR, DGH, and MS designed the study. AM, KT, PA, DGH, and CEV collected the samples from Keri Lake. HW analyzed the hydrocarbon content and composition. AM, GV, and AS generated and analyzed the sequence data. AM, RR, MS, and AS conceptualized and wrote the manuscript. All authors contributed to revisions and approved the final manuscript.

To:

KT, RR, DGH, and MS designed the study. PA designed the geological study. AM, KT, PA, DGH, and CEV collected the samples from Keri Lake. HW analyzed the hydrocarbon content and composition. AM, GV, and AS generated and analyzed the sequence data. AM, RR, MS, and AS conceptualized and wrote the manuscript. All authors contributed to revisions and approved the final manuscript.

The original article has been corrected.

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