

Correction

Correction: Molecular Phylogenetics and Systematics of the Bivalve Family Ostreidae Based on rRNA Sequence-Structure Models and Multilocus Species Tree

The *PLOS ONE* Staff

There is an error in the seventh sentence of the Abstract. The subfamily name is misspelled. The correct sentence is: Therefore, the subfamily *Crassostreinae* (including *Crassostrea*), *Saccostreinae* **subfam. nov.** (including *Saccostrea* and tentatively *Striostrea*) and *Ostreinae* (including *Ostreinae* and *Lophinae* taxa) are recognized.

There is a typographical error in the second sentence of the second paragraph in the “Dataset, DNA extraction, amplification and sequencing” section of the Methods. The correct sentence is: It is worth mentioning that, despite some oysters show a split of the 16S rRNA gene into two segments, e.g. in the Asian *Crassostrea*, *C. virginica*, *Saccostrea mordax*, and *Ostrea edulis* [26], [36], our target 16S fragment (3' half portion of the 16S rRNA gene corresponding to domains IV and V) is entirely located in the second segment.

Reference

1. Salvi D, Macali A, Mariottini P (2014) Molecular Phylogenetics and Systematics of the Bivalve Family Ostreidae Based on rRNA Sequence-Structure Models and Multilocus Species Tree. *PLoS ONE* 9(9): e108696. doi:10.1371/journal.pone.0108696

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