CORRECTION

Correction: Integrated Taxonomy Reveals Hidden Diversity in Northern Australian Fishes: A New Species of Seamothe (Genus *Pegaseus*)

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In [Fig 10](#), the COI GenBank Accession Numbers for the *Pegaseus* species individuals in Plate B are incorrect. Please see the correct [Fig 10](#) here.

OPEN ACCESS

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Fig 10. Molecular species identification of *Pegasus* species using Genetic treeML trees. (A) sequences from the 16S gene; (B) sequences from the COI gene. Trees are based on the K2 evolutionary distance model and are shown here with mined *Pegasus* and *Eurypegusus* sequences from GenBank. The trees are shown here with an *E. draconis* outgroup. Bootstrap support values (following 1000 replicates) are shown above the nodes.

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The COI GenBank Accession Numbers for the *Pegasus* species individuals in [S2 Table](#) are incorrect. Please view the correct [S2 Table](#) below.

Supporting information

S2 Table. Genetic sample information.
(XLSX)

Reference

1. Osterhage D, Pogonoski JJ, Appleyard SA, White WT (2016) Integrated Taxonomy Reveals Hidden Diversity in Northern Australian Fishes: A New Species of Seamothe (Genus *Pegasus*). PLoS ONE 11 (3): e0149415. <https://doi.org/10.1371/journal.pone.0149415> PMID: 26934529