



Whole Mitochondrial Genomes Applied to Phylogenetics of Aquatic Animals

Guest Editor:

Prof. Dr. Joseph Quattro

Department of Biological
Sciences, University of South
Carolina, Columbia, South
Carolina 29045, USA

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Message from the Guest Editor

Dear Colleagues,

Recent technological advances have facilitated the acquisition of mitochondrial genomes and their use to answer questions regarding the phylogenetic relationships of aquatic organisms. We seek manuscripts that utilize mitochondrial genomes to help support or refute phylogenetic hypotheses that require wider coverage than a handful of targeted loci might provide. The focus of these manuscripts can be methodological (e.g., data acquisition and phylogenetic analyses) or, more narrowly, applications of these sorts of data to interesting questions in aquatic biology. We ask that focused manuscripts outline a clear hypothesis supporting the rationale for the study, rather than the simple accumulation of mitochondrial genomes without adequate justification.

Prof. Dr. Joseph Quattro

Guest Editor





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Editor-in-Chief

Prof. Dr. Maria Angeles Esteban

Department of Cell Biology and Histology, Faculty of Biology, University of Murcia, 30100 Murcia, Spain

Message from the Editor-in-Chief

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Contact Us

Fishes Editorial Office
MDPI, Grosspeteranlage 5
4052 Basel, Switzerland

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