

Accepted Manuscript

Phylogenetic analysis shows the general diversification pattern of deep-sea notacanthiforms (Teleostei: Elopomorpha)

David Barros-García, Elsa Froufe, Rafael Bañón, Juan Carlos Arronte, Alejandro De Carlos

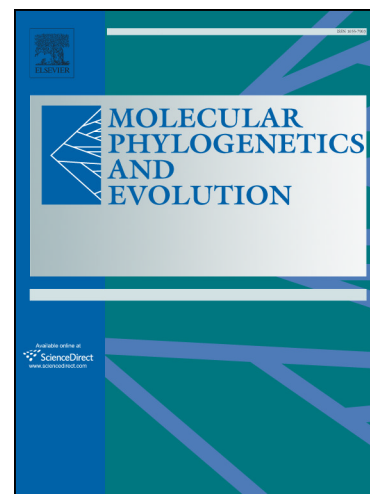
PII: S1055-7903(17)30850-3
DOI: <https://doi.org/10.1016/j.ympev.2018.03.007>
Reference: YMPEV 6079

To appear in: *Molecular Phylogenetics and Evolution*

Received Date: 22 November 2017
Revised Date: 2 February 2018
Accepted Date: 7 March 2018

Please cite this article as: Barros-García, D., Froufe, E., Bañón, R., Carlos Arronte, J., De Carlos, A., Phylogenetic analysis shows the general diversification pattern of deep-sea notacanthiforms (Teleostei: Elopomorpha), *Molecular Phylogenetics and Evolution* (2018), doi: <https://doi.org/10.1016/j.ympev.2018.03.007>

This is a PDF file of an unedited manuscript that has been accepted for publication. As a service to our customers we are providing this early version of the manuscript. The manuscript will undergo copyediting, typesetting, and review of the resulting proof before it is published in its final form. Please note that during the production process errors may be discovered which could affect the content, and all legal disclaimers that apply to the journal pertain.



Phylogenetic analysis shows the general diversification pattern of deep-sea notacanthiforms (Teleostei: Elopomorpha)

DAVID BARROS-GARCÍA^{1,2} ELSA FROUFE³ RAFAEL BAÑÓN^{4,5}, JUAN CARLOS ARRONTE⁶, ALEJANDRO DE CARLOS¹

¹Department of Biochemistry, Genetics and Immunology, University of Vigo, C/ Fonte das Abelleiras s/n, 36310 Vigo, Spain. E-mails: davbarros@uvigo.es; adcarlos@uvigo.es

²Programa de Doctorado en Metodología y Aplicaciones en Ciencias de la Vida, Facultad de Biología. Universidad de Vigo, C/Fonte das Abelleiras s/n, 36310 Vigo, Spain.

³Centro Interdisciplinar de Investigação Marinha e Ambiental (CIIMAR/CIMAR). Terminal de Cruzeiros do Porto de Leixões, Avenida General Norton de Matos, S/N 4450-208 Matosinhos, Portugal. E-mail: elsafroufe@gmail.com

⁴Instituto de Investigaciones Marinas, Consejo Superior de Investigaciones Científicas (IIM-CSIC), C/ Eduardo Cabello, 6, 36208 Vigo, Spain. E-mail: anoplogaster@yahoo.es

⁵Grupo de Estudos do Medio Mariño (GEMM), Puerto Deportivo s/n, 15960 Ribeira, A Coruña, Spain

⁶Instituto de Hidráulica Ambiental (IH Cantabria). Universidad de Cantabria, PCTCAN, C/Isabel Torres nº 15, 390011, Santander, Spain. E-mail: j_aronnte@hotmail.com

Corresponding author: davbarros@uvigo.es

Abstract

The Notacanthiformes is an ancient group of deep-sea ray-finned fishes comprising 27 species in two families; Halosauridae and Notacanthidae. Although many studies have tried to reconstruct the phylogenetic relationships among the major clades of Elopomorpha, little is known about the evolutionary history of notacanthiforms. Molecular and morphological data were used to test previous hypotheses regarding the phylogenetic relationships among notacanthiform taxa, and to unravel the origin and evolution of this group. The molecular analyses of notacanthids showed similar results to those previously obtained employing osteological data, which proposed the existence of the Lipogenyinae (*Lipogenys*) and Notacanthinae (*Notacanthus* + *Polyacanthonotus*) subfamilies. Nevertheless, when the external morphology data is considered *Lipogenys* is more related to *Notacanthus* than *Polyacanthonotus*. The analyses could not fully resolve the inner relationships of the

Download English Version:

<https://daneshyari.com/en/article/8648898>

Download Persian Version:

<https://daneshyari.com/article/8648898>

[Daneshyari.com](https://daneshyari.com)