Accepted Manuscript

Title: A Deep Hidden Diversity of

Dictyostelia<!--<RunningTitle>Deep Hidden Diversity of

Dictyostelia</RunningTitle>->

Authors: Sandra L. Baldauf, Maria Romeralo, Omar

Fiz-Palacios, Nahid Heidari

PII: S1434-4610(18)30001-4

DOI: https://doi.org/10.1016/j.protis.2017.12.005

Reference: PROTIS 25605

To appear in:

Received date: 4-9-2017 Revised date: 19-12-2017 Accepted date: 30-12-2017

Please cite this article as: Baldauf, Sandra L., Romeralo, Maria, Fiz-Palacios, Omar, Heidari, Nahid, A Deep Hidden Diversity of Dictyostelia.Protist https://doi.org/10.1016/j.protis.2017.12.005

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.



ACCEPTED MANUSCRIPT

ORIGINAL PAPER

A Deep Hidden Diversity of Dictyostelia

Running title: Deep Hidden Diversity of Dictyostelia

Sandra L. Baldauf¹, Maria Romeralo, Omar Fiz-Palacios, and Nahid Heidari

Department of Systematic Biology, Evolutionary Biology Centre, Norbyvägen 18D, University of Uppsala, SE-75236 Uppsala, Sweden

¹Corresponding author; fax +46 184 71 64 57 e-mail sandra.baldauf@ebc.uu.se (S. L. Baldauf).

Submitted September 4, 2017; Accepted December 30, 2017

Monitoring Editor: C. Graham Clark

Dictyostelia is a monophyletic group of transiently multicellular (sorocarpic) amoebae, whose study is currently limited to laboratory culture. This tends to favour faster growing species with robust sorocarps, while species with smaller more delicate sorocarps constitute most of the group's taxonomic breadth. The number of known species is also small (~150) given Dictyostelia's molecular depth and apparent antiquity (>600 myr). Nonetheless, dictyostelid sequences are rarely recovered in culture independent sampling (ciPCR) surveys. We developed ciPCR primers to specifically target dictyostelid small subunit (SSU or 18S) rDNA and tested them on total DNAs extracted from a wide range of soils from five continents. The resulting clone libraries show mostly dictyostelid sequences (~90%), and phylogenetic analyses of these sequences indicate novel lineages in all four dictyostelid families and most genera. This is especially true for the species-rich Heterostelium and Dictyosteliaceae and but also the less species-rich Raperosteliaceae. However, the most novel deep branches are found in two very species-poor taxa, including the deepest branch yet seen in the highly divergent Cavenderiaceae. These results confirm a deep hidden diversity of Dictyostelia, potentially including novel morphologies and developmental

Download English Version:

https://daneshyari.com/en/article/8392888

Download Persian Version:

https://daneshyari.com/article/8392888

Daneshyari.com