New insights on the phylogenetic relationships of the Proseriata (Platyhelminthes), with proposal of a new genus of the family Coelogynoporidae

MARCO CURINI-GALLETTI¹, BONNIE L. WEBSTER², TINE HUYSE²,³, MARCO CASU¹, ERNEST R. SCHOCKAERT⁴, TOM J. ARTOIS⁴ & D. TIMOTHY J. LITTLEWOOD²

¹Dipartimento di Zoologia e Genetica Evoluzionistica, University of Sassari, via Maroni 25, I 07100 Sassari, Italy
²Parasitic Worms Group, Department of Zoology, The Natural History Museum, Cromwell Road, London SW7 5BD, UK
³Katholieke Universiteit Leuven, Laboratory of Animal Diversity and Systematics, Ch. Deberiotstraat 32, B-3000 Leuven, Belgium
⁴Research group Biodiversity, Phylogeny and Population Studies, Centre for Environmental Sciences, Hasselt University, Campus Diepenbeek, Agoralaan Building D, B-3590 Diepenbeek, Belgium

Abstract

Proseriata is a diverse and species-rich taxon of Neoophora (Platyhelminthes). The group is divided into two morphologically well-characterised taxa: Lithophora and Unguiphora. Previous molecular analyses, aimed at ascertaining in-group relationships of the Proseriata, were equivocal in supporting the monophyly of the Lithophora, and of one of its families, the Coelogynoporidae. Here we present a new phylogeny of the Proseriata, based on complete ssrDNA and partial (D1-D6) lsrDNA. The analysis includes 36 proseriate species. 13 new sequences from seven species are used, four of which from species of the Coelogynoporidae and one from a new species of the recently established family Calviriidae. Phylogenetic analysis was performed using Bayesian Inference (BI), maximum likelihood (ML) and maximum parsimony (MP). Clades were considered to have high nodal support if BI posterior probability and ML and MP bootstrap percentages were \( > 90 \% \). The resulting cladogram strongly supports the monophyly of the Lithophora. In addition, the monophyly of the Coelogynoporidae and their inclusion in the Lithophora are corroborated. Within the Lithophora, two sister clades are supported, including Coelogynoporidae+Calviriidae, and Otoplanidae+Archimonocelididae+Monocelididae respectively. In the Coelogynoporidae, there was a poor correspondence between estimates of phylogeny and the monophyly of genera, suggesting the family is in need of systematic revision. Current morphology-based systematic arrangements of Otoplanidae and Monocelididae are not supported by molecular results. The enigmatic taxon Ciliopharyngiella (formerly included in the Rhabdocoela) clusters with the Unguiphora, although with low support. One new species of Coelogynoporidae (Parainvenusta englarorum n. gen n. sp.) is described and discussed in the appendix.

Key words: phylogeny; taxonomy; ssrDNA; lsrDNA; Unguiphora; Lithophora; Ciliopharyngiella

Introduction

The first comprehensive molecular study specifically designed to assess both in-group and out-group phylogenetic relationships of the Proseriata, a species-rich and widespread taxon of Neoophoran flatworms, left two major problems unsolved. In fact, the monophyly of the Proseriata themselves and of one of its major subtaxa, the Lithophora were unsupported by the molecular trees obtained (Littlewood et al., 2000). These results were in stark contrast with most of the morphology-based phylogenies of the group, which instead clearly supported their monophyly (Sopott-Ehlers, 1985; Ehlers, 1985; Ehlers & Sopott-Ehlers, 1990). However, molecular data partially agreed with alternative, morphology based reconstructions, which hypothesized paraphyletic Lithophora (Martens & Schockaert, 1988) and Proseriata (Brüggemann, 1986).