



Comparative phylogenetic study of Stichotrichia (Alveolata: Ciliophora: Spirotrichea) based on 18S-rDNA sequences

T.S. Paiva¹, B.N. Borges², M.L. Harada² and I.D. Silva-Neto¹

¹Departamento de Zoologia, Laboratório de Protistologia,
Instituto de Biologia, CCS, Universidade Federal do Rio de Janeiro,
Rio de Janeiro, RJ, Brasil

²Laboratório de Biologia Molecular Francisco Mauro Salzano,
Instituto de Ciências Biológicas, Cidade Universitária Prof. José Silveira Netto,
Universidade Federal do Pará, Belém, PA, Brasil

Corresponding author: T.S. Paiva
E-mail: tpaiva@biologia.ufrj.br

Genet. Mol. Res. 8 (1): 223-246 (2009)
Received December 8, 2008
Accepted January 19, 2009
Published March 3, 2009

ABSTRACT. Since molecular phylogenies of stichotrich ciliates started to be published, some remarkable contradictions to morphology-based classifications have been reported, such as the Convergent Evolution of Urostylids and Uroleptids (CEUU) hypothesis, the *Halteria* paradox, the polyphyly of *Oxytricha* and of Stichotrichia. We hypothesized the internal phylogeny of 18S-rDNA from 53 morphological species of stichotrichs and their relationships with Hypotrichia and Oligotrichia using parsimony and neighbor-joining methods, including new data from *Pseudouroleptus caudatus* and *Strongylidium pseudocrassum*. Competing phylogenetic scenarios were compared using statistical tests, and the results suggest the reconsideration of both CEUU and the position of *Halteria* among flexible-body oxytrichids. The polyphyly of *Oxytricha* was not rejected and the monophyly of Stichotrichia was accepted based on parsimony analysis if *Pseudoamphisiella* is considered an external (discocephalid related) taxon.

Key words: CEUU; *Halteria*; *Oxytricha*; *Strongylidium*; *Pseudouroleptus*