Phylogenetic relationships, based on SSU rDNA sequences, among the didelphic genera of the family Trichodoridae from Portugal

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Summary – A survey of virus vector trichodorid nematodes was carried out in the central and northern regions of Portugal. Morphobiometric identification showed the presence of trichodorid species previously reported from Portugal, except for *Paratrichodorus porosus*, which is reported for the first time in Continental Europe. Small subunit ribosomal DNA (SSU rDNA) sequences of ten different species occurring in Portugal were obtained and a phylogenetic analysis based on their alignment was performed to infer relationships among the different Portuguese trichodorid species and three non-indigenous populations (*Nanidorus minor*, *P. allius* and *P. teres*). The re- sulting phylogenetic tree is in agreement with the currently accepted classification for Trichodoridae, except for *Nanidorus*, which clusters together with *Trichodorus* species, while the genera *Paratrichodorus* and *Trichodorus* appear as two distinct groups. A better understanding of the generic groupings in the family Trichodoridae was found. Based on the new molecular analyses we herein accept *Nanidorus* as a valid genus.

Keywords – 18S rDNA, molecular, *Nanidorus*, *Paratrichodorus*, *Paratrichodorus porosus*, phylogeny, tobacco rattle virus (TRV), *Trichodorus*, *Tylolaimophorus minor*.