

Molecular variability in *Brycon cf. pesu*Müller and Troschel, 1845 (Characiformes: Characidae) from the Araguaia-Tocantins Basin

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ABSTRACT. Brycon pesu is a small-sized fish distributed throughout the Amazon and Orinoco Basins and other coastal basins of northeastern South America. Brycon cf. pesu specimens from the Araguaia-Tocantins Basin are currently separated into two morphotypes, Brycon sp1 and Brycon sp2, owing to different coloration of their anal fin. Brycon sp2 has a reddish margin stripe on the anal fin which morphologically distinguishes it from Brycon sp1. In the present research, nuclear and mitochondrial markers were used to test the hypothesis that the Brycon sp1 and Brycon sp2 morphotypes are distinct species. Specimens from the two morphotypes were collected from the Lajeado Hydroelectric Plant and the Palmas River in the Araguaia-Tocantins Basin. Thirty-five loci obtained by the amplification of five inter-simple sequence repeat primers were analyzed but no species-specific bands were detected. Electrophoretic profiles obtained from 5S rDNA non-transcribed spacer amplification failed to show any differentiation in morphotypes. These results were corroborated by nucleotide sequence analysis of the mtDNA control region, in which 24 polymorphic nucleotide sites, representing a polymorphism rate of only 5%, were detected. The low rates of polymorphism detected by inter-simple sequence repeat,

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non-transcribed spacer and mtDNA D-loop markers strongly reject the hypothesis that the two morphotypes *Brycon* sp1 and *Brycon* sp2 represent distinct species within *Brycon cf. pesu*. Further studies are needed to obtain conclusive data on the notion that the coloration of the anal fin is an intraspecific polymorphism, possibly related to environmental factors.

Key words: *Brycon cf. pesu*; Araguaia-Tocantins Basin; Morphotypes; Inter-simple sequence repeat; 5S rDNA non-transcribed spacer; mtDNA