

Molecular characterization of *Leishmania* sp. (Kinetoplastida: Trypanosomatidae) from cases of cutaneous leishmaniasis in the Municipalities of Rio Preto da Eva and Manaus, Amazonas State, Brazil*

Caracterização molecular de *Leishmania* sp. (Kinetoplastida: Trypanosomatidae) proveniente de casos de leishmaniose cutânea dos Municípios de Rio Preto da Eva e Manaus, Estado do Amazonas, Brasil

Caracterización molecular de *Leishmania* sp. (Kinetoplastida: Trypanosomatidae) proveniente de casos de leishmaniasis cutánea de los Municipios de Rio Preto da Eva y Manaus, Estado de Amazonas, Brasil

Luanda de Paula Figueira

Programa de Pós-graduação Mestrado Multidisciplinar em Patologia Tropical da Universidade Federal do Amazonas, Manaus, Amazonas, Brasil

Antonia Maria Ramos Franco

Laboratório de Leishmaniose e Doença de Chagas, Coordenação de Pesquisas em Ciências da Saúde, Instituto Nacional de Pesquisas em Ciências da Saúde, Manaus, Amazonas, Brasil

Introduction: In the Amazon Region, deforestation resulting from the construction of roads and the establishment of new villages is often associated with epidemic outbreaks of diseases such as leishmaniasis. Because of the wide variety of leishmaniasis vector species and reservoir animals in this region, there is high genetic polymorphism in parasites of the genus *Leishmania*. **Objective:** This aim of this study was to identify the species of *Leishmania* isolated from humans in two municipalities of Amazonas State (Rio Preto da Eva and Manaus) and to evaluate their genetic polymorphisms. **Materials and Methods:** Twenty-three isolates were analyzed using Random Amplified Polymorphic DNA (RAPD) and isoenzyme electrophoresis. Eight loci were analyzed and compared to known diagnostic loci to assign species to each isolate, based on the similarity of bands. Interspecific variability among the samples was determined by RAPD and analyzed using phenetic analysis and genetic distance to establish the taxonomic relationships among the species/isolates of this genus. **Results:** Five different zymodemes were identified using a variety of numerical techniques and methods. The zymodeme analysis classified the samples into two species of the subgenus *Viannia*: *L. guyanensis* and *L. naiffi*. All *L. guyanensis* strains were clustered within the same zymodeme and were found to be polymorphic through RAPD analysis; *L. naiffi* showed similar patterns. **Conclusion:** The results revealed apparent isoenzyme homogeneity between samples of *L. guyanensis* and heterogeneity between samples of *L. naiffi*. This suggests that these species circulate in both municipalities. The RAPD showed that there was genetic variability among the isolate strains of *L. guyanensis* and *L. naiffi*. The relationship between the isolates and its consequences were analyzed, particularly with reference to the importance of selection on genetic markers.

Keywords: Polymorphism, Genetic, *Leishmania*, isoenzymes; Eletroforese; Random Amplified Polymorphic DNA Technique.

Financial Support: Instituto Nacional de Pesquisas da Amazônia (INPA) and Fundação de Amparo as Pesquisas do Estado do Amazonas (FAPEAM/Projeto Temático).

* Summary of dissertation submitted to the Post-Graduation Program in Tropical Pathology (Programa de Pós-graduação em Patologia Tropical) of the Universidade Federal do Amazonas under the guidance of Professor Antonia Maria Ramos Franco, Ph.D., towards completion of the requirements for the Master's degree on March 3, 2006. Manaus, Amazonas, Brazil.

Correspondência / Correspondence / Correspondencia:

Antonia Maria Ramos Franco

Instituto Nacional de Pesquisas da Amazônia, Laboratório de Leishmaniose e Doença de Chagas

Av. André Araújo, nº 2936 Manaus-Amazonas-Brasil

E-mail: afranco@inpa.gov.br

Recebido em / Received / Recibido en: 26/8/2010

Aceito em / Accepted / Aceito en: 29/3/2011