Mem. Inst. Oswaldo Cruz, Rio de Janeiro, Vol. 86 (4): 477-478 oct./dec. 1991

SCHIZODEME ANALYSIS WITH THE RESTRICTION ENDONUCLEASE RSA I DIFFERENTIATES BETWEEN TRYPANOSOMA RANGELI AND TRYPANOSOMA CRUZI

ANTONIO M. GONÇALVES; NÉDIA S. NEHME; NANCY SARAVIA*; IRIS SEGURA* & CARLOS M. MOREL

Instituto Oswaldo Cruz, Departamento de Bioquímica e Biologia Molecular, Caixa Postal 926, 20001 Rio de Janeiro, RJ, Brasil *Centro Internacional de Entrenamiento e Investigaciones Medicas (CIDEIM), Avenida 1a. Norte No. 3-03, Apartado Aereo 5390, Cali, Colombia

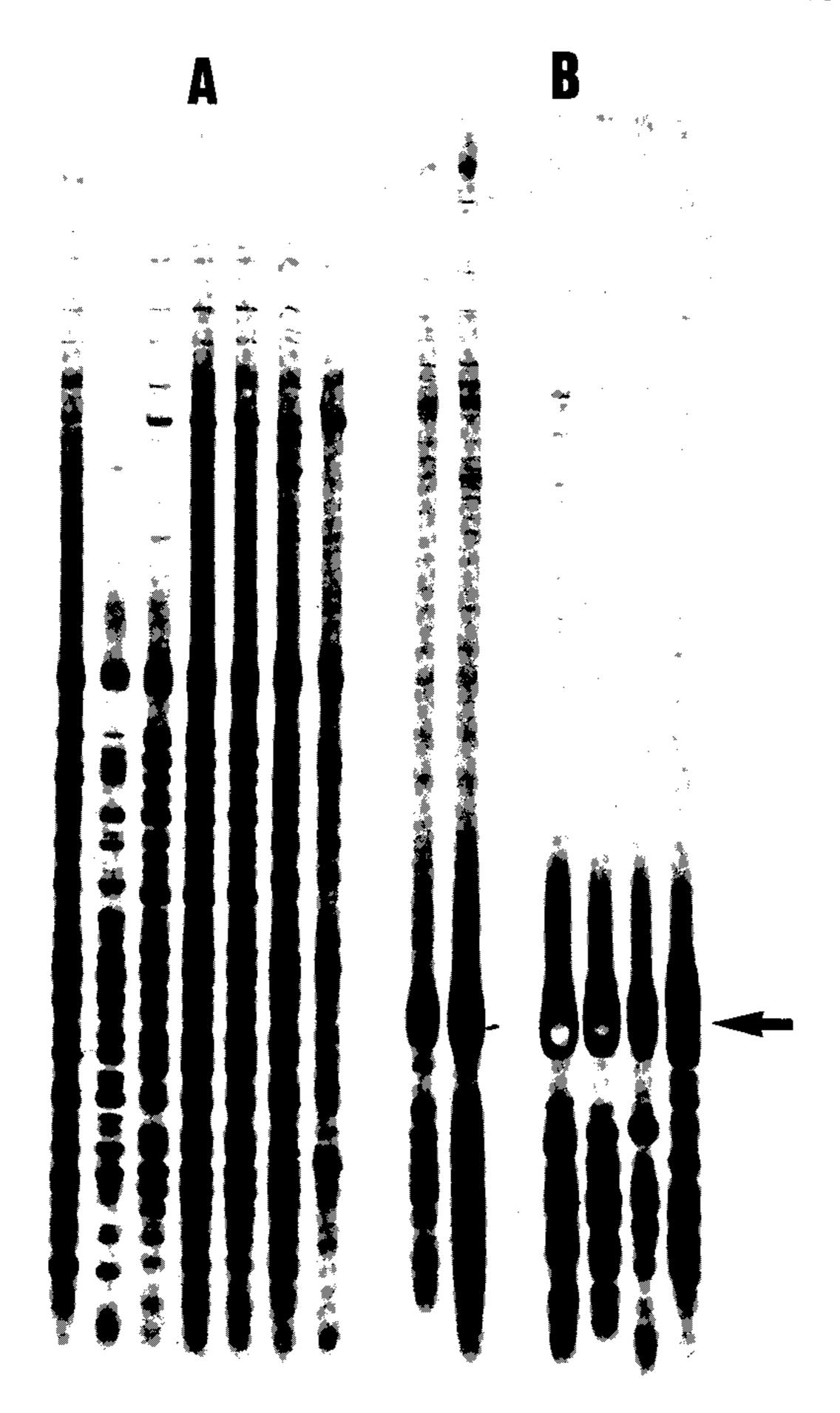
The differentiation between the two American trypanosomes Trypanosoma cruzi and T. rangeli is an important issue in the geographical areas where they coexist. For this purpose serology, morphology and behaviour in various biological systems need to be assessed (A. D'Alessandro, 1976, p. 328-393 in W. H. R. Lumsden & D. A. Evans (eds) Biology of the Kinetoplastida, Academic Press, London). Recent approaches have used monoclonal antibodies (L. Hudson et al., 1987, Acta Tropica, 44: 387-394) and complement lysis, lectin agglutination and isoenzyme profiles (M. Steindel et al., 1991, Mem. Inst. Oswaldo Cruz, 86: 73-79).

While characterizing several isolates of these parasites from Colombia and Brazil by schizodeme analysis (C. M. Morel et al., 1980, *Proc. Nat Acad. Sci. USA*, 77: 6810-6814), we found that they could be easily distinguished using the endonuclease *Rsa* I. The figure shows that

digestion of the kinetoplast DNA minicircles from T. cruzi with this enzyme originates a major band of around 350 base pairs (bp) (arrow). This is consistent with the restriction site for Rsa I (GT/AC) being located in the minirepeat, a constant sequence of 120 bp repeated four times along the 1400 bp minicircle molecule of T. cruzi (W. Degrave et al., 1988, Mol. Biochem. Parasitol., 27: 63-70). By contrast, a similar treatment of T. rangeli kinetoplast DNA originates a more complex restriction fingerprint with fragments distributed over a wide range of molecular weight.

DNA sequence analysis of kinetoplast DNA minicircles from *T. rangeli* is being carried out in our laboratories to investigate their structure and determine the distribution of *Rsa* I sites. The present results, however, already indicate that *Rsa* I schizodeme analysis can be a simple and reliable method for the differentiation of this parasite from *T. cruzi*.

This work received financial support from the UNDP/World Bank/WHO Special Programme for Research and Training in Tropical Diseases (TDR), CNPq and COLCIENCIAS.



Rsa I schizodeme analysis of isolates of Trypanosoma rangeli and T. cruzi. Kinetoplast DNA samples were prepared and processed as described by C. M. Morel et al. (1980, Proc. Nat. Acad. Sci. USA, 77: 6810-6814) and A. M. Gonçalves et al. (1984, p. 95-109 In C. M. Morel Genes and Antigens of Parasites, Oswaldo Cruz Foundation, Rio de Janeiro) using Rsa I from New England Biolabs. The restriction fragments were analyzed on a 6-10% polyacrylamide silver-stained gel according to A. M. Gonçalves et al. (1990, Mem. Inst. Oswaldo Cruz, 85: 101-106). A: T. rangeli (seven isolates from Colombia, two from human cases, five from the insect host Rhodnius prolixus). B: T. cruzi (three isolates from Brazil – two from human cases, one from a triatomine; three isolates from Colombia – one from a human case and two from Didelphis marsupialis). The arrow indicates the fragment of MW = 350 bp (1/4 th the MW of a minicircle).