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## Natural infection by *Trypanosoma cruzi* in dogs located in Ituberá, Southern Bahia, Brazil

### Infecção natural por *Trypanosoma cruzi* em cães domiciliados de Ituberá, Sul da Bahia, Brasil

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#### Resumo

A Doença de Chagas, causada pelo protozoário flagelado *Trypanosoma cruzi*, é uma antropozoonose de grande importância para a saúde pública. Diversas espécies de mamíferos são reservatórios do parasita, incluindo o cão. O objetivo deste trabalho foi verificar a infecção natural por *T. cruzi* na população de cães do município de Ituberá, Bahia. Foram avaliados 392 cães domiciliados, em todos os bairros do município, dos quais foram coletados 5mL de sangue para realização do diagnóstico molecular. A amplificação do DNA de *T. cruzi* foi através da Reação em Cadeia da Polimerase (PCR), com os primers P35 e P36, que amplificam um fragmento de 330 pb. Os produtos das PCRs foram submetidos à eletroforese em gel de agarose a 2% contendo Sybr (Invitrogen®). Dos 392 cães avaliados, apenas 2 (0,51%) animais, um macho e uma fêmea, foram positivos no diagnóstico molecular de *T. cruzi*. Conclui-se com esse estudo que há cães naturalmente infectados pelo *T. cruzi* no município de Ituberá-Bahia e que este achado constitui um alerta aos veterinários, profissionais da saúde e autoridades sanitárias locais, cujos cães podem atuar como reservatórios da doença.

**Palavras-chave:** *Canis familiaris*. Diagnóstico molecular. Doença de Chagas. Tripanossomíase.

#### Abstract

Chagas disease, caused by the flagellate protozoan *Trypanosoma cruzi*, is an anthroponosis of great importance for public health. Several species of mammals are reservoirs for this parasite, including dogs. The objective of this work was to verify the natural infection by *T. cruzi* in the population of dogs of the municipality of Ituberá, Bahia. A total of 392 domiciled dogs from all districts of the city were evaluated; five milliliters of blood was collected from the dogs for molecular diagnosis. *T. cruzi* DNA was amplified through the polymerase chain reaction (PCR); the primers P35 and P36, which amplify a fragment of 330 bp, were used. The PCR products were subjected to 2% agarose gel electrophoresis containing Sybr (Invitrogen). Of the 392 dogs evaluated, only 2 (0.51%) animals, one male and one female, tested positive for *T. cruzi*. This study concluded that there are dogs naturally infected by *T. cruzi* in the municipality of Ituberá-Bahia and that this finding is an alert to veterinarians, health professionals, and local health authorities that their own dogs can act as reservoirs of the disease.

**Key words:** *Canis familiaris*. Molecular diagnostics. Chagas disease. Trypanosomiasis.

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Chagas' Disease or American Trypanosomiasis (COURA; BORGES-PEREIRA, 2010) is caused by the protozoan, *Trypanosoma cruzi* (SOUZA et al., 2010). This anthroponosis is endemic in some Latin American countries; however, there has been an increased detection of the parasite in non-endemic countries, such as the United States, Canada, several European countries, and some Western Pacific countries (WHO, 2017).

Transmission to vertebrate hosts occurs mainly after stinging and blood ingestion, through direct contact with feces (BAHIA et al., 2002) contaminated with hematophagous insects of the *Reduviidae* family (SOUZA et al., 2010), and by oral and transplacental transmission (OPAS, 2009), blood transfusion, and organ transplantation (SOUZA et al., 2010).

It is estimated that 6-7 million people in the world are infected with *T. cruzi* (WHO, 2017). This chronic disease affects 2-3 million people in Brazil and has become a major public health problem in the country (MINISTÉRIO DA SAÚDE, 2014). An epidemiological study conducted in Brazil between 1975 and 1980 by Camargo et al. (1984) showed that the states with the highest seroprevalence of infection in the country were Rio Grande do Sul (8.84%), Minas Gerais (8.83%), Goiás (7.40%), Sergipe (5.97%), and Bahia (5.44%). A more recent study indicates that Bahia still has a high prevalence of *T. cruzi* infection in humans with a seroprevalence that varies from 5.4-25% (ARAS et al., 2002).

Previous studies have already shown that dogs are naturally infected by this protozoan (GÜRTLER et al., 2007; LEÇA-JÚNIOR et al., 2013). The main clinical signs observed in these animals are anorexia, fever, lymphadenopathy, and cardiac alterations such as myocarditis, tachycardia, severe cardiac arrhythmia (CAMACHO, 2003; NELSON; COUTO, 2006), congestive heart failure, and diastolic dysfunction (PASCON et al., 2010; SANTANA et al., 2012).

The municipality of Ituberá is endemic for

Chagas disease (SESAB, 2013). Thus, the dogs of this region may act as reservoirs and sentinels of the disease, and play an important epidemiological role. The objective of the present study was to verify the natural infection by *T. cruzi* in the population of dogs in the municipality of Ituberá-BA.

The municipality of Ituberá is located in the Southern Lowlands of Bahia state (13° 43' S 39° 08' O). The region has a hot climate with average temperatures of 25.3 °C and an annual thermal amplitude of 5.6 °C, with annual rainfall varying between 1800 and 2400 mm, distributed throughout the year. The biome of the municipality is represented by the Atlantic forest. The municipality has an approximate population of 29,273 inhabitants, a total area of 417 km<sup>2</sup> and density of 63.73 inhabitants/km<sup>2</sup> (IBGE, 2016).

The sample size was calculated using the Epi Info 3.5.2 software, with an expected prevalence of 50%, a 5% error, and a 95% confidence level considering the size of the canine population (392 dogs) is 12% of the population of the municipality (CIFUENTES, 1988). The samples were collected homogeneously from all the districts of the municipality, covering both the rural and urban areas from July to November 2015; this collection was based on the proportion of the population of each neighborhood in relation to the total population of the municipality. For each residence visited, a maximum of two dogs were examined. The study was approved by the Ethics Committee on Animal Use (CEUA) of the State University of Santa Cruz, under protocol 028/12.

Five milliliters of blood were collected by puncturing the cephalic or jugular vein of dogs. The samples were conditioned in tubes with anticoagulant (EDTA), refrigerated, and sent to the Veterinary Genetics Laboratory of the State University of Santa Cruz (UESC), where they were later subjected to molecular biology procedures.

Blood stored in EDTA tubes was centrifuged to obtain the leukocyte layers and the DNA was extracted by an Easy-DNA kit (Invitrogen). DNA

was stored at -20 °C prior to use. The DNA concentration of each sample was quantified with a NanoDrop 2000 (Thermo Scientific) for subsequent PCR. The mean and standard deviation of the DNA concentration were 140.96 and 115.08, respectively, and the mean and standard deviation of the purity of the samples were 1.98 and 0.38, respectively.

First, the integrity of the DNA was verified by using primers to amplify enzyme GAPDH (glyceraldehyde 3-phosphate dehydrogenase) (5' CCAAAGTTGTCATGGATGACC 3' and 5' CCTTCATTGACCTCAACTACAT 3'), which amplify a 400-bp fragment, as described by Birkenheuer et al. (2003). The reagents used were obtained from Invitrogen (Carlsbad, California, USA).

The *T. cruzi* DNA was amplified using primers P35: 5' AAATAATGTACGGGGGAGATGCATGA 3' and P36: 5' GGGTTCGATTGGGGTTGGTGT 3', which amplify a 330-bp fragment from the region of the mini-circle of mitochondrial DNA (kDNA) (AVILA et al., 1990). The reaction conditions used were similar to those described by Avila et al. (1990). Reaction mixtures at a final volume of 25 µL were composed of 1.0X Taq DNA polymerase buffer, 1.5 mM of MgCl<sub>2</sub>, 2 mM of each dNTP, 10 pmol of each primer, 1.5 U of Taq DNA polymerase, and 100 ng of genomic DNA. Invitrogen reagents (Carlsbad, California, USA) were used. The PCR conditions were as follows: denaturation at 94 °C for 1 min, 61 °C for 1 min for primer annealing, and final extension at 72 °C for 1 min for a total of 35 cycles.

DNA extracted from the epimastigote forms of the *T. cruzi* strain Tc II (former strain y) was used as a positive control. This DNA was kindly provided by researcher, Danielle Oliveira dos Anjos, from the Gonçalves Muniz Research Center (FIOCRUZ). Ultrapure water was used as a negative control.

The PCR products were subjected to 2% agarose gel electrophoresis and probed with Sybr (Invitrogen). The presence of bands was

analyzed with the aid of a transilluminator (Loccus Biotecnologia). All extracted DNA samples were positive in the PCR for the GAPDH enzyme.

Of the 392 dogs evaluated in the present study, 2 (0.51%), a male and a female, both from the urban area, tested positive.

There are few studies on the natural infection of dogs by *T. cruzi*; most research has been carried out on experimental infections and used serological tests as a diagnostic technique (BAHIA et al., 2002). Serological tests have been used due to the high costs of the PCR technique, which is used only for experimental purposes or to confirm discrepant serological results (OPAS, 2009). However, PCR results showed a low prevalence (0.5%) of natural *T. cruzi* infection in dogs, corroborating the results obtained by Leça-Júnior et al. (2013), who also investigated the natural infection in domiciled dogs with the same technique and found a 0.7% infection rate.

It should be noted that according to Avila et al. (1990) and Chiari (1999), the PCR technique used is very sensitive for diagnosing infections in patients with chronic Chagas disease, and its sensitivity varied from 96 to 100% when compared to serological tests, for example (AVILA et al., 1993; WINCKER et al., 1994).

The low positivity observed in this study may be related to the fact that blood was collected only once from each animal and a single PCR reaction of the sample was performed. According to Araújo et al. (2002), these factors reduce the sensitivity of the technique. Another possibility is that some infected animals in this study were in the chronic phase of the disease with parasitemia below the detection threshold of the PCR reaction used or that they were in fact not infected. The *T. cruzi* positivity of these domiciled dogs has great epidemiological importance, since these animals can act as reservoirs of the parasite and are important sources of feeding for *Triatoma infestans* and other triatomines, attracting these insects into

the domestic environment (GÜRTLER et al., 2007).

According to Gonçalves et al. (2012), of the 62 species of triatomines in Brazil, 25 are present in the state of Bahia; among these, the main vector species of Chagas disease in Brazil can be found, namely, *Panstrongylus megistus*, *Triatoma brasiliensis*, *Triatoma sordida*, *Triatoma pseudomaculata*, and *T. infestans*.

The region of Ituberá is located in an area that presents a transmission risk of Chagas disease (SESAB, 2013) and is home to remnants of the main vector, *T. infestans*, even after intense vector control in Brazil (MINISTÉRIO DA SAÚDE, 2014). In addition to *T. infestans*, there are other triatomine species in the Northeast region, such as *Triatoma tibiamaculata*, which was previously considered to be distributed in areas of wilderness but is currently found in households infected with *T. cruzi*; this suggests a potential risk of disease transmission to the inhabitants of the region (RIBEIRO JÚNIOR et al., 2015).

The results of this work lead to the conclusion that there are dogs naturally infected with *T. cruzi* in the municipality of Ituberá-Bahia; therefore, there is a possibility that zoonosis may occur from these infected dogs.

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