Molecular detection of *Trypanosoma cruzi* in *Mus musculus* in a rural town in Mérida, Yucatán, México

Detección molecular de *Trypanosoma cruzi* en *Mus musculus* en una localidad rural de Mérida, Yucatán, México

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American trypanosomiasis is a disease caused by the protozoan *Trypanosoma cruzi* and affects approximately 6 million people in the Americas. Commensal rodents are a food source for the vector and potential reservoirs of *T. cruzi*. The objective of this study was to estimate the prevalence of *T. cruzi* in commensal rodents that inhabit the rural town of Molas, Yucatán, México. Rodents were captured in households using Sherman traps. DNA extracted from heart samples from the captured rodents was used to detect *T. cruzi* using a PCR test. The prevalence of *T. cruzi* was compared by considering the sex and age of the rodents with the Chi-square test. A total of 114 *Mus musculus* mice were captured. The prevalence of *T. cruzi* was 21.1 % (n = 24) in the individuals examined. The comparison of the prevalence of *T. cruzi* between males and females and between adults and juveniles of *M. musculus* did not show statistically significant differences (P > 0.05). The prevalence of *T. cruzi* in *M. musculus* was high compared to other previous studies in México, and infection was observed regardless of rodent sex and age. These results show that *M. musculus* participates in the biological cycle of *T. cruzi* at Molas. Further studies are needed to understand the type of involvement (e.g., reservoir) of these rodents in the transmission dynamics of this parasite.

Key words: American trypanosomiasis; commensal rodents; molecular biology; southeastern México; tropical.

La tripanosomiasis americana es una enfermedad causada por el protozoario *Trypanosoma cruzi* y afecta aproximadamente a 6 millones de personas en las Américas. Los roedores comensales son una fuente de alimento para el vector y potenciales reservorios de *T. cruzi*. El objetivo del presente estudio fue estimar la prevalencia de *T. cruzi* en roedores comensales de la localidad rural de Molas, Yucatán, México. Se capturaron roedores en viviendas usando trampas Sherman. El DNA extraído de muestras de corazón de los roedores capturados fue utilizado para la detección de *T. cruzi* por medio de la técnica de PCR. Se comparó la prevalencia de *T. cruzi* considerando el sexo y edad de los roedores con la prueba de Chi-cuadrada. Un total de 114 ratones *Mus musculus* fueron atrapados. La prevalencia de *T. cruzi* fue 21.1 % (*n* = 24) en los individuos examinados. La comparación de las prevalencias de *T. cruzi* entre machos y hembras y entre los adultos y juveniles de *M. musculus* no arrojó diferencias estadísticas significativas (*P* > 0.05). La prevalencia de *T. cruzi* en *M. musculus* fue alta, en comparación con otros estudios previos en México, y la infección se presentó independientemente del sexo y la edad de los roedores. Estos resultados demuestran que *M. musculus* participa en el ciclo biológico de *T. cruzi* en Molas. Es necesario realizar mayores estudios para entender el tipo de participación (*e.g.*, reservorio) de estos roedores en la dinámica de transmisión de este parásito.

Palabras clave: Biología molecular; roedores comensales; sureste de México; tripanosomiasis americana; tropical.

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American trypanosomiasis, known as Chagas disease, is caused by the flagellated protozoan *Trypanosoma cruzi*. It is a life-threatening disease that has been neglected, which affects approximately 5,742,167 people and puts 70,199,360 people at risk of infection in Latin America (WHO 2015). *Trypanosoma cruzi* is transmitted through hematophagous hemipteran insects (vectors), such as *Rhodnius prolixus*, *Triatoma dimidiata*, and *Triatoma infestans*, accounting for 80 % of reported cases (OPS 2018). Vector transmission occurs when an infected vector feeds on a mammal and, after feeding, defecates on it, contaminating the feeding site or adjacent mucous membranes with parasites (WHO 2015). Other modes of transmission are transfusion of contaminated blood, congenital (vertical), transplantation of infected organs, and consumption of food contaminated with the parasite (Coura 2014).

Trypanosoma cruzi has been reported in more than 100 species of wild mammals, including marsupials, rodents, bats, armadillos, and primates (Jansen et al. 2017). In wild environments, rodents have been suggested to play a secondary role as reservoirs for this parasite due to the low infection frequencies reported (usually 0.4 %–5 %) compared to other mammals such as marsupials (11 %–90 %; Jansen et al. 2017). In anthropized environments, there are rodents known as commensals, such as the house mouse Mus musculus and the black rat Rattus rattus, which take advantage of food sources and shelter that favor large populations and infestations in households throughout the year (Battersby et al. 2008). This has suggested that commensal rodents may play an important role in domestic and peridomestic transmission of T. cruzi (Pinto et al.

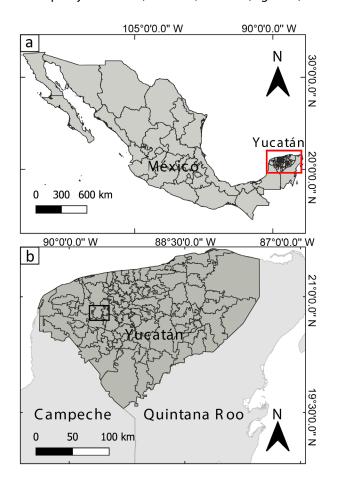
2006; Ayaqui and Ruelas 2020). There are reports of T. cruzi infection in M. musculus and R. rattus in these environments in Ecuador and Chile, with contrasting results. Although M. musculus was the rodent most frequently captured (n=79) in households in Guayas, Ecuador, all specimens tested negative, while in R. rattus (n=61), the second most captured species, the infection rate was 11.5 %. In Coquimbo, Chile, R. rattus (n=46) was captured more frequently than M. musculus (n=5), but both species showed similar infection rates, 83.6 % and 83.3 %, respectively.

In the city of Mérida, Yucatán, México, and its rural surroundings, people positive for *T. cruzi*, have been reported, as well as vectors (Guzmán-Tapia et al. 2007; García-Montalvo 2011) and the rodents *M. musculus* and *R. rattus* (Panti-May et al. 2017) also positive for *T. cruzi*. Although experimental infections with this protozoan in *M. musculus* are abundant, few studies have reported the natural infection rate in this rodent compared to *R. rattus* (Jansen et al. 2017). Therefore, the present study aimed to determine the prevalence of *T. cruzi* infection in *M. musculus* living in a rural town in Yucatán, México.

As part of a multidisciplinary study on zoonotic pathogens (<u>Pacheco-Castro et al. 2013</u>), commensal rodents were sampled in the rural town of Molas (20° 48' 55.42" N, 89° 37' 48.40" W), ca. 8 km south of the city of Mérida, in the municipality of Mérida, Yucatán, México (Figure 1). During

2009–2010, 200 Sherman traps were placed monthly in 20 households for 3 consecutive nights. The bait used was a mixture of oat flakes and artificial vanilla essence. The captured rodents were euthanized with sodium pentobarbital administered intraperitoneally (Leary et al. 2020). Subsequently, a mid-thoracic incision was performed to remove the heart, which was preserved in 10 % buffered formalin for subsequent molecular testing and histopathology analyses. The species, sex, and age of each rodent specimen were recorded; age was determined according to Panti-May et al. (2012). All field and laboratory protocols were approved by the Bioethics Committee of the Campus of Biological and Agricultural Sciences, Autonomous University of Yucatán (protocol no. CB-CCBA-L-2009-001).

From each heart sample, a 0.5 g section was obtained and placed in 200 µL of phosphate buffer (PBS) and incubated at room temperature for 24 hr. Afterward, the PBS was discarded, and the DNA was purified with the Insta-Gene™ Matrix resin (Bio-Rad®, Hercules, California) following the manufacturer's protocol. For the detection of *T. cruzi*, primers 121 and 122 were used, which amplify a region of the kinetoplast DNA (kDNA) of *T. cruzi* of approximately 330 base pairs (Wincker et al. 1994). Each PCR reaction included 50-100 ng of DNA, 10 µL of GoTaq Green Master Mix (Promega, Madison, Wisconsin), 200 ng of each primer and nuclease-free water to a final volume of 20 µL.



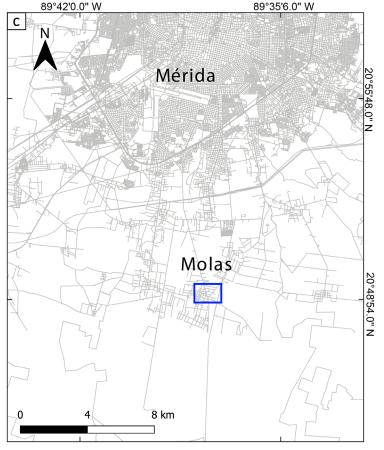


Figure 1. Geographic location of the town of Molas, Yucatán, México. a) Location of the state of Yucatán in México (red box); b) location of the city of Mérida (black box) in Yucatán; c) location of the town of Molas (blue box) in relation to the city of Mérida.

The amplification conditions were 5 min at 94 °C; 35 cycles of 94 °C for 60 seconds, 57 °C for 60 seconds, and 72 °C for 60 seconds; and a final extension at 72 °C for 10 minutes. Positive controls (*T. cruzi* DNA) and negative controls (molecular-grade water) were included in each reaction. Electrophoresis of PCR products was performed on 1.5 % agarose gels stained with ethidium bromide. The results were visualized and recorded on a photo-documenter (Bio-Rad®, Hercules, California).

The prevalence of *T. cruzi* infection and its 95 % confidence intervals (CI) were estimated with the Quantitative Parasitology 3.0 program (Rózsa et al. 2000). Likewise, the prevalence of infection was compared between host sexes and between age groups using the Chi-square test and the program mentioned above.

A total of 114 individuals of M. musculus were captured, including 47 females and 67 males; of these, 89 were adults and 25 were juveniles. Twenty-four samples of M. musculus (21.1 %, CI = 14.4 %–29.7 %) tested positive for *T. cruzi* (Figure 2). The comparison of the prevalence of *T. cruzi* between males (25.4 %, CI = 16.3 %-37.2 %) and females (14.9 %, CI = 7.1 % - 28.5 %), was not significantly different $(\chi 2 = 1.8, d.f. = 1, P = 0.18)$. In addition, no significant differences were found ($\chi 2 = 3.3$, d.f. = 1, P = 0.07) in the infection rate with *T. cruzi* between adults (27.7 %, CI = 16.7 % - 34.8%) and juveniles (8 %, CI = 1.5 %–25.6 %).

The presence of *T. cruzi* in *M. musculus* through molecular methods has been reported in some regions of the Americas. The prevalence of *T. cruzi* found in *M. musculus* in Molas (21.1 %) is higher than the 6.8 % reported in the United States of America (Herrera et al. 2015), but lower than 83.3 % in Chile (Yefi-Quinteros et al. 2018). In México, a 7.9 % prevalence of *T. cruzi* in this rodent was reported in the state of Morelos (Ramsey et al. 2012). Particularly in the city of Mérida, Panti-May et al. (2017) determined the infection of M. musculus with T. cruzi in an urban and a suburban area located ca. 11 km and 8 km north of Molas, respectively. In the urban area of Mérida, no specimens of M. musculus were positive for T. cruzi, while in the suburban area, the infection rate was 15.7 %. Another study conducted in Yucatán investigated T. cruzi infection in M. musculus and R. rattus living in a rural community in the municipality of Cenotillo; no specimens of M. musculus were positive (Hernández-Cortazar et al. 2018). This shows the wide variability of infection with *T. cruzi* in commensal rodents, even within the same region.

The present study recorded infection with *T. cruzi* in M. musculus regardless of rodent sex or age. This involves the exposure of this rodent to the parasite through several transmission routes, such as vectorial, oral, vertical, and sexual. Blood from M. musculus has been found in the gut of vectors such as T. dimidiata in Guatemala

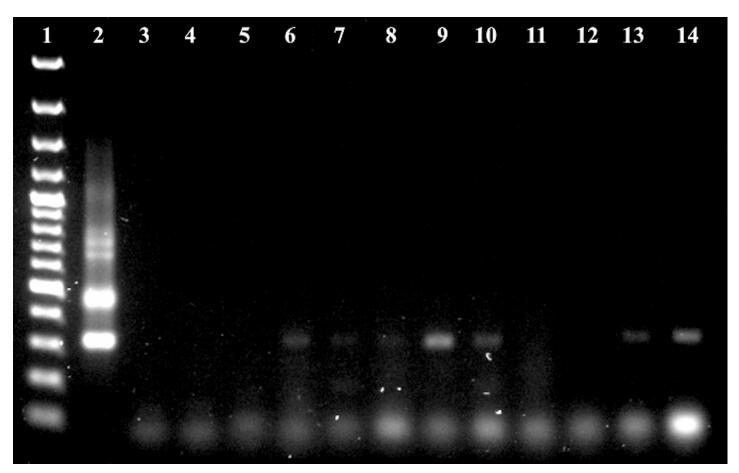


Figure 2. 1.5 % agarose gel showing the amplified 330 base pairs (bp) corresponding to the Trypanosoma cruzi kDNA in Mus musculus in Molas, Yucatán, México. Lane (L) 1, 100-bp molecular weight marker; L2, positive control; L3, negative control; L4, L5 and L12, negative samples; L6-L10, L13, L14, positive samples.

(Bustamante et al. 2014) and México (Torres-Montero et al. 2012) and Panstrongylus geniculatus in Venezuela (Segovia et al. 2023). The foregoing shows that M. musculus is a food source for vectors, which can promote vector-borne transmission. Experimental studies have shown that M. musculus can be infected through oral inoculum with blood and metacyclic trypomastigotes of *T. cruzi* (Dias et al. 2013), which may favor the maintenance of the parasite in rodents through the consumption of vectors that carry the metacyclic forms and contact with the blood of mice infected with trypomastigotes during fights. Vertical and sexual transmission routes have been experimentally demonstrated in M. musculus, but it has been concluded that they do not contribute significantly to the transmission and maintenance of the parasite (Rios et al. 2018; Faral-Tello et al. 2023).

This study used heart tissue to detect *T. cruzi* DNA in naturally infected mice. Infection with *T. cruzi* is characterized by an acute phase in which the parasite is found in the blood and various tissues, including the heart, and a chronic phase in which the parasite infects several tissues, such as the heart, skeletal muscle, kidney, liver, and spleen (Cencig et al. 2011; Caldas et al. 2012). Although *T. cruzi* can parasitize multiple tissues, the parasite load can vary in the different infection stages, depending on the parasite strain, the inoculum, and reinfections (Andrade et al. 1999; Cummings and Tarleton 2003; Cencig et al. 2011).

Commensal rodents have been identified as potential sentinels of vector transmission of T. cruzi because they are susceptible to infection, can be a source of food for vectors, are abundant, easy to capture, and live in proximity with the local inhabitants (Pinto et al. 2006; Bustamante et al. 2014; Panti-May et al. 2017; Ayaqui and Ruelas 2020). Mus musculus is the most frequently reported commensal rodent species in the city of Mérida (Panti-May et al. 2012, 2016). However, R. rattus may be more abundant in areas with tree vegetation due to its semi-arboreal behavior (Battersby et al. 2008). Therefore, R. rattus should be considered in studies addressing pathogens in commensal rodents. The results obtained in the present work highlight the need to study the participation of M. musculus in the biological cycle of T. cruzi in the city of Mérida and the rest of the Yucatán state.

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