



Leishmania infantum in the reproductive organs of dogs

Diogo Tiago da Silva^{1,2}  Maria Luana Alves^{1,3}  Julio Cesar Pereira Spada^{1,2}
João Augusto Franco Leonel^{1,3}  Julia Cristina Benassi³  Wilma Aparecida Starke-Buzetti⁴
Helena Lage Ferreira^{1,3} Lara Borges Keid^{1,3} Rodrigo Martins Soares^{1,5}
Trícia Maria Ferreira de Sousa Oliveira^{1,3*} 

¹Programa de Pós-graduação em Epidemiologia Experimental Aplicada às Zoonoses, Departamento de Medicina Veterinária Preventiva e Saúde Animal, Faculdade de Medicina Veterinária e Zootecnia, Universidade de São Paulo (USP), Pirassununga, SP, Brasil. E-mail: tricia@usp.br.

*Corresponding author.

²Faculdade de Ciências Agrárias de Andradina, Fundação Educacional de Andradina, Andradina, SP, Brasil.

³Laboratório de Medicina Veterinária Preventiva Aplicada, Departamento de Medicina Veterinária, Faculdade de Zootecnia e Engenharia de Alimentos, Universidade de São Paulo (USP), Pirassununga, SP, Brasil.

⁴Departamento de Biologia e Zootecnia, Faculdade de Engenharia de Ilha Solteira, Universidade do Estado de São Paulo (UNESP), Ilha Solteira, SP, Brasil.

⁵Departamento de Medicina Veterinária Preventiva e Saúde Animal, Faculdade de Medicina Veterinária e Zootecnia, Universidade de São Paulo (USP), Pirassununga, SP, Brasil.

ABSTRACT: *Leishmania infantum* causes canine leishmaniasis. Using parasitological and molecular analyses, we identified *L. infantum* in the reproductive organs of male and female dogs. Using histochemistry, immunohistochemistry, and PCR, we examined tissue samples from the reproductive organs of 8 male dogs and 16 female dogs diagnosed with leishmaniasis. Despite the absence of macroscopic or microscopic lesions in these organs, we observed *L. infantum* amastigotes in tissue samples from the testis and the uterus. PCR and sequencing of these tissues revealed sequences that matched 100% with *L. infantum* DNA available at GenBank. The presence of *L. infantum* amastigotes and DNA in testicular and uterine tissue samples suggested that these organs can harbor the parasite without associated macroscopic or microscopic lesions, and this can be especially important in the vertical and venereal transmission of leishmaniasis in dogs.

Key words: histochemical, immunohistochemical, leishmaniasis, testis, uterus.

Leishmania infantum em órgãos reprodutivos de cães

RESUMO: *Leishmania infantum* é agente etiológico da leishmaniose canina. Por meio de análises parasitológicas e moleculares, a presença do parasita foi investigada em órgãos reprodutivos de cães machos e fêmeas. Amostras de tecidos dos órgãos reprodutivos de 8 cães machos e 16 fêmeas diagnosticados com leishmaniose foram avaliadas por histoquímica, imunohistoquímica e PCR. Apesar de não terem sido observadas lesões macroscópicas ou microscópicas nos órgãos reprodutivos desses cães, formas amastigotas de *L. infantum* foram observadas em amostras teciduais do testículo e útero. A PCR e o sequenciamento do DNA extraído desses tecidos revelaram sequências 100% idênticas a *L. infantum* depositadas no GenBank. Nossos resultados sugerem que os testículos e o útero podem abrigar o parasita, sem associação com lesões macroscópicas ou microscópicas, o que pode ter uma grande importância na transmissão venérea e vertical da leishmaniose entre cães.

Palavras-chave: histoquímica, imunohistoquímica, leishmaniose, testículo, útero.

INTRODUCTION

Leishmania infantum causes human and animal cases of visceral leishmaniasis. Dogs are the main epidemiological reservoir of the parasite in the domestic environment, and vectorial transmission by sand flies is the most important means of infection (SOLANO-GALLEGO et al., 2011).

Many infected dogs remain without clinical signs; however, canine leishmaniasis (CanL) is a systemic disease characterized by nonspecific clinical signs (SOLANO-GALLEGO et al., 2011). *L. infantum* infection or DNA has been reported in several canine tissues, organs, and body fluids (TAFURI et al., 2004; SOLANO-GALLEGO et al., 2011; SILVA et al., 2016; FERNANDES et al., 2019), including those

present in the genital system (DINIZ et al., 2005; SILVA et al., 2008; AMARA et al., 2009; MANNA et al., 2012; BOECHAT et al., 2016; OLIVEIRA et al., 2016a, 2016b; MAGRO et al., 2017). In particular, infection of the canine reproductive tract has received substantial attention since venereal and vertical transmission were reported (PANGRAZIO et al., 2009; SILVA et al., 2009). Early vertical transmission to the embryo in a pregnant dog infected with *L. infantum* was detected by molecular means (OLIVEIRA et al., 2017). In the United States, where vectorial transmission *L. infantum* has not been proven, cases of vertical transmission in dogs appear to be important for the maintenance of the parasite circulation in the population (BOGGIATTO et al., 2011; TOEPP et al., 2019). This phenomenon highlights alternative routes of *L. infantum* transmission that can be important for the maintenance of parasite communities in canine populations.

In this study, using parasitological and molecular methods, we searched for *L. infantum* amastigote forms and DNA in the reproductive organs of dogs diagnosed with leishmaniasis from an endemic area of Brazil.

MATERIALS AND METHODS

During an epidemiological survey conducted by the Zoonotic Disease Control Center (ZDCC) of Ilha Solteira, São Paulo, Brazil, investigators diagnosed *Leishmania* spp. infections in 24 dogs of several breeds, both sexes (8 males and 16 females) and various ages (mean age: 3.2 years). The detection methods included (a) direct parasitological examination of popliteal or pre-scapular lymph node aspirate stained with Panótico® staining kit (Laborclin); (b) immunochromatographic testing (*Dual Parth Platform* – DPP®, Biomanguinhos); and (c) indirect ELISA testing (Biomanguinhos). All dogs presented with clinical signs suggestive of CanL, including skin lesions, lymphadenopathy, loss of body weight, lethargy, mucous membrane pallor, ocular lesions, diarrhea, splenomegaly, and hepatomegaly (SOLANO-GALLEGO et al., 2011). The dogs' owner transported the dogs to the ZDCC for euthanasia in accordance with the dictates of the Brazilian visceral leishmaniasis control program. We collected tissue samples from reproductive organs of all 24 dogs as donations from the ZDCC. The study was approved by the Ethics Committee for Animal Use (CEUA) of Faculdade de Engenharia de Ilha Solteira, Universidade do Estado de São Paulo (FEIS/UNESP) under protocol number 06/2014-

CEUA.

We collected fragments of tissue approximately 1 cm in size from the reproductive organs (ovaries and uteruses from females and testes from males). We fixed some of the tissue in 10% formalin with 0.01 M phosphate-buffered saline for histochemical (HC) and immunohistochemical (IHC) analyses. We stored the remaining tissue in 1.5-mL sterile microtubes at -20 °C for molecular analyses.

About HC, the tissues were fixed and then embedded in paraffin. We created histological sections of 5 µm thickness and stained them with hematoxylin and eosin for histopathological examination. Then, we evaluated three sections of each tissue under an optical microscope at 40× or 100× magnification.

For IHC, we performed immunostaining of *L. infantum* amastigotes according to procedures described by TAFURI et al. (2004) with modifications by QUEIROZ et al. (2011). We hydrated the deparaffinized slides and incubated them in hydrogen peroxide (Fluka Chemie AG, CH-9470 Buchs, Switzerland) to block endogenous peroxidase activity. Next, we incubated the slides in normal goat serum (1:50 dilution) to block nonspecific immunoglobulin absorption in the tissue. The primary antibody was immune serum from dog that was naturally infected with *L. infantum* (ELISA and IFAT titers = 1:2.560). The secondary antibody was a biotinylated rabbit anti-IgG antibody produced in goat (BA-1000, Vector Laboratories, Inc., Burlingame, CA, USA). After appropriate washes, we incubated tissue sections in a solution of avidin-biotin-peroxidase complex (Vectastain ABC Kit PK 6200; Vector Laboratories Inc., Burlingame, CA, USA), followed by incubation with the peroxidase substrate (Nova REDTM substrate Kit SK-4800; Vector Laboratories Inc., Burlingame, CA, USA). Finally, we counterstained the sections with Mayer's hematoxylin and mounted them on glass slides. The tissue samples were positive when extra or intramacrophagic amastigote forms were specifically immunostained with red/brown color in at least one tissue section under an optical microscope (at 40× or 100×).

We performed DNA extraction from the tissue samples using the Illustra™ Blood Genomic Prep Mini Spin kit (GE Healthcare) in accordance with the manufacturer's recommendations.

The PCR reactions of each sample proceeded using the internal transcribed spacer 1 (ITS-1) region of the rDNA of the Trypanosomatidae family using the primers LITSR (5'-CTGGATCATTTTCCGATG-3') and L5.8S (5'-TGATACCACTTATCGCACTT-3') that

annealed to the conserved sequences SSU and 5.8S. as described by EL TAI et al. (2000). We used an *L. infantum* (syn. *Leishmania chagasi*, MCAN/BR/1984/CCC-17.481) DNA sample provided by the Fundação Oswaldo Cruz (FIOCRUZ), Rio de Janeiro, Brazil, as a positive control. Sterilized ultrapure water was the negative control. We performed PCR gel electrophoresis on 1.5% agarose gels and subjected PCR products to DNA sequencing as described by FERNANDES et al. (2019).

RESULTS

All 24 infected dogs presented some clinically suggestive signs of CanL. We recorded onychogryphosis (75%; 18/24), skin lesions (70.8%; 17/24), weight loss (62.5%; 15/24), lymph node enlargement (50%; 12/24), alopecia (45.8%; 11/24), splenomegaly (45.8%; 11/24), hepatomegaly (25%; 6/24), corneal opacity (25%; 6/24) and diarrhea (12.5%; 3/24). Dogs were euthanized at the ZDCC and tissue samples from the reproductive organs were collected and subjected to parasitological (HC and IHC) and molecular (PCR) methods to identify parasites.

First, it is important to highlight that we observed neither macroscopic nor microscopic lesions on the genital organs we sampled in this study. HC and IHC showed *L. infantum* amastigote forms present in the connective tissue of the seminiferous tubules from the testes of two dogs (25%; 2/8) (Table 1; Figure 1C-D). We observed amastigotes in the endometrium of the uterus in two females using IHC (12.5%; 2/16) (Table 1; Figure 1A-B). There were no *L. infantum* amastigote forms in ovarian tissue samples, according to HC and IHC (Table 1).

Tissue samples from 10 dogs (41.7%; 10/24), four testis tissue samples (50%; 4/8), and

six uterus tissue samples (37.5%; 6/16) were PCR positive for the ITS-1 region (Table 1). As in HC and IHC, none of the ovarian tissue samples were ITS-1 PCR-positive (Table 1). Upon sequencing, all sequences showed 100% identity with *L. infantum* (similarity with sequence deposited on GenBank under accession number KM677131.1).

DISCUSSION

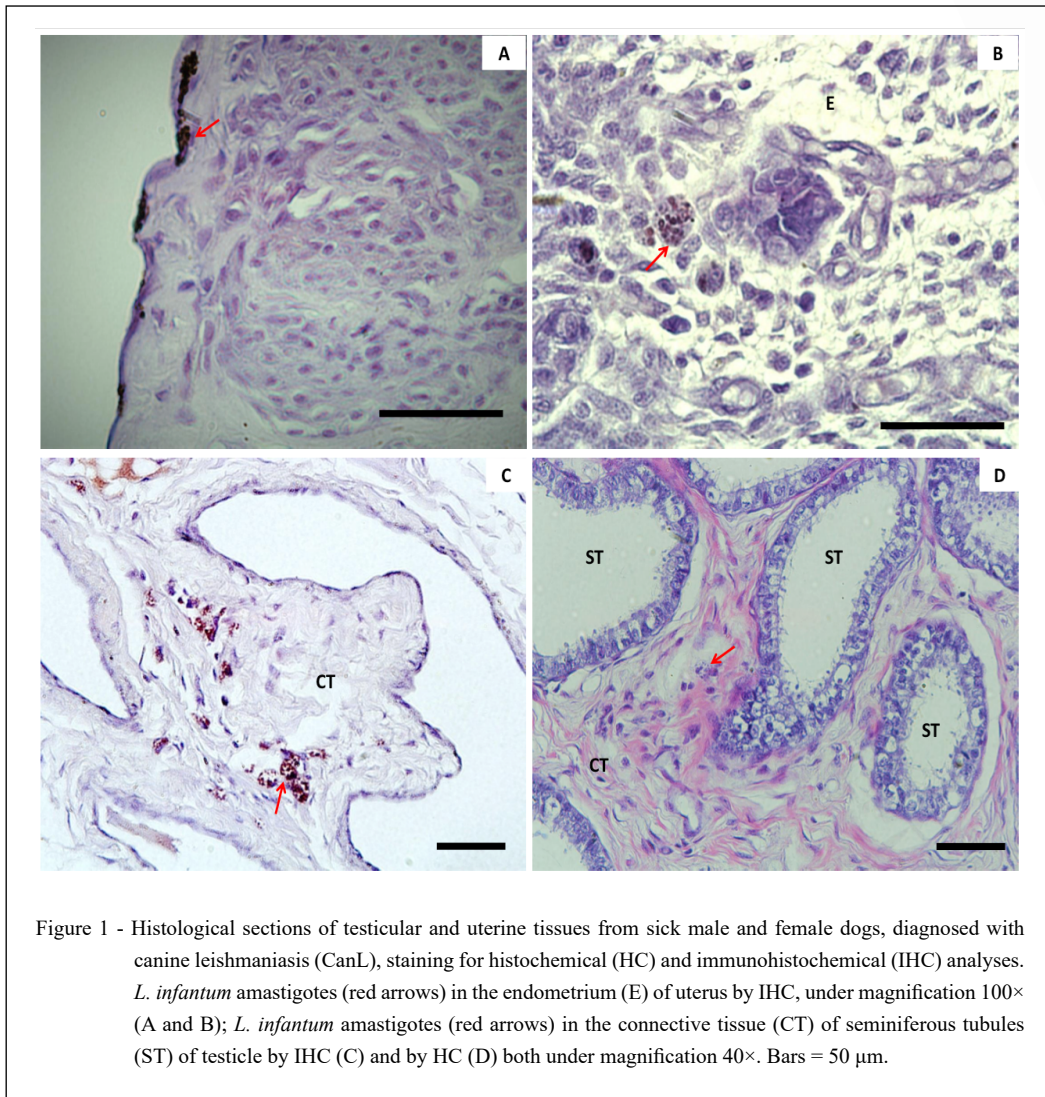
Ten dogs diagnosed with leishmaniasis were positive for DNA or amastigotes of *L. infantum* in their reproductive organs (testis and uterus), without macroscopic or microscopic alterations.

The presence of *Leishmania* spp. in the male genital system was reported by other authors in external (glans, urethral, prepuce, scrotum) and internal genitalia (testis, epididymis, prostate) of infected dogs without clinical signs or overt illness (DINIZ et al., 2005; AMARA et al., 2009; MANNA et al., 2012; BOECHAT et al., 2016; OLIVEIRA et al., 2016a). In the present study, we identified *L. infantum* amastigotes and DNA in tissue samples of testis from infected and sick male dogs (Figure 1C-D), without macroscopic or microscopic alterations observed in the testes. Orchitis was associated with amastigote forms in canine testicular tissues, where they probably acted as factors triggering chronic inflammatory responses (DINIZ et al., 2005; MANNA et al., 2012). BOECHAT et al. (2016) observed testicular degeneration, atrophy, absence of spermatogenesis, and necrosis related to the presence of the infection. It is clear that testis involvement could result in shedding of the amastigotes in the semen, favoring venereal transmission of the disease (MANNA et al., 2012; OLIVEIRA et al., 2016a).

In the female genital system, external (vulva, vagina, and mammary glands) and internal

Table 1 - *L. infantum* amastigotes and DNA detection by histochemical (HC), immunohistochemical (IHC), and PCR methods in the genital organs of sick male and female dogs, diagnosed with leishmaniasis.

Dogs	Number	Clinical signs	Organs	HC	IHC	PCR	Sequencing
Male	8	100% (8/8)	Testis	25% (2/8)	25% (2/8)	50% (4/8)	100% <i>L. infantum</i>
Female	16	100% (16/16)	Uterus	0	12.5% (2/16)	37.5% (6/16)	100% <i>L. infantum</i>
			Ovaries	0	0	0	-



genitalia (ovary, uterus, cervix) were reported to harbor *Leishmania* spp. in infected female dogs without clinical signs or overt illness (SILVA et al., 2008; BOECHAT et al., 2016; OLIVEIRA et al., 2016b; BOECHAT et al., 2020; MAGRO et al., 2017). In our study, we identified *L. infantum* amastigotes and DNA in uterine tissue samples from infected and sick female dogs (Figure 1A-B). Unlike the present study, *Leishmania* amastigotes forms and DNA were reported in ovaries in other studies (SILVA et al., 2008; BOECHAT et al., 2016; OLIVEIRA et al., 2016a).

In the present study, we reported a higher proportion of *L. infantum* infection in the reproductive organs of male dogs (50%) than in females (37.5%). Other studies suggested a tropism for the canine male

genital system, particularly the epididymis, prepuce, and glans penis, resulting in inflammation of these organs, with shedding of the organism into the semen. This contributed to venereal transmission of the parasite during copulation from male to female dogs (SILVA et al. 2008). In fact, the venereal transmission of *L. infantum* in dogs was demonstrated only from male to female (SILVA et al., 2009). However, BOECHAT et al. (2020) demonstrated similar *L. infantum* active infections in female and male dogs, with vagina, testis, vulva, and epididymis showing the highest frequencies of active infection.

In females, the presence of the infection in the uterus, without lesions, reinforces the notion that vertical transmission of canine leishmaniasis should receive more attention. These routes of infection

appear to be responsible for the maintenance of the parasite in areas of the United States where vectorial transmission is unproven, with a high risk of transmission from mother to offspring (BOGGIATTO et al., 2011; TOEPP et al., 2019).

Our study highlights the visceralization pattern of *L. infantum* infection reaching the canine reproductive organs from naturally-infected male and female dogs. The presence of *L. infantum* amastigotes and DNA in testicular and uterine tissue samples from dogs with CanL suggests that these organs can harbor the parasite without associated macroscopic or microscopic lesions.

CONCLUSION

We found evidence for *L. infantum* infection in the testis and uterus of dogs. Our findings reinforce the notion of venereal and vertical transmission as a means of transmission of CanL in endemic areas.

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BIOETHICS AND COMMITTEE APPROVAL BIOSSECURITY

The study was approved by the Ethics Committee for Animal Use (CEUA) of Faculdade de Engenharia de Ilha Solteira, Universidade do Estado de São Paulo (FEIS/UNESP) under protocol number 06/2014-CEUA.

DECLARATION OF CONFLICT OF INTERESTS

The authors declare no conflict of interest. The founding sponsors had no role in the design of the study; in the collection, analyses, or interpretation of data; in the writing of the manuscript, and in the decision to publish the results.

AUTHORS' CONTRIBUTIONS

All authors contributed equally to the design and writing of the manuscript.

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