

Ehrlichia canis morulae and DNA detection in whole blood and spleen aspiration samples

Detecção de mórulas e DNA de *Ehrlichia canis* em sangue e em aspirado de baço em cães naturalmente infectados

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Abstract

The aim of this study was to compare the detection of *Ehrlichia canis* morulae and DNA by nPCR in whole blood and spleen aspiration. The sample included 40 dogs showing thrombocytopenia associated to clinical signs suggestive of canine ehrlichiosis. Morulae detection showed that in 35 of the dogs studied, 17 had morulae in spleen tissue, and two in buffy coat smears. *E. canis* DNA was detected in 29/40 blood samples. We verified that morulae detection is more efficient in cytological preparations from spleen aspiration. On the other hand, nPCR on spleen and blood samples were equally efficient for disease diagnosis.

Keywords: *Ehrlichia canis*, morulae, spleen, nPCR, dogs.

Resumo

O objetivo desse estudo foi comparar a pesquisa de mórulas de *Ehrlichia canis* e a nPCR em sangue total e em aspirado de baço. Selecionaram-se 40 cães apresentando trombocitopenia associada a sinais e sintomas sugestivos de erliquiose canina. A pesquisa de mórula mostrou que dentre 35 amostras, 17 apresentaram mórulas nas preparações do baço, e duas nos esfregaços feitos a partir da papa leucocitária. O DNA de *Ehrlichia canis* foi detectado em 29 de 40 amostras de baço e em 30 de 40 no sangue. No presente estudo observou-se que a pesquisa de mórula é mais eficiente nas preparações citológicas obtidas da punção aspirativa do baço e que tanto a nPCR de baço quanto a de sangue foram eficientes no diagnóstico da doença.

Palavras-chave: *Ehrlichia canis*, mórula, baço, nPCR, cães.

Introduction

Canine ehrlichiosis is an infectious disease caused by the gram-negative bacterium *Ehrlichia canis* and transmitted by the brown dog tick *Rhipicephalus sanguineus* (LEWIS et al., 1977). The disease is reported in many countries, being more common in tropical and subtropical regions (NEER; HARRUS, 2006).

The incubation period of canine ehrlichiosis can range from 8 to 20 days (CASTRO et al. 2004; NEER; HARRUS, 2006), followed by the acute, subclinical, and chronic phases of the disease (HARRUS et al., 1999). The acute phase lasts 2 to 4 weeks (NEER; HARRUS, 2006) and is characterized by fever, weight loss, anorexia, depression, lymphadenomegaly, splenomegaly, vasculites, and ocular and musculoskeletal signs (WANER et al., 1999; DAGNONE; MORAIS; VIDOTTO, 2001; CASTRO et al. 2004). Thrombocytopenia is the most common abnormality in naturally or experimentally infected dogs in this disease phase (WANER et al., 1995).

The subclinical phase has a duration that varies from months to years (CODNER; FARRIS-SMITH, 1986; NEER;

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HARRUS, 2006). During this phase the animal could show high anti-*E. canis* antibody titers (WANER et al., 1997), persistent thrombocytopenia, and leucopenia with no other clinical signs (CODNER; FARRIS-SMITH, 1986; WANER et al., 1997; HARRUS et al., 1998a). It has been suggested that during this phase the organism stays housed in the animal's spleen (HARRUS et al., 1998a, 2004). Severe pancytopenia, hemorrhagic diatesis, and organism debilitation can characterize the chronic phase (NEER; HARRUS, 2006). Although the conditions that can lead the animal to this phase are not totally clear, it has been suggested that factors such as breed, immune system deficiency, stress conditions, co-infections, virulence strain, and geographical region could have some influence (HARRUS et al., 1998a).

Diagnosis can be done by direct identification of inclusion bodies or morulae of *E. canis* in leucocytes from blood smears (ELIAS, 1991), or from buffy coat smears and lymphonode aspiration (MYLONAKIS et al., 2003). Detection of specific antibodies by the immunofluorescent antibody test (IFAT) and Dot-ELISA (OLIVEIRA et al., 2000; CADMAN et al., 2004; NAKAGHI et al., 2008), and by molecular techniques such as nPCR (WEN et al., 1997; NAKAGHI et al., 2008) are also used as diagnostic methods.

This study sought to investigate whether the presence of morulae in splenic aspirate smears could be used as diagnosis of canine ehrlichiosis, since there is evidence that the spleen could shelter the parasite and participate in the pathogenesis of this disease. In addition, we also investigated the presence of parasite DNA in blood samples and spleen tissue of dogs with clinical suspicion of naturally occurring canine ehrlichiosis.

Material and Methods

1. Animals

Forty dogs suspected of being infected with *E. canis* cared for at the Veterinary Teaching Hospital of Universidade Estadual Paulista (UNESP) in Jaboticabal, SP, Brazil from April 2004 to May 2005 were used in this study. The inclusion criteria of these dogs were the presence of thrombocytopenia with one or more clinical signs of the disease: anorexia, apathy, splenomegaly, lymphadenopathy, uveitis, hemorrhagic diatesis, or tick infestation. From each dog we collected 10mL of blood, one aliquot with ethylenediaminetetraacetic acid (EDTA) to be used for the buffy coat smear and nPCR. The other aliquot without EDTA was used to obtain serum samples used to detect anti-*Babesia canis* antibodies by IFAT and anti-*E. canis* antibodies by Dot-ELISA. Spleen fine needle aspiration samples were used in morulae search and in DNA extraction for nPCR. All the procedures were performed with the dog owner's permission, and the rules of Comitê de Ética, Bioética e Bem Estar Animal were followed as instructed by the UNESP.

2. Morulae detection

Morulae detection was performed using the buffy coat blood smears and the splenic aspirated samples.

The fine needle aspiration was performed in 35 among 40 samples. The dogs were restrained by hand in dorsal decubitus and ultrasound-guided fine needle aspiration biopsies were performed, with a 2.0 mL sample collected with EDTA. Then sample smears were done and let dry at room temperature.

The buffy coat was obtained from 1.5 mL of EDTA whole blood sample transferred to a sterile microtube and centrifuged at 2,500 g for 10 minutes. The buffy coat was collected to make the smears in the same way as described for the spleen samples. Both the smears (spleen and buffy coat) were methanol fixed for 3 minutes and Giemsa stained. The slides were visualized by optic microscopic at 1,000×.

3. DNA extraction and nPCR

DNA extraction from blood and spleen samples was performed using QIAmp DNA Blood Mini kit (Qiagen®) according to manufacturer's instructions. The amplification was performed in two steps in a thermocycler (MJ-Research-PTC-200) as previously described (HARRUS et al., 1998a; MURPHY et al., 1998; NAKAGHI et al., 2008). The first round used 5.0 µL of template DNA in 5.0 µL PCR buffer 10× (100 mM Tris-HCl, pH 9.0, 500 mM KCl), 0.2 mM each dNTP, 2.5 mM MgCl₂ (Invitrogen®), 1 pmol each primer ECC (5'- GAACGAACGCTGGCGGCAAGC -3') and ECB (5'- CGTATTACCGCGGCTGCTGGCA -3'), 1.25 U of Taq DNA polymerase (Invitrogen®) and water to complete 50 µL. The PCR cycles consisted of 3 minutes at 94 °C, 30 cycles of 94 °C for 45 seconds, 65 °C for 2 minutes, 72 °C for 2 minutes, and a final extension at 72 °C for 5 minutes. The nPCR assay was performed using the same reaction conditions as the first amplification, but species-specific primers ECAN (5'- CAATTATTTATAGCCTCTGGCTATAGGA -3') and HE3 (5'- TATAGGTACCGTCAATTATCTTCCCTAT -3'), and 1.0 µL from the initial PCR was used as template. The PCR consisted of 3 cycles of 1 minutes at 94, 55 °C for 2 minutes, and 72 °C for 1 minute and 30 seconds, followed by 37 cycles of 1 minute at 92, 55 °C for 2 minutes, and 72 °C for 1 minute and 30 seconds, and a final extension at 72 °C for 5 minutes. The PCR products were visualized on a 1.5% agarose gel stained with ethidium bromide and UV light. The positive sample used as a control was obtained from a dog that was experimentally infected with *E. canis* Jaboticabal strain, and ultrapure water was used as negative control. The amplified products were measured using a 100 bp DNA ladder (Invitrogen®). DNA extraction and master-mix preparation were processed in different laboratories to avoid contamination.

4. Anti-*E. canis* and anti-*B. canis* antibodies detection by serology

In order to evaluate the presence of anti-*E. canis* antibodies we used the Immunocomb kit® (Biogal, Israel) following the manufacturer's instructions. To detect anti-*B. canis* antibodies, a IFAT protocol previously described by Furuta et al. (2009) was used. Briefly, 2.0 µL of serum samples were diluted (1:40, 1:80, 1:160, 1:320, 1:640, 1:1280) and were dropped in the marked

areas with *B. canis* antigen sensitized, and then incubated for 45 minutes in 32 °C in a humid chamber. Then followed a series of washes and a final drying step by addition of the conjugated (Sigma® - F7884) diluted in 1:80 according to manufacturer's instructions. All the slides were viewed in an optic microscope with fluorescent light (Olympus BX60), and the serum samples were considered positive when they reacted in dilutions greater than 1:40.

5. Statistical analyses

The results were analyzed using the chi-square test with 5% significance level in an SAS software program (Statistical Analysis System, 2006).

Results

1. Morulae detection

Among 35 samples collected from dogs with clinical signs of ehrlichiosis, 17 (48.6%) showed intracytoplasmic morulae in spleen aspiration (Figure 1) and two (5.7%) showed in buffy coat.

2. nPCR

The nPCR results showed that from all 40 samples analyzed, the *E. canis* DNA was isolated in 29 (72.5%) spleen samples and in 30 (75%) whole blood samples. The ehrlichial DNA was not isolated in 11 (27.5%) spleen samples and in 10 (25%) whole blood samples. These results can be seen in Table 1. The product of 398 base pairs corresponding to the amplified *E. canis* DNA is shown in Figure 2 (a and b). The results do not show a statistical difference ($p > 0.05$) in ehrlichial DNA detection in spleen aspirates and whole blood by the nPCR reaction, showing that both techniques are effective in the diagnosis of naturally occurring canine ehrlichiosis.

3. Presence of anti-*E. canis* and anti-*B. canis* antibodies

Amid the 40 samples evaluated, 33 (82.5%) presented antibody titers for *E. canis* and 28 (70%) for *B. canis*. Amid the 33 samples with *E. canis* titers, in two of them we could not isolate the bacteria DNA in blood and spleen samples and no morulae were found, except in one where morulae were found in spleen tissue and leukocyte smear.

4. Positivity frequency between diagnostic tests

In 30 positive nPCR whole blood samples, 28 were also spleen sample nPCR positive, 26 in Dot-ELISA, and 21 in anti-*B. canis* IFAT test. Fifteen samples showed morulae in spleen cytology, and one of these was also positive in buffy coat smear. Among 29 spleen nPCR positive animals, 28 were also whole blood nPCR positive, 22 in Dot-ELISA, and 22 in anti-*B. canis* IFAT.

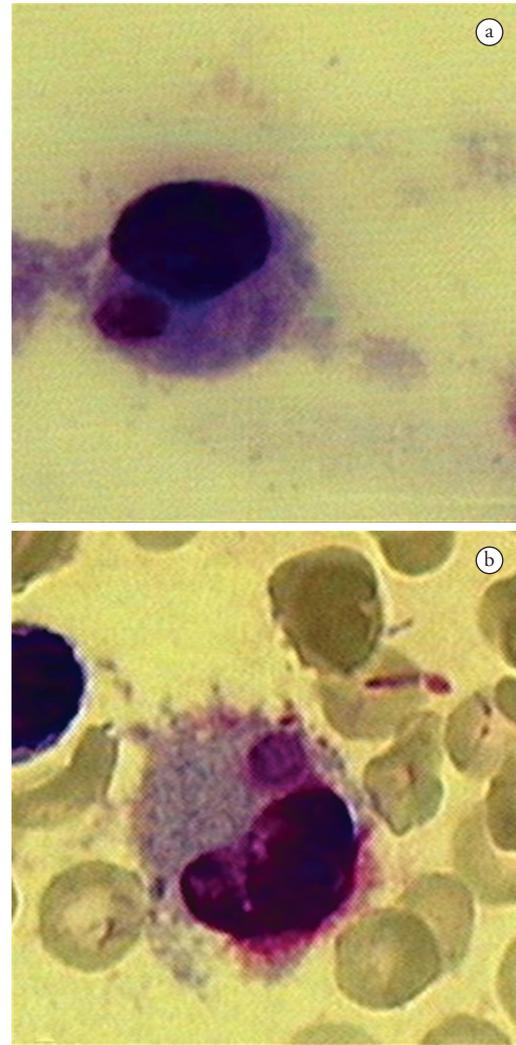


Figure 1. Photomicrography of cytological smears of spleen aspiration showing a morula in the mononuclear cells in a and b. 100x. Giemsa stained.

Table 1. Dogs naturally infected and positive or negative for *E. canis* DNA.

	Positive animals	Negative animals	Total of animals
nPCR (blood)	30	10	40
nPCR (spleen)	29	11	40

Fourteen had morulae in spleen cytology and one in buffy coat smear. These results can be seen in Table 2. From the 40 evaluated animals, two were positive only in Dot-ELISA, one in blood nPCR, one in IFAT, and none were positive only in spleen nPCR or morulae search.

Among 17 samples in which morulae was detected in spleen aspiration, 15 were also nPCR positive in blood samples, 14 were nPCR positive in spleen samples and in Dot-ELISA, and 10 in anti-*B. canis* IFAT.

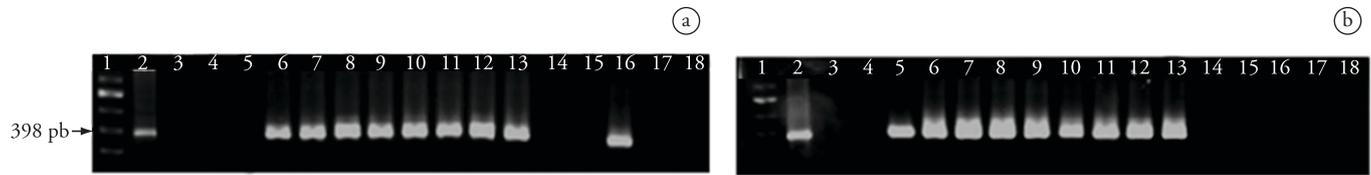


Figure 2. Amplified products by nPCR to detect *E. canis* DNA in spleen aspirates a) and whole blood b) samples from dogs with clinical suspicion of ehrlichiosis. Lane 1 molecular size ladder (100 pb), lane 2 positive control, lane 3 negative control (ultrapure water), lanes 4 to 18 dog samples.

Table 2. Positive animals in molecular, serological, and parasitological diagnostic tests in dogs with naturally occurring ehrlichiosis.

PCR	Positive animals	
nPCR b + nPCR s	28/30	
nPCRb (blood) (n = 30)	nPCR b + Dot-ELISA	26/30
	nPCR b + IFAT	21/30
	nPCR b + Morulae	15/30
nPCR s (spleen) (n = 29)	nPCR s + nPCR b	28/29
	nPCR s + Dot-ELISA	27/29
	nPCR s + IFAT	22/29
	nPCR s + Morulae	14/29

Discussion

The results found in this study have shown that morulae search in spleen aspirates is more efficient for the parasite detection than the search in buffy coat smear. Other studies have also observed a low frequency of morulae in buffy coat smears and they have attributed this feature to the low parasitemia that occurs under natural infection (ELIAS, 1991; HARRUS; BARK; WANER, 1997; FRANK; BREITSCHWERDT, 1999; OLIVEIRA et al., 2000). The low parasitemia could be due to the bacteria remaining lodged in the spleen longer than in the blood, justifying the higher number of morulae in this organ.

The high number of morulae-positive samples in spleen aspiration shown in this study is in disagreement with what Harrus et al. (1998a) previously reported. One possible explanation could be associated to the time of sample collection. While Harrus et al. (1998a) evaluated dogs during the experimental and sub-clinical phase of the disease, in our study we studied naturally infected animals that could be presenting any of the three possible phases of the disease. Even though some animals in our study could also be presenting the sub-clinical phase of the disease, the knowledge we currently have on how the bacteria behaves in naturally infected animals and on its cycle in the host is still poor.

Some researchers believe that the spleen could be the shelter for ehrlichial organisms for periods longer than the bacterial time in the blood (FOLEY et al., 1999; HARRUS et al., 2004). This feature could justify the higher number of morulae detected in the spleen in this experiment. However the results of this search didn't show better results in the detection of ehrlichial DNA in spleen than in blood and they disagree with the results of Harrus et al. (1998a). These researchers showed superiority in the ehrlichial DNA detection extracted from spleen of dogs in the subclinical

phase of the disease. In another study, Harrus et al. (2004) showed that the early detection of the disease after infected blood inoculation in dogs as equally positive in samples of spleen and of blood. In later evaluations the spleen samples showed superiority over the blood samples in these same dogs after sixteen days of doxycycline therapy. In this study, the detection of *E. canis* DNA in spleen tissue and in blood samples was equally efficient, even in dogs that presented morulae in the splenic aspirate but not in the buffy coat smears. Nevertheless, in some of the dogs where *E. canis* DNA was isolated in spleen tissue and blood samples, no morulae were found. Further studies are needed to explain this controversy. The exact mechanism of *E. canis* and some rickettsiae to stay sheltered in the host is not completely understood (FOLEY et al., 1999). Some studies suggest that the spleen produces inflammatory mediators and other splenic substances that could collaborate in the bacterial persistence in the host and therefore they could act in the disease pathogenesis (HARRUS et al., 1998b). In addition, it was demonstrated that splenectomized dogs experimentally infected with *E. canis* suffer from a milder form of the disease, when compared to non-splenectomized dogs (HARRUS et al., 1998b). On the other hand, splenectomized human beings had a more severe type of ehrlichiosis than non-splenectomized patients (RABINSTEIN et al., 2000), probably because there is a reduction in the antigen's clearance, a decrease in responses against new antigens, and a decrease in phagocytosis and in other protective responses (DAVIDSON; WALL, 2001). These different spleen functions against pathogenic agents could be due to multiple differences in the structure and function of the organ in the species, as suggested by Bowdler (2002).

In one dog, *E. canis* DNA was found in spleen tissue but not in blood samples. In two others, the opposite was true: DNA was isolated from blood but not from spleen samples. The dog with spleen nPCR positive was pancitopenic and had titers of antibodies anti-*E. canis* of 1:1280 and anti-*B. canis* of 1:160. The morulae search of this animal was not done. The other blood-positive and spleen-negative dogs didn't have anti-*E. canis* and anti-*B. canis* antibodies, were anemic and thrombocytopenic, and had leukocytosis (data not shown). One of them had *E. canis* morulae in spleen cytology. These anomalies may suggest that the nPCR could have false negatives results caused by low parasitemia, as reported by Harrus et al. (2004). Although the nPCR is a highly sensitive and specific assay, it should not be the only diagnostic test (WEN et al., 1997). The low frequency of positive dogs in only one of the tests reinforces the idea that the definitive diagnosis of canine ehrlichiosis should be one by more than one test, combined with clinical signs and symptoms presented by the animal.

The present study showed that among the assays for the diagnosis of naturally occurring canine ehrlichiosis, the search for morulae in splenic aspirates is more efficient than in buffy coat smears when it is intended to provide parasitological diagnosis of canine ehrlichiosis. It is important to note that the search for morulae in the spleen tissue is a low-cost assay with easy execution, and which can be safely applied by the clinician to the diagnosis of canine ehrlichiosis. Additionally the spleen nPCR is as efficient as the blood nPCR in the molecular diagnosis of naturally occurring canine ehrlichiosis.

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