

UNIVERSIDADE FEDERAL DO PARANÁ

THIAGO GALLO BIZARI

DETECÇÃO MOLECULAR DE HEMOPLASMAS E PATÓGENOS
TRANSMITIDOS POR CARRAPATOS EM OURIÇO-PIGMEU-AFRICANO
(*ATELERIX ALBIVENTRIS*) NO PARANÁ, SUL DO BRASIL.

CURITIBA

2024

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Dissertação apresentada ao Programa de Pós-graduação em Ciências Veterinárias, Setor de Ciências Agrárias, Universidade Federal do Paraná, como requisito à obtenção do título de Mestre em Ciências Veterinárias.

Orientador: Prof. Dr. Rafael F. C. Vieira

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“Não existe saber mais, ou saber menos. Há saberes diferentes.”

-Paulo Freire

RESUMO

O mercado de *pets* não convencionais cresce cada vez mais, e com isso observa-se novos animais selvagens que começam a ser enquadrados nesse setor. Entre eles, há uma popularidade crescente pelo ouriço-pigmeu-africano *Atelerix albiventris* (MAMMALIA: ERINACEIDAE). Estes animais são originários do continente africano, sendo uma espécie exótica no Brasil. Por seu potencial invasor, assim como o atual conhecimento de seu papel como transmissor de doenças, é importante compreender como estes animais se comportam na cadeia epidemiológica de diversas doenças. O objetivo deste trabalho foi caracterizar um perfil hematológico, bioquímico e de microbitoia cutânea e fecal, assim como pesquisar agentes infecciosos por meio de técnicas moleculares. Amostras de sangue, fezes e swab de pele foram coletadas de vinte e cindo (25) ouriços apreendidos no estado do Paraná. Destas, vinte e duas (22) amostras de sangue foram selecionadas e submetidas a análises moleculares por ensaios de PCR convencional para *Theileria/Babesia* spp. (18S rRNA) e *Ehrlichia/Anaplasma* spp e PCR em tempo real (qPCR) para *Mycoplasma* spp. (16S rRNA). Uma parte do sangue coletado para avaliação hematológica apresentou coágulo e fibrina, dificultando a avaliação de alguns parâmetros. Os dados hematológicos e bioquímicos estão sumarizados nas Tabelas 1 e 2, respectivamente. Os exames microbiológicos da pele mostraram crescimento de *Staphylococcus* spp. (25/25) e *Streptococcus* spp. (13/25). Nas fezes, o isolamento de *Candida tropicalis* (15/25), *Mucor* spp. (12/25) e *Proteus vulgaris* (10/25) foi proeminente. A avaliação molecular dos 22 ouriços resultou negativa para os genes pesquisados, indicando que estes animais não estavam infectados pelos patógenos estudados. Uma vez que são animais mantidos sob tratamento humano, é possível que não tenham tido contato prévio com o vetor destas doenças, ou até mesmo que tenham um protocolo sanitário terapêutica/preventivo realizado frequentemente. Buscar compreender a cadeia de transmissão e determinar protocolos para anteceder doenças nesses animais se faz necessário, uma vez que programas de vigilância são necessários para mitigar ações de saúde pública eficientes e evitar grandes danos ao ambiente e à saúde dos humanos e animais.

Palavras-chave: hedgehogs; carapatos; doenças transmitidas por vetores.

ABSTRACT

The market for unconventional pets is growing more and more, and as a result, new wild animals are beginning to be included in this sector. Among them, there is growing popularity for the African pygmy hedgehog *Atelerix albiventris* (MAMMALIA: ERINACEIDAE). These animals originate from the African continent, being an exotic species in Brazil. Due to their invasive potential, as well as the current knowledge of their role as disease transmitters, it is important to understand how these animals behave in the epidemiological chain of various diseases. The objective of this work was to characterize a hematological, biochemical and cutaneous and fecal microbiota profile, as well as research infectious agents using molecular techniques. Blood, fecal, and skin swab samples were collected from twenty-five (25) hedgehogs seized in the state of Paraná. Twenty-two (22) blood samples were selected and subjected to molecular analysis by conventional PCR assays for *Theileria/Babesia* spp. (18S rRNA) and *Ehrlichia/Anaplasma* spp and real-time PCR (qPCR) for *Mycoplasma* spp. (16S rRNA). (16S rRNA). Some of the blood collected for hematological evaluation showed clots and fibrin, making it difficult to evaluate some parameters. Hematological and biochemical data are summarized in Tables 1 and 2, consecutively. Microbiological examinations of the skin showed growth of *Staphylococcus* spp. (25/25) and *Streptococcus* spp. (25/13). In the feces, the isolation of *Candida tropicalis* (15/25), *Mucor* spp. (25/12) and *Proteus vulgaris* (25/10) were prominent. The molecular evaluation of the 22 hedgehogs was negative for the genes studied, indicating that these animals were not infected by the pathogens studied. Since they are animals kept under human treatment, it is possible that they have not had previous contact with the vector of these diseases, or even that they have a therapeutic/preventive health protocol frequently carried out. Seeking to understand the transmission chain and determine protocols to precede diseases in these animals is necessary, since surveillance programs are necessary to mitigate efficient public health actions and avoid major damage to the environment and the health of humans and animals

.Keywords: hedgehogs; ticks; tick-borne-disease.

LISTA DE ABREVIATURAS

CHGM - Average globular hemoglobin
DNA – Deoxyribonucleic acid
EDTA – Ethylenediaminetetraacetic acid
Gapdh - Glyceraldehyde 3-fosfate dehydrogenase gene
Hb - Hemoglobin's concentration
Hm - Hematimetry
IAT – Instituto Água e Terra
IBGE – Instituto Brasileiro de Geografia e Estatística
IQR – Interquartile Range
MCHC – Mean Corpuscular Haemoglobin Concentration
MCV – Mean Corpuscular Haemoglobin
NaCl – Cloreto de sódio
PCR – Polymerase Chain Reaction
PCV – Packed cell volume
PIB – Produto Interno Bruto
PPT – Proteína plasmática total
RNA – Ribonucleic acid
 s – Desvio padrão
VGM - Average globular volume
 \bar{x} - Média

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1. INTRODUÇÃO

A interação humano-animal vem aumentando cada vez mais, fazendo com que haja um contato próximo entre eles e permitindo que patógenos circulem interespécies. A identificação de zoonoses e o entendimento da epidemiologia destes agentes é de suma importância para criar políticas de controle e prevenção destas doenças (Natterson-Horowitz et al., 2022).

Dentre as zoonoses, as doenças transmitidas por vetores possuem grande relevância em decorrência da ampla disposição geográfica de seus vetores e fácil disseminação, uma vez que havendo os meios adequados, o ciclo se completa. Agentes como bactérias, vírus e protozoários se enquadram nestas categorias e possuem grande relevância na saúde pública (Madison-Antenucci et al., 2020). Muitas vezes estas doenças são negligenciadas e subdiagnosticadas, enquanto isso seus números aumentam e ações efetivas de controle não são colocados em prática.

Além disso, o crescente mercado de pet exóticos coloca em foco a discussão sobre as doenças que podem ser compartilhadas entre os animais e os humanos. Os ouriços-pigmeus-africanos (*Atelerix albiventris*), conhecidos como hedgehogs, tem popularidade crescente no Brasil, apesar de seu comércio ainda não ser autorizado pelos órgãos legais.

A participação destes animais na epidemiologia de doenças é ainda pouco conhecida, havendo poucos dados no mundo com a detecção de agentes infecciosos e as alterações clínicas causadas nestes animais. (Riley & Chomel, 2005).

2. REVISÃO BIBLIOGRÁFICA

2.1 Saúde pública

Zoonoses são doenças que afetam os seres humanos e são transmitidas por animais, relacionadas muitas vezes com classes sociais baixas e áreas menos urbanizadas. Muitas destas doenças acabam não sendo diagnosticadas e favorecem a subnotificação (Natterson-Horowitz et al., 2022).

A expansão das áreas urbanas para onde antes havia mata nativa possibilita maior proximidade entre animais selvagens e os humanos, o que além de aumentar a transmissão de doenças entre ambos, favorece interações que muitas vezes podem causar o óbito dos animais (Richini-Pereira et al., 2010).

Os agentes transmitidos por vetores possuem grande relevância quando falamos de zoonoses. Encontrando o vetor no ambiente, as chances de disseminação da doença são muito grandes (Dantas-Torres et al., 2012). Doenças como Leishmaniose, Febre amarela e Doença de Lyme ganharam destaque nos últimos anos e afetam milhares de indivíduos anualmente (Gorem et al, 2023).

Entender como esses agentes se comportam, quais os possíveis hospedeiros e o papel na transmissão entre humanos são essenciais para que políticas públicas sejam criadas e colocadas em prática, visando diminuir os casos e controlar possíveis endemias. Para tanto, é necessário que a devida atenção seja dada e diagnósticos corretos sejam realizados (Richini-Pereira et al., 2010).

2.2 Animais silvestres

A presença de agentes zoonóticos em animais silvestres torna-se importante quando vemos a população humana interagindo cada dia mais com áreas devastadas e entrando em contato com a fauna (Bicca-Marques e De Freitas, 2010). Em alguns casos, doenças que não apresentam manifestação nestes animais acabam causando sinais clínicos nos animais domésticos e em humanos.

Entender como estas espécies coexistem nos ajuda a prever aumentos de casos e muitas vezes servem como sentinelas de doenças emergentes. Macacos, por exemplo, são grandes indicadores do aumento de casos de febre amarela e ajudam a nortear políticas para evitar que os casos em humanos não cresçam exponencialmente (Bicca-Marques e De Freitas, 2010).

Quando pensamos nos animais silvestres exóticos que se tornam pets, temos um problema ainda maior levando em consideração que esses animais muitas vezes são espécies com potencial para se tornar invasoras e quebrar o equilíbrio do nosso ecossistema (Lockwood et al, 2019). Ainda, a falta de acompanhamento médico-veterinário e o fato de muitas vezes serem espécies ilegais dificulta o controle sanitários sobre esses animais, dificultando medidas de controle e prevenção da disseminação de diversas doenças.

Esses pets exóticos acabam se tornando grande fonte de dispersão de zoonoses conhecidas e em alguns casos, permitem que novos agentes entrem em contato e se tornem problemas de saúde pública (Chomel et al, 2007), aumentando

não apenas os riscos, mas também favorecendo o desenvolvimento de doenças novas e mais adaptadas.

2.3 Ouriços-pigmeus-africanos (*Atelerix albiventris*)

Os ouriços pigmeus africanos (*A. albiventris*), também conhecidos como hedgehogs, são animais encontrados normalmente no continente africano, sendo disseminado pelo mundo como um pet não convencional. É o menor entre os animais do gênero *Atelerix*, tendo comportamento onívoro e noturno predominantemente. (Santana et al, 2010)

A relação destes animais com diversos patógenos que acometem outros animais e são transmitidas por vetores está cada vez mais em pauta. A identificação destes micro-organismos, associando eles a doenças clínicas e buscar correlacionar com a epidemiologia destas doenças se faz necessária para compreender o papel dos hedgehogs na transmissão de zoonoses (Riley & Chomel, 2005).

2.4 Patógenos

Os hemoparasitos da ordem Piroplasmida (*Babesia*, *Cytauxzoon* e *Theileria*) possuem relevância epidemiológica. São organismos intracelulares obrigatórios que necessitam de um hospedeiro vertebrado e invertebrado para concluir seu ciclo de vida. Estes agentes muitas vezes causam sinais clínicos em animais domésticos, mas costumam ser assintomáticos em animais selvagens. (Schnittger et al., 2022).

Os protozoários do gênero *Babesia* comumente são encontrados nos sangues de mamíferos. São transmitidos por várias espécies de carrapatos da família Ixodidae, nos quais também realizam parte do ciclo biológico (Scott e Scott, 2018). Estes agentes infectam os eritrócitos dos hospedeiros definitivos, levando em alguns casos há anemias secundárias a captura destas hemácias.

Os organismos do gênero *Cytauxzoon* acometem mais comumente felinos selvagens e costumam causar doenças mais graves nos felinos domésticos. Novas espécies são frequentemente identificadas em todo o mundo (Gallusová et al., 2016). Com parasitismo em macrófagos, cada vez mais são reportados casos de infecção vinda da fauna silvestre (Squerre et al., 2020).

O gênero *Theileria* se distingue por infectar inicialmente leucócitos e posteriormente eritrócitos. Assim como os anteriores, são transmitidos por carrapatos

e acometem diferentes espécies animais. Com a evolução do diagnóstico molecular, aumentou-se o número de casos, uma vez que na microscopia muitas vezes os organismos eram confundidos entre os diferentes gêneros (Mans et al., 2015).

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4. Objetivos

4.1 Objetivo Geral

O presente estudo teve por propósito pesquisar a infecção de *Atelerix albiventris*, uma espécie exótica no Brasil, por agentes zoonóticos comumente encontrada em nosso território e determinar um perfil sanitário.

4.2 Objetivos Específicos

- a) Pesquisar patógenos transmitidos por carrapatos e hemoplasmas em *A. albiventris*;
- b) Caracterizar o perfil hematológico e bioquímico dos *A. albiventris*
- c) Caracterizar a microbiota cutânea e fecal dos *A. albiventris*

5. Artigo: Health profile of captive Four-toed hedgehogs (*Atelerix albiventris*) from Paraná state, southern Brazil

Abstract

Four-toed hedgehogs are omnivorous mammals native to sub-Saharan Africa and popular as pets worldwide. In Brazil, while their trade is illegal, confiscations remain frequent. This study aimed to assess the health status of 25 hedgehogs from illegal breeding sites, including hematological and biochemical analysis, as well as fecal and skin pathogen screenings, and to investigate the occurrence of haemotropic *Mycoplasma* spp. and tick-borne pathogens in these animals. Real-time PCR (qPCR) was used to detect the 16S rRNA gene of hemoplasmas, while conventional PCR targeting 18S rRNA and 16S rRNA genes were used to identify *Theileria/Babesia* spp. and *Ehrlichia/Anaplasma* spp., respectively. No ectoparasites were found in these animals, and the hematological and biochemical evaluation indicated a good general state of health. Microbiological examinations of the skin showed growth of *Staphylococcus* spp. (25/25) and *Streptococcus* spp. (13/25). In the feces, the isolation of *Candida tropicalis* (15/25), *Mucor* spp. (12/25) and *Proteus vulgaris* (10/25) were prominent. All samples tested negative for *Mycoplasma* spp., *Theileria/Babesia* spp., and *Ehrlichia/Anaplasma* spp. The data obtained suggest that the animals evaluated were not infected by harmful pathogens. Understanding their role in zoonotic transmission is necessary, especially when about exotic species.

Keywords: *Ehrlichia* spp.; *Anaplasma* spp.; *Mycoplasma* spp.; Hedgehog.

Introduction

Four-toed hedgehogs (*Atelerix albiventris*, Mammalia: Erinaceidae) are nocturnal, omnivorous mammals widely distributed across sub-Saharan Africa (Harrison & Bates 1991; Kock & Ebenau 1996; Hutterer 2005) and are considered an exotic species in Brazil. European and African hedgehogs are increasingly sought as exotic companion animals (Ruszkiowski et al., 2021), although keeping them without special permits is illegal in several countries. Despite the prohibition on the commercialization and ownership of hedgehogs in Brazil, confiscation reports are

frequent, underscoring the need for policies to prevent their uncontrolled breeding and spread.

Hedgehogs are commonly infested by a diversity of ectoparasites such as hard ticks (Acari: Ixodidae) and fleas (Souza et al., 2006) which are regarded as vectors of zoonotic pathogens, such as *Rickettsia* sp., *Borrelia* sp., *Ehrlichia* sp., *Anaplasma* sp. and *Bartonella* sp. species (Bezerra-Santos et al., 2021), *Leishmania* sp. (Pourmohammadi and Mohammadi-Azni, 2019) and tick-borne encephalitis viruses. Considering the role of other animals of the Erinaceidae family in disseminating these agents, there is a potential risk for infection in these species as well. Investigating hedgehog pathogens is essential to identify clinicopathological changes (Balti et al., 2021), and detecting the gastrointestinal parasites helps clarify these animals' sanitary and epidemiological profiles. Furthermore, the close relationship between humans and animals raises concerns about the potential of zoonotic diseases. Also, the increasing demand for unconventional pets for companionship presents a challenge to public health, especially regarding diseases that are often overlooked (Riley & Chomel, 2005). Therefore, this study aimed to investigate the presence of haemotropic Mycoplasma and tick-borne pathogens (TBP), such as *Ehrlichia/Anaplasma* and *Babesia/Theileria*, while also evaluating the hematological and biochemical profiles of four-toed hedgehogs apprehended by the Água e Terra Institute (IAT), in the state of Paraná, southern Brazil.

Material and Methods

Sampling

A total of 25 four-toed hedgehogs from a governmental apprehension by request of the Água e Terra Institute (IAT) in the state of Paraná, southern Brazil, were selected for the study while kept under the responsibility of the State agency.

Blood samples

After chemical restrained, with an association of ketamine (5 mg/kg) and xylazine (1 mg/kg), the animals were laid on a flat surface, and blood samples (up to 3 mL) were collected by venipuncture of the proximal jugular or cranial vein and placed into sterile tubes containing EDTA (BD Vacutainer®, Franklin Lakes, NJ, EUA) and

stored at –20 °C prior the molecular analysis. Three milliliters were stored in tubes containing a serum separator gel (BD Vacutainer®) and kept at room temperature (25 °C) until a visible clot appeared. The samples were centrifuged at 1500 × g for 5 min, serum separated and stored at – 20 °C for biochemical analysis.

Complete blood count and biochemical analysis

The differential leukocyte count was performed by counting 100 leukocytes on a blood smear. Packed cell volume (PCV) was determined by the microhematocrit method, using capillary tubes and centrifugation at 11,000 RPM for 5 minutes (Farrand, 1976). Hemoglobin (HGB) concentration was measured using the spectrophotometric cyanmethemoglobin method. Total red blood cell (RBC), white blood cell (WBC), platelet counts, and hemoglobin (Hgb) concentrations were measured using a hematology analyzer (BC-2800 Vet, Myndray®). The hematimetric indices, including mean corpuscular volume (MCV) and mean corpuscular hemoglobin concentration (MCHC), were calculated using Wintrobe's methodology (1990).

Serum samples were thawed at room temperature before analysis on an automated biochemical analyzer (BS-200, Myndray®), previously calibrated using a commercial control serum (Control Lab®). The following parameters were evaluated using the respective methods: total protein concentration (biuret method, Katal®), albumin (bromocresol green method, Katal®), total globulins (calculated by the difference between the total proteins and albumin), urea (UV enzymatic method of urease-GLDH, Kovalent®), creatinine (Jaffé method, Kovalent®), aspartate aminotransferase (AST; UV kinetic method of AST, Katal®), alanine aminotransferase (ALT; UV kinetic method), and alkaline phosphatase (UV kinetic method).

DNA extraction and PCR assays

The genomic DNA from 200µL of whole blood was extracted using a commercially available kit (ReliaPrep™ gDNA blood Miniprep System, Promega, Madison, Wisconsin, USA), following the manufacturer's instructions. Ultrapure water was used as a negative control in parallel to monitor cross-contamination. A conventional PCR assay (cPCR) for the mammal endogenous gene glyceraldehyde-3-fosphate dehydrogenase (*gapdh*) was performed to ensure successful DNA extraction (Rebouças et al., 2013). Thereafter, the hedgehog's DNA samples were screened

using a universal SYBR green real-time PCR (qPCR) assay targeting the 16S rRNA gene of hemoplasmas, as previously described (Willi et al., 2009). A standard curve was calibrated using serial dilutions of gBlock™ (Integrated DNA Technologies, Coralville, IA, USA). All parameters were analyzed according to the standards established by MIQE (Minimum Information for Publication of Quantitative Real-time PCR Experiments) (Bustin et al., 2009). Additionally, DNA samples from hedgehogs were screened using cPCR assays targeting a 551bp fragment of the 18S rRNA gene of *Theileria/Babesia* spp. (Almeida et al., 2012), and a 349bp fragment of the 16S rRNA gene of *Ehrlichia/Anaplasma* spp. (Parola et al., 2000). DNA from *Babesia vogeli* and *Ehrlichia canis* obtained from naturally infected dogs and nuclease-free water were used as positive and negative controls, respectively.

Skin swabbing

The skin samples of 25 animals were individually collected by swabbing the interscapular skin with sterile flexible rods. These swabs were then plated onto a blood agar plate. Colonies that grew were further identified by light microscopy and biochemical tests following the methodology described by Koneman et al (2006).

Feces culture

25 feces samples were collected from the coop, without isolation of each animal, packed in sterile tubes, and sent for microbiological isolation. The samples were inoculated into tubes containing 2mL of BHI broth (Brain-Heart infusion), incubated under shaking at 37°C overnight. Subsequently, aliquots were plated on MacConkey agar plates supplemented with 2 µg/mL of ceftriaxone and colistin. Isolated colonies were seeded again on MacConkey agar plates without antibiotics and incubated at 37°C for 18 hours. The collected colonies were evaluated under optical microscopy and for morphological identification. (Koneman et al., 2006)

Statistical analysis

The Chi-square test was used to determine the average values and deviations from the standard hematological patterns. A 95% confidence interval and p-values were calculated. For samples rejected normality (RN), robust data analysis was

performed, and the interquartile range (IQR) was calculated. Data compilation and analysis were performed using Microsoft Excel™ and the MedCalc® software.

Results

Blood count and biochemical analysis

The mean (\bar{x}) and standard deviation (s) of the blood counting and hematocrit were $5.9:0.99 \times 10^{12}/L$ and $40.7:4.64\%$, respectively. The statistical analysis of the blood counting and biochemistry results are summarized in Tables 1 and 2, respectively.

After evaluation, some samples were excluded from the white blood series counting and differential due to the presence of clogs. Additionally, some samples had insufficient volume to repeat or confirm some biochemical tests and were excluded from the results.

PCR results

The *gapdh* gene was consistently amplified in all hedgehog samples. All samples tested negative for hemoplasmas, *Ehrlichia/Anaplasma* spp., and *Theileria/Babesia* spp.

Skin swabbing

Nine pathogens were detected in the animals, with some cases exhibiting co-infections. *Candida tropicalis* (15/25, 60%; 95% CI), *Mucor* spp. (12/25, 48%; 95% CI), *Proteus vulgaris* (10/25; 40%; 95% CI) and *Enterobacter sakazakii* (7/25; 28%; 95% CI) were the most frequent pathogens, followed by *Proteus mirabilis* (4/25; 16%; 95% CI), *Citrobacter amalonaticus* (2/25; 8%; 95% CI), *Enterobacter* spp. (2/25; 8%; 95% CI), *Geotrichum* spp. (1/25, 4%; 95% CI), and *Aspergillus* spp. (1/25, 4%; 95% CI).

Feces culture

From the 30 fecal samples, specimens of *Bacillus* spp. (4/25; 3,3%; 95% CI), *Micrococcus* spp. (2/25; 8%; 95% CI); *Staphylococcus* spp. (25/25; 100%; 95% CI), and *Streptococcus* spp. (13/25; 52%; 95% CI) were identified.

Discussion

In microbiological examinations of the skin, the detection of *Candida* spp. in 60% of patients is relevant when considering the zoonotic nature of this pathogen, more effectively in the treatment of direct handlers, such as zookeepers and veterinarians, and in the case of contact with immunosuppressed individuals. Furthermore, the other pathogens identified may have been found in skin diseases or even systemic diseases as secondary opportunistic agents, since they appear to be present on the skin.

In an analysis of vaginal swabs from some sea urchins in Indonesia, specimens of *Proteus* spp. and *E. coli* were isolated (Ash-Santri et al, 2021). When compared with our data from skin swabs, there is a greater variety of agents present in the skin than in the genital tract.

The fecal analysis yielded a different result from another study in Indonesia (Amanbayeva, 2021). This could be associated with the difference in location and the consequent change in the microbiota. *Salmonella* spp. has been described in *A. albiventris* (Perez et al., 2021), causing concern about the role of these animals in its transmission. In the present study, this bacterium was not observed, which may again be related to the environment in which the evaluated animals were found.

On the other hand, the observation of *Staphylococcus* spp. in all samples, as well as *Streptococcus* spp. in 52% of the samples, shows an incidence of these agents in the digestive tract of these animals, which may also act as pathogens in cases of dysbiosis or even affect contacts of their species or of different species that are immunocompromised.

The hematological values of four-toe hedgehogs analyzed in this study showed greater variations compared to previous studies (Okorie-Kanu et al., 2014). Although all animals were clinically healthy, some cases of leukocytosis were observed, suggesting an active inflammatory process. The difficulty in determining the exact age of the animals and the limited literature on the hematological patterns of these species complicated the accurate interpretation of the data obtained. Additionally, the serum biochemical analyses showed discrepancies when compared to studies on this species or similar ones (Okorie-Kanu et al., 2015; Gabriele Rossi et al., 2014). For example, compared to other domestic animals, elevated AST values in one of the sampled hedgehogs may indicate significant liver damage or muscle

damage, especially related to increased CK activity. Similarly, the mild hypoalbuminemia observed in this study also suggests the presence of an inflammatory process since albumin is a negative acute-phase protein. During inflammation, the synthesis of specific globulins (acute-phase proteins) increases by hepatocytes, while B lymphocytes synthesize immunoglobulins, leading to a relative decrease in albumin levels (Stocham & Scott, 2011).

Hedgehogs have been suggested as potential hosts or reservoirs of zoonotic vector-borne pathogens (Krawczyk et al., 2015) and are frequently infested with ticks and fleas (Iacob & Iftinca, 2018). The ticks commonly found on hedgehogs belong to the genera *Amblyomma*, *Rhipicephalus*, *Dermacentor*, *Hyalomma*, and *Ornithodoros* (Estrada-Peña et al., 2018; Guglielmone et al., 2014), all of which are known vectors of zoonotic pathogens such as *Rickettsia* spp., *Borrelia* spp., *Ehrlichia* spp., *Anaplasma* spp., and *Bartonella* spp. (Regnery et al., 1992; Bouyer et al., 2001; Millán et al., 2019). These pathogens are of significant medical and veterinary importance in neotropical regions (Barros-Battesti et al., 2006).

Wild animals, mainly those presenting synanthropic behavior, are considered important reservoirs of zoonotic pathogens (Hassell, Begon, Ward, & Fèvre, 2017). In Brazil, there are several studies identifying tick-borne diseases in synanthropic animals, some with great zoonotic importance (Collere et al., 2022; Orozco et al., 2022; Vieira et al., 2009). In a previous study, *A. phagocytophilum* and *Borrelia* spp. were detected in ticks collected from European hedgehogs (*Erinaceus europaeus*) (Krawczyk et al., 2015). Despite the detection of various zoonotic pathogens (bacterial, viral, protozoal, and fungal) in hedgehogs in previous studies (Riley & Chomel, 2005; Dumitrache et al., 2013; Kocan et al., 2010; Krawczyk et al., 2015; Pourmohammadi & Mohammadi-Azni, 2019), all animals in this study tested negative for hemoplasmas and tick-borne pathogens.

Hedgehogs, which thrive in urban, rural, and natural environments alongside domestic animals and humans (Skuballa et al., 2007), can contribute to environmental disturbance when introduced into new ecosystems. Since the hedgehogs were seized animals from captive breeding environment, they may not have been exposed to arthropods that could have infected them. In general, animals bred for trade are often subjected to veterinary care and medications, which may help

maintain a controlled health status. This could explain the absence of pathogens and ectoparasites in the animals studied.

5.0 Conclusion

The present study collected hematological and biochemical data that indicated a certain degree of active inflammation, despite the absence of clinical signs and perceptible clinical disease. It is likely that these animals have greater resistance to small physiological changes. Furthermore, the results of fecal and skin cultures show a diverse microbiota, with possible opportunistic pathogens for the individual or contacts. Vector-borne diseases were not identified. It is necessary to expand scientific knowledge about the hematological and biochemical profiles, as well as the diseases that affect *A. albiventris*, to better understand its unique health and the role of zoonotic diseases in this species.

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Conflict of interest

The authors declare no conflicts of interest.

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Table 1 – Haematological profile of Captive *Atelerix albiventris* from the state of Paraná, south Brazil

Analyte	UNIT	N	Mean	SD	Median	Min	Max
RBC	$10^{12}/L$	25	5.9	0.99	5.68	4.3	7.43
PCV	%	24	40.70	4.64	40.5	33	53
Hemoglobin	g/L	24	120.17	18.68	118.5	92	161
WBC	$10^6/L$	17	12.3	5.1	12	6	21.8
Neutrophils	%	17	52.29	12.5	53	32	72
Bands	%	17	0.29	0.58	0	0	2
Lymphocytes	%	17	28.9	9.91	30	16	47
Monocyte	%	17	6.35	4.13	5	1	18
Eosinophils	%	17	9.94	5.91	9	0	19
Basophils	%	17	1.70	1.72	1	0	6
Neutrophils	$10^6/L$	17	6.58	3.37	5.73	1.92	13.46
Bands	$10^6/L$	17	0.043	0.1	0	0	0.38
Lymphocytes	$10^6/L$	17	3.47	1.8	3.36	1.08	8.93
Monocyte	$10^6/L$	17	0.78	0.6	0.67	0.12	2.25
Eosinophils	$10^6/L$	17	1.16	0.84	0.93	0	3.4
Basophils	$10^6/L$	17	0.21	0.22	0.18	0	0.88

Table 2 – Serum biochemistry profile of captive *Atelerix albiventris* from the state of Paraná, south Brazil -

Analyte	UNIT	N	Mean	SD	Median	Min	Max
Urea	mmol/L	22	4.20	0.95	4.04	2.44	6.79
Creatinine	mmol/L	22	40.99	8.43	44.21	26.52	53.05
ALP	U/L	22	64.97	27.41	57.9	10.1	133.9
ALT	U/L	21	67.22	23.89	60.6	32.9	120.7
AST	U/L	21	29.76	14.22	28.6	14.3	80.4
Total protein	g/L	23	69	13.54	69	47	113
Albumin	g/L	23	26.34	5.17	27	10	25
Globulin	g/L	23	43.04	17.46	41	20	112
A:G RATIO		23	0.69	0.25	0.64	0.08	1.35

6. CONSIDERAÇÕES FINAIS

A pesquisa por agentes infecciosos com potencial zoonótico em espécies exóticas que estão sendo inseridas em nosso país é fundamental para evitar grandes danos e prever o comportamento epidemiológico destas doenças nesses animais. Apesar deste estudo não apresentar positividade para os patógenos selecionados, a vigilância passiva deve ser mantida, uma vez que *A. albiventris* é sabidamente uma possível fonte de infecção de diversos patógenos.