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Article

Ehrlichia canis and *Rickettsia conorii* Infections in Shelter Dogs: Seropositivity and Implications for Public Health

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Abstract: A cross-sectional study was conducted to gain insight into the epidemiology of canine ehrlichiosis and rickettsiosis in northern Portugal. Specific IgG antibodies to *Ehrlichia canis* were analysed using a commercial indirect ELISA test, and antibodies to *Rickettsia conorii* using a commercial IFAT. A total of 113 dogs from two different shelters were sampled, and seroprevalence values of 0.9% (95% confidence [CI]: 0.2–4.8%) for *E. canis* and 9.7 (95% CI: 5.5–16.6%) for *R. conorii* were found. Univariable models investigated risk factors for seropositivity. The odds ratios (OR) of *R. conorii* seropositivity were higher for female dogs (OR = 2.32; 95% CI: 1.58–3.39) and for dogs belonging to Shelter 2 (OR = 3.0; 95% CI: 2.28–3.95). Seropositive dogs for co-infection (*E. canis* + *R. conorii*) were more frequently observed in females (OR = 2.4; CI 95%: 1.66–3.47) and in Shelter 2 (OR = 2.72; 95% CI: 1.97–3.76). These findings show that shelter dogs in northern Portugal are exposed to *E. canis* and *R. conorii*, which can affect both canines and humans. It is imperative to adopt a One Health approach to educate the public about the hazards of canine zoonoses and develop legislation and procedures to control their spread and preserve public health.

Keywords: Antibodies; dog; *Ehrlichia canis*; Portugal; *Rickettsia conorii*; shelter.

1. Introduction

In the past few years, there has been an increasing recognition of the significance of canine vector-borne diseases (CVBD) as a growing threat to the health of both humans and animals across the globe. Factors such as climate change, globalization, rising international mobility and trade, and the rapid growth of human and canine populations have all played a part in causing a shift in the distribution of CVBD [1,2]. The aetiology of CVBD are multifaceted and could encompass a range of disease-causing agents, such as viruses, bacteria, protozoa, and helminths. These harmful pathogens are typically transmitted through vectors like fleas, ticks, mosquitoes, and lice. [1,2]. *Ehrlichia canis*, member of the order Rickettsiales and an obligate intracellular Gram-negative bacterium, is the primary causative agent of canine monocytotropic ehrlichiosis (CME), a severe and sometimes fatal immunosuppressive disease in temperate and tropical regions of Africa, Europe, and United States of America (USA). It is transmitted globally by the brown dog tick, *Rhipicephalus sanguineus sensu lato* [3–5]. In this vector, *E. canis* transmission is feasible transstadially but not transovarially [6]. In Europe, only the species *E. canis* has been identified in dogs [4].

Ehrlichia canis causes a wide range of clinical signs in dogs, with infection ranging from subclinical to fatal illness [7]. Common clinical manifestations include anorexia, epistaxis, fever, lethargy, weight loss, other hemorrhagic signs, pale mucous membranes, and lymph node enlargement. [8]. Rickettsioses, caused by Gram-negative obligate intracellular bacteria also in the

order Rickettsiales and transmitted by ticks, represent a relevant causative factor within CVBD [9]. The Spotted Fever Group (SFG) illnesses in Mediterranean, southern Europe, North and Sub-Saharan Africa, the Middle and Far East, America, and Australia are most commonly caused by *Rickettsia conorii*. Mediterranean spotted fever (MSF) is a disease that strikes suddenly and in humans typically causes high fever, flu-like symptoms, a black eschar at the site of the tick bite, and a maculopapular rash. In serious cases, the disease may cause severe neurological symptoms and affect multiple organs. The mortality rate is estimated to be around 2.5%. Elderly age, cirrhosis, chronic alcoholism, and glucose-6-phosphate dehydrogenase deficiency are traditional risk factors for severe forms of the disease [10,11]. This seasonal human disease predominantly occurs from April to October, reaching its peak from June to August [10]. The first cases of human infection by *R. conorii* in Portugal were described in 1910 with a disease characterized by high fever and skin spots [12]. The primary vector of *R. conorii* is the brown dog tick *R. sanguineus* s.l. [13]. This tick species exhibits a global geographic range, a high capacity for pathogen transmission and a remarkable ecological adaptation [2]. However, other species of *Rhipicephalus* and *Ixodes* ticks may also serve as vectors for *R. conorii* [14]. Due to their high tick exposure, dogs serve as a sentinel for human infection. Since dogs live near humans and frequently share the same living space, the presence of seropositive dogs can indicate endemic locations and risk factors for illness occurrence in humans [15].

To confirm a rickettsial infection (including *E. canis* and *R. conorii*), it is necessary to either directly detect the presence of the bacteria through molecular methods or perform serological testing to identify the presence of specific antibodies. However, it is important to note that if the test is done too early in the course of the bacterial infection before the production of antibodies, it may yield false negative results [16–18].

Shelter medicine plays an important role in the health of animals, people and the environment, making it a compelling example of the One Health concept. This integrated approach recognises the interconnectedness of human, animal and environmental health and emphasises how they influence each other [19]. Shelter medicine contributes to human health by preventing the spread of zoonotic diseases. Providing medical care and vaccination to shelter animals reduce the risk of these diseases spreading to shelter staff and potential adopters [20]. Shelter medicine in Portugal extends its impact beyond the country borders through international adoptions. Many shelters and rescue organisations in Portugal facilitate the adoption of animals by individuals and families from other countries [21,22]. This provides homes for animals and promotes global cooperation in animal welfare and health. Shelter animals, especially dogs, are particularly susceptible to tick infestations and the pathogens transmitted by these vectors, including *E. canis* and *R. conorii* [23–25]. These dogs serve as a critical reservoir for these vector-borne agents, potentially contributing to their transmission to other animals and even humans. The confined and often overcrowded conditions within shelter environments can facilitate close contact between infected and susceptible animals, increasing the risk of disease spread [24,26]. Furthermore, as dogs are known to share strong bonds with humans and often become adopted into households, the potential for zoonotic transmission becomes a relevant concern. Thus, shelter dogs play a vital role in the epidemiology of these vector-borne diseases, warranting attention to disease prevention and control measures to safeguard the health of both animals and humans.

Conducting seroepidemiological surveys in shelter animals is essential for the animal's welfare, reducing the risk of disease transmission to humans and maintaining community health. This embodies the One Health concept, recognising the interdependence between human and animal health. This study aimed to conduct a serological survey for *E. canis* and *R. conorii* infections in dogs from two animal shelters in the North of Portugal.

2. Materials and Methods

2.1. Study Area

This study was conducted in two shelters, one in Braga district (Shelter 1) and the other in Bragança district (Shelter 2), located in the North region of Portugal. Braga is located in the former province of Minho, and Bragança is part of the historical province of Trás-os-Montes e Alto Douro.

The geographical area of the North of Portugal spans 21.286 km² and has a resident human population of 3.587.074 inhabitants, according to the 2021 census [27].

2.2. Animals and samples

This study was based on a convenience sample of 113 dogs from two shelters. All these dogs were available for adoption. Based on a physical examination, veterinarians classified the animals as apparently healthy dogs. All dogs were examined for ectoparasite infestation (ticks, fleas and lice).

Dogs were sampled from March to May 2022. Blood samples were collected from dogs of two shelters in the scope of regular testing. The samples were taken aseptically by puncturing the cephalic vein, and 2 ml of blood was collected into anti-coagulant-free vacutainers. Information was recorded on the animals' sex, age, and shelter localisation. Travel histories were not available. After 15 minutes of centrifugation at 1500 g, sera were separated and stored at -20 °C for further analysis. Serological analysis included determining the presence of specific antibodies to *E. canis* by an enzyme-linked immunosorbent assay (ELISA) and *R. conorii* by an indirect immunofluorescence antibody test (IFAT).

For *Ehrlichia* diagnosis, serum samples from dogs were diluted at 1:100 in sample buffer and screened for qualitative detection of circulating IgG antibodies for *E. canis*, with the Euroimmun® test (Euroimmun Medizinische Labordiagnostika AG, Lübeck, Germany). The reported sensitivity/specificity of the Euroimmun® test were 92%/100% for *E. canis*. This ELISA was operated according to the manufacturer's instructions in the product package insert. Ratios were stratified into three rising categories: samples with a ratio < 0.8 were considered negative, samples between ≥ 0.8 and < 1.1 were considered borderline, and samples with a ratio ≥ 1.1 were considered positive.

The same samples were further tested by IFAT using commercial IFA slides (MegaFLUO® RICKETTSIA conorii, MEGACOR Diagnostik GmbH, Hoerbranz, Austria) for the detection of specific IgG antibodies to *R. conorii* according to the manufacturer's instructions. Sera were tested at a cut-off dilution of 1:80, which was considered positive.

This study was approved by the Ethics Committee of UTAD (process reference: Doc6-CE-UTAD-2022).

2.3. Data analysis

An exact binomial test was used to calculate confidence intervals (CI) for the proportions, with a 95% confidence level. Chi-square and Fisher's exact tests compared proportions of positivity related to categorical dependent variables. Analyses were done with SPSS® 29.0 software for Windows. A probability (*p*) value < 0.05 was regarded as statistically significant. Case definition: a dog tested positive for *E. canis* or *R. conorii* antibodies and was considered infected.

Binomial variables showing a significant difference between categories were selected for univariate logistic regression analysis to identify independent risk factors of exposure to the *E. canis* or *R. conorii*, calculating odds ratios (OR) and their 95% CI.

3. Results

A total of 113 dogs were studied (Figure 1). There were no detectable ticks in any dogs at visual inspection. Regarding sex, 45 (39.8%) were females and 68 (60.2%) were males. There were 57 (50.4%) animals aged 12 months or less and 56 (49.6%) older than 12 months. All dogs were mongrels, i.e. they did not belong to any officially recognized breed.

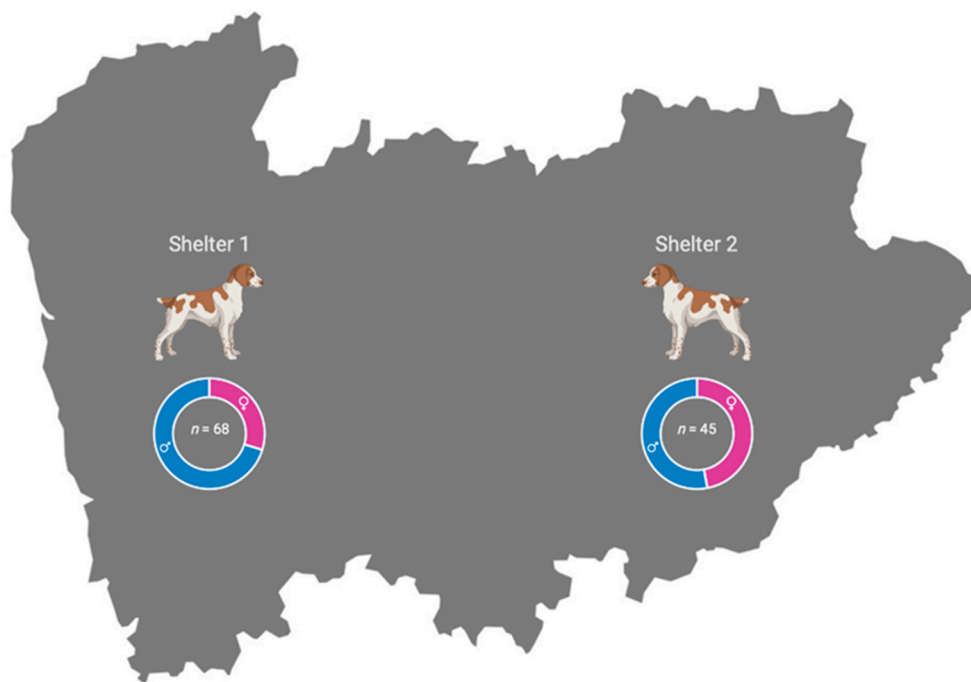


Figure 1. A total of 113 dogs were sampled from two shelters in Braga (Shelter 1) and in Bragança (Shelter 2) districts, in the North of Portugal.

Seroprevalence values of 0.9% ($n = 1$) (95% CI: 0.2–4.8%) for *E. canis* and 9.7 ($n = 11$) (95% CI: 5.5–16.6%) for *R. conorii* were found. Of the dogs tested, 3.5% ($n = 4$) (95% CI: 1.4–8.7%) and 10.6% ($n = 12$) (CI: 6.2–17.6%) had inconclusive results for *E. canis* and *R. conorii*, respectively. Twelve dogs were positive for all agents (10.6%; 95% CI: 5.6–17.8%) (Table 1).

Table 1. Seroprevalence of *Ehrlichia canis* and *Rickettsia conorii* infections in two shelter dog populations prepared to be adopted ($n = 113$ dogs) from the North of Portugal.

Pathogen	No. of infected Dogs	Inconclusive results	Prevalence (%)	95% CI *
<i>Ehrlichia canis</i>	1	4	0.9	0.02–4.8
<i>Rickettsia conorii</i>	11	12	9.7	4.9–16.8
Co-infection	12	16	10.6	5.6–17.8

*95% CI (confidence interval).

3.1. Seropositivity to *Ehrlichia canis*

Among the *E. canis* positive samples, the prevalence in females (2.2%; 95% CI: 0.06–11.8%) was higher than in males (0.0%; 95% CI: 0.0–5.3%), but the difference was not statistically significant ($p = 0.173$). Regarding age, the prevalence found in dogs with 12 months or less was 0.0% (95% CI: 0.0–6.3%), and in older than 12 months was 1.8% (95% CI: 0.04–9.6%), but these differences were not statistically significant ($p = 0.235$). Regarding origin, the lowest value of prevalence was found in Shelter 2 (0.0%; 95% CI: 0.0–7.9%) and the highest in Shelter 1 (1.5%; 95% CI: 0.04–7.9%), with these differences not being statistically significant ($p = 0.312$) (Table 2).

Sex, shelter and age were not found to be significantly associated with seroprevalence of *Ehrlichia canis*.

3.2. Seropositivity to *Rickettsia conorii*

The seroprevalence of antibodies to *R. conorii* was significantly different between females (20.0%; 95% CI: 9.6–34.6%) and males (2.9%; 95% CI: 0.36–10.2%) ($p = 0.003$). When comparing results among different age groups, the prevalence found in dogs with 12 months or less was 5.3% (95% CI: 1.1–14.6%), and 14.3% in dogs older than 12 months (95% CI: 6.4–26.2%), but these differences were not

statistically significant ($p = 0.106$). Regarding origin, the lowest value of prevalence was found in the Shelter 1 (0.0%; 95% CI: 0.0–5.3%) and the highest in Shelter 2 (24.4%; 95% CI: 12.9–39.5%) (Table 2), with these differences being statistically significant ($p < 0.001$). Age was not significantly associated with the seroprevalence of *R. conorii*.

3.3. Seropositivity to co-infections (*E. canis* + *R. conorii*)

The seroprevalence of antibodies to co-infections (*E. canis* + *R. conorii*) was significantly different between females (22.2%; 95% CI: 11.2–37.1%) and males (2.9%; 95% CI: 0.36–10.2%) ($p = 0.001$). When comparing results among different age groups, the prevalence found in dogs with 12 months or less was 5.3% (95% CI: 1.1–14.6%), and 16.1% in dogs older than 12 months (95% CI: 7.6–28.3%), but these differences were not statistically significant ($p = 0.057$). Regarding origin, the lowest value of prevalence was found in Shelter 1 (1.5%; 95% CI: 0.04–7.9%) and the highest in Shelter 2 (24.4%; 95% CI: 12.9–39.5%) (Table 2), with these differences being statistically significant ($p < 0.000$). Age was not significantly associated with seroprevalence of co-infection (*E. canis* + *R. conorii*).

Table 2. Seroprevalence of infection by *Ehrlichia canis*, *Rickettsia conorii* and co-infection in shelter dogs from the North of Portugal.

Variables/categories	<i>E. canis</i> , positive/ total (%)		<i>R. conorii</i> , positive/total (%)		Coinfection (<i>E.</i> <i>canis</i> + <i>R.</i> <i>conorii</i>), positive/total (%)	
		95% CI		95% CI		95% CI
Sex	$p = 0.173$		$p = 0.003^*$		$p = 0.001^*$	
Male	0/68 (0.0%)	0.0–5.3%	2/68 (2.9%)	0.36–10.2%	2/68 (2.9%)	0.36–10.2%
Female	1/45 (2.2%)	0.06–11.8%	9/45 (20.0%)	9.6–34.6%	10/45 (22.2%)	11.2–37.1%
Age	$p = 0.235$		$p = 0.106$		$p = 0.057$	
≤ 12 months	0/57 (0.0%)	0.0–6.3%	3/57 (5.3%)	1.1–14.6%	3/57 (5.3%)	1.1–14.6%
> 12 months	1/56 (1.8%)	0.04–9.6%	8/56 (14.3%)	6.4–26.2%	9/56 (16.1%)	7.6–28.3%
Origin	$p = 0.312$		$p \leq 0.000^*$		$p \leq 0.000^*$	
Shelter 1	1/68 (1.5%)	0.04–7.9%	0/68 (0.0%)	0.0–5.3%	1/68 (1.5%)	(0.04–7.9%)
Shelter 2	0/45 (0.0%)	0.0–7.9%	11/45 (24.4%)	12.9–39.5%	11/45 (24.4%)	12.9–39.5%

* $p < 0.05$.

3.4. Risk factors for *Rickettsia conorii* and the co-infection (*E. canis* + *R. conorii*) in sheltered dogs

Univariable models results in shelter dogs are shown in Table 3. Two variables were associated ($p < 0.05$) with seropositivity to *R. conorii* in shelter dogs in the univariable model. At the individual level, the odds of *R. conorii* seropositivity were found to be higher for female dogs (OR = 2.32; 95% CI: 1.58–3.39) and for dogs belonging to Shelter 2 (OR = 3.0; 95% CI: 2.28–3.95).

Seropositive dogs for co-infection (*E. canis* + *R. conorii*) were more frequently observed in females (OR = 2.4; CI 95%: 1.66–3.47) and in Shelter 2 (OR = 2.72; (95% CI: 1.97–3.76).

Table 3. Risk factors for *Rickettsia conorii* and the co-infection (*E. canis* + *R. conorii*) in sheltered dogs.

Dependent variable/ /Risk factor	<i>p</i> value	OR	95% CI
Positivity to <i>R. conorii</i>			
Sex	$p = 0.003$		
Male		1	

Female		2.32	1.58–3.39
Origin	$p \leq 0.000$		
Shelter 1		1	
Shelter 2		3.0	2.28–3.95
Positivity to mixed infection			
Sex	$p = 0.001$		
Male		1	
Female		2.4	1.66–3.47
Origin	$p \leq 0.000$		
Shelter 1		1	
Shelter 2		2.72	1.97–3.76

4. Discussion

Canine vector-borne diseases (CVBD) result from various pathogens transmitted to dogs by arthropods, such as ticks and insects. Numerous CVBD agents have zoonotic implications, as dogs can act as carriers/reservoirs and early indicators of infections in humans. The study aimed to evaluate the prevalence of seropositive infection or exposure to *E. canis* and *R. conorii* in the dog population of two shelters in the North of Portugal.

The present results revealed the presence of antibodies to *E. canis* and *R. conorii* in dogs in the study area. The seroprevalence of *E. canis* in our study was 0.9%. Another study carried out in dogs in 120 veterinary medical centres from all the regions of mainland and insular Portugal, using a rapid test ELISA, reported an apparent seroprevalence of 0.7% in the North of Portugal. This same study found an overall seroprevalence of 4.1% in apparently healthy dogs and 16.4% in dogs suspected of CVBD [28]. In the North of Portugal, molecular techniques previously confirmed the infection in a dog with clinical signs [29,30]. Other epidemiological studies performed by molecular techniques confirmed the infection in Central [31] and southern Portugal [32,33]. In Portugal, there is also a report of *E. canis* infection in foxes [34]. The individual seroprevalence found in dogs in this study was similar to previously reported values in the North of Portugal, but much lower than other studies in the country. Our findings are higher than the lower seroprevalence in some European countries, ranging from 0.2% in Hungary [35] to 0.3% in Finland and France [35,36]. Our findings are apparently lower than the reported seroprevalences among dogs in Europe. Previous studies in Spain reported seroprevalence values of 3.1–19.2% in Spain [37], 6.4–46.7% in Italy [38], 58.3% in Greece [39], 20.7% in Turkey [40], 17.9% in Albania [41], 2.1% in Romania [42], 0.9–10.1% in Germany [43], 11.1% in Serbia [44] and 29.8% in public shelter and 12.3% owned dogs, in Montenegro [45].

Other studies outside Europe performed with molecular techniques found values ranging from 3.1% in Qatar [46] to 1.9%–5.8% in Angola [47,48].

In the present study, we found an overall seroprevalence of 9.7% for *R. conorii*, indicating some history of exposure to or active infection with this pathogen. This value is much lower than previous findings in Spain. Prior studies recorded a prevalence of 56.4% in northeastern Spain [49] and 24.6% in the northwest part of the country [37]. Studies performed in Italy, an endemic country, reported anti-*R. conorii* antibodies in dogs with seroprevalences ranging from 15.5% to 74% [38,50]. A study on police dogs reported a seroprevalence of 72% in Albania [51]. Other seroprevalences in dogs range from 23% in Croatia [52] to 44.8% (in hunting dogs) in Serbia [53]. In the study performed in Montenegro, the prevalence of *R. conorii* was higher in owned dogs (81.9%) than in dogs from a shelter (60.6%) [45].

We additionally studied other risk factors for seropositivity. The OR of being positive to *R. conorii* and of co-infection were significantly higher in females than males. Our results do not agree with previous reports. Specific literature has reported higher seropositivity in males due to their increased likelihood of contact with tick species compared to females, owing to behavioural characteristics [54].

The present study found an overall seroprevalence of 10.6% for co-infections. Previous studies indicated that co-infections in vertebrate hosts, often arising from concurrent or sequential exposure

to distinct tick species or the transmission of multiple pathogens by a single tick species, are frequent and can complicate both diagnosis and treatment, potentially increasing the likelihood of severe disease. Typically, co-infections involve pathogens that share a common vector and/or have overlapping geographical ranges [55–57].

Concerning ehrlichiosis, there is no long-lasting or adequately efficient immune response to ensure the host protection in the event of recurring infection [58]. Distinguishing between potential reinfections and persistent subclinical conditions is difficult. In most cases, dogs are returned to the environment where they lived before infection, creating conditions for repeated exposure to infected ticks. This circumstance emphasises the need for planned tick management strategies to keep susceptible dogs safe [59].

Our results showed no difference between dogs age and seropositivity for the studied pathogen species. This finding is not in line with previous studies [24,60] which found an association between older dogs and seropositivity, as antibodies to *R. conorii* can remain detectable for a long period [61]. Elderly dogs have a higher likelihood of exposure throughout their lives, and dogs in suboptimal physical condition may experience weakened immune systems, which increases the risk of infection [24,60].

In dogs with acute disease, PCR techniques for *E. canis* DNA are more sensitive than ELISA or IFAT for early detection of CME. For routine diagnosis of *E. canis* infection, PCR tests are commonly accessible. Several laboratories provide panels that include PCR assays for various vector-borne diseases. *E. canis* PCR tests can be done on blood, lymph node aspirates, splenic aspirates or bone marrow. For the diagnosis of chronic CME, convalescent IFAT or ELISA testing is far more sensitive than PCR assays [62,63].

Obtained results should be analysed with attention. An ELISA or IFAT positive result only indicates a past or present infection and does not necessarily reflect the current disease status. A positive outcome can be obtained based on antibody titres even if the condition has been resolved, as antibodies may persist in the body for several months or even years after initial infection [49]. Regardless of whether an active infection is present, an animal may be serologically negative, especially during the incubation period or the early stages of the illness. In ehrlichiosis, antibody synthesis typically commences 12 to 14 days after infection [49,64,65].

The sensitivity of detecting *Rickettsia* spp. in blood appears to be moderate to low in humans [67,68] and dogs [68,69]. The differences between serological and molecular tests are likely due to *Rickettsia* spp. circulating in low levels in the blood during the acute phase of illness [70] and being promptly removed from blood. Experimental infections of dogs with *R. conorii* resulted in a brief rickettsiemia lasting 2–10 days [71,72] which did not return even after immunosuppression [73].

The prevalence of ehrlichiosis is most significant in regions characterised by a high concentration of primary population vectors, specifically "hard" ticks belonging to genera *Rhipicephalus*, *Amblyomma* or *Dermacentor* [4]. The Mediterranean area offers an ideal habitat for the proliferation of numerous tick species. *R. sanguineus* and *Ixodes ricinus* are prevalent across Portugal and have been observed feeding on diverse hosts, including humans [74]. The same occurs for *R. conorii*, for which vectors are the primary mode of infection, and infected ticks are likely the most significant risk factor for infections in dogs and people [68]. However, in our study, no animal had ticks during blood collection. Since no ticks were observed, the low exposure of dogs in the two shelters, mainly to *R. sanguineus* ticks associated with adequate tick control programs, may explain the low seroprevalence in this study.

Because dogs in this study were apparently healthy, positive animals were most likely in the chronic phase of infection [75]. Shelters can serve as a public health warning system for zoonotic diseases like those caused by *E. canis* and *R. conorii*. Positive serological results in dogs serve as a valuable warning system for veterinarians. These results may indicate prior exposure to a pathogen. While they do not confirm the presence of disease, they trigger a need for further investigation to ensure the welfare of the individual dog. This proactive approach to healthcare helps veterinarians to promptly address any underlying health issues, contributing to their canine patients' overall health and welfare [26].

This work has some limitations since it was a cross-sectional study, with a small sample size of shelters and wide CI of the generated OR and self-selection bias. There are other limitations in the study: travel histories of these dogs were not obtained, so it is not possible to say that exposure to infected ticks was restricted to the location of the shelter of origin. The findings were considered preliminary until confirmed by molecular techniques, which can provide a high level of specificity in determining the particular rickettsia species responsible for these antibody responses.

Dogs kept at animal shelters are well-known for harbouring or transferring virulent pathogens to animals and humans. Because of their origin as unwanted animals, filthy living circumstances in shelters, high population density, stress and exposure to rodents and arthropod vectors, shelter dogs are excellent sentinels for several vector-borne and zoonotic diseases [26,76]. In our study, the seroprevalence values observed for *E. canis* were lower than those reported previously in owned dogs. This finding concerning lower seropositivity in shelters is quite surprising, as we could expect that those who live in shelters had high seropositivity. Due to increased environmental exposure and a lack of preventatives, dogs who do not receive veterinary care and those who enter shelters, of whom strays make up most cases, are likely more susceptible to contracting CVBD [76–78].

5. Conclusions

The findings of this study show that shelter dogs in northern Portugal are exposed to *E. canis* and *R. conorii*, which can affect both canines and humans. A One Health approach is necessary in order to educate the public about the hazards of canine zoonoses and developing legislation and procedures to control their spread and to preserve public health. Regular screening can provide valuable data on the distribution and prevalence of rickettsial diseases in shelter animals. This information should be used for epidemiological studies and surveillance, helping to understand disease patterns and to improve prevention strategies.

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Informed Consent Statement: Legal detainers or owners signed informed consent for the inclusion of dogs in this study.

Data Availability Statement: The data presented in this study are available upon request from the corresponding author.

Conflicts of Interest: The authors declare no conflict of interest. The funders had no role in the design of the study; in the collection, analyses, or interpretation of data; in the writing of the manuscript; or in the decision to publish the results.

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