

Survey of Vector-borne Comorbidities in a Leishmaniaendemic Canine Population from Paraguay

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Introduction

Canine Visceral Leishmaniasis (CVL) is an endemic parasitic disease in Paraguay with a majority of seropositive dogs observed to be asymptomatic for clinical signs consistent with CVL. Little is known about the full landscape of canine vectorborne diseases (CVBD) infecting dogs in Paraguay and the role concurrent infections may play in the onset of CVL. The objective of this study was to assess the incidence of Leishmania and other CVBD agents in a population of pet dogs from Leishmania-endemic regions of Paraguay.

Methods

A total of 100 serum samples and 400 whole blood samples were collected. The samples originated from 32 unique postal codes distributed across the eastern region of Paraguay. Nucleic acid was extracted from the whole blood samples and realtime PCR was run to detect the following pathogens: Leishmania; Ehrlichia canis; Ehrlichia chaffeensis; Ehrlichia ewingii; Anaplasma platys; Anaplasma phagocytophilum; Babesia vogeli; Babesia gibsoni; Babesia canis; Rickettsia spp.; Bartonella spp.; Mycoplasma haemocanis; Candidatus Mycoplasma haematoparvum; Hepatozoon canis; Hepatozoon americanum; Dirofilaria immitis; and Trypanosoma cruzi. ELISAs were performed on serum to detect the presence of antibodies to Leishmania, Ehrlichia spp., E. canis, E. ewingii, E. chaffeensis, Anaplasma spp., A. platys, A. phagocytophilum, Babesia gibsoni, Babesia canis/vogeli, and Borrelia burgdorferi, as well as antigen from D. immitis and Angiostrongylus vasorum.

Results

Serology Results (n=100)				
CVBD Pathogen	Pos, n	% Pos, n		
Leishmania	46	46%		
Ehrlichia spp.	51	51%		
Ehrlichia canis	51	51%		
E. chaffeensis	5	5%		
E. ewingii	3	3%		
Anaplasma spp.	46	46%		
A. platys	23	23%		
A. phagocytophilum	3	3%		
Babesia canis / B. vogeli	25	25%		
B. gibsoni	1	1%		
Borrelia burgdorferi	0	0%		
Dirofilaria immitis	1	1%		
Angiostrongylus vasorum	0	0%		
Leishmania + other CVBD	37	37%		



PCR Results (n=400)			
CVBD Pathogen	Pos, n	% Pos, n	
Leishmania	71	17.8%	
Trypanosoma cruzi	0	0.0%	
Ehrlichia canis	110	27.5%	
E. chaffeensis	0	0.0%	
E. ewingii	0	0.0%	
Anaplasma platys	45	11.3%	
A. phagocytophilum	0	0.0%	
Babesia vogeli	37	9.3%	
B. canis	0	0.0%	
B. gibsoni	0	0.0%	
Rickettsia spp.	1	0.3%	
Bartonella spp.	0	0.0%	
Mycoplasma haemocanis	15	3.8%	
Cand. M. haematoparvum	12	3.0%	
Hepatozoon canis	6	1.5%	
H. americanum	0	0.0%	
Dirofilaria immitis	0	0.0%	
Leishmania + other CVBD	28	7.0%	

Occurrence of CVBD co-infections Leish- positive samples by whole blood PCR (n=71)			
CVBD Pathogen	Pos, n	% Pos, n	
None	43	60.6%	
Ehrlichia canis	20	28.2%	
Anaplasma platys	5	7.0%	
Babesia vogeli	4	5.6%	
M. haemocanis	2	2.8%	
Cand. M. haematoparvum	2	2.8%	
Hepatozoon canis	1	1.4%	

Discussion

Real-time PCR detected Leishmania infections in this population at a rate of 18%. Additional CVBD pathogens detected by PCR included E. canis (28%), A. platys (11%), B. vogeli (9%), M. haemocanis (4%), Cand. M. haematoparvum (3%), and H. canis (2%). All of these pathogens are known to be transmitted by Rhipicephalus sanguineus ticks. Overall, 7% of samples were PCR positive for Leishmania and at least one other CVBD pathogen with E. canis being the most common co-infecting agent. Of the serum samples tested by ELISA, 46% were seropositive for Leishmania. High seropositive rates were also observed for Ehrlichia spp. (51%), E. canis (51%), Anaplasma spp. (46%), A. platys (23%), and Babesia spp. (26%). Overall, 37% of samples were seropositive to Leishmania and at least one additional CVBD pathogen. These findings indicate that co-infections with Leishmania and other CVBD pathogens, particularly those transmitted by Rhipicephalus sanguineus ticks, are relatively common in Paraguayan dogs. The potential for co-infection with other CVBD pathogens should be taken into consideration when diagnosing and treating CVL.

