

Short Communication

Prevalence of *Bartonella henselae* and *Bartonella clarridgeiae* in cats and dogs in Korea

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Blood, saliva, and nail samples were collected from 54 dogs and 151 cats and analyzed for the presence of *Bartonella henselae* with a novel nested polymerase chain reaction (PCR) method. *Bartonella (B.) henselae* was detected in feral cat blood (41.8%), saliva (44.1%), and nail (42.7%) samples. *B. henselae* was also detected in pet cat blood (33.3%), saliva (43.5%), and nail (29.5%) samples and in pet dog blood (16.6%), saliva (18.5%), and nail (29.6%) samples. Nine samples were infected with *B. clarridgeiae* and 2 were co-infected with *B. henselae* and *B. clarridgeiae* of blood samples of dogs. This report is the first to investigate the prevalence of *B. henselae* and *B. clarridgeiae* in dogs and cats in Korea, and suggests that dogs and cats may serve as potential *Bartonella* reservoirs.

Keywords: Bartonella, cats, cat-scratch disease, dogs, Korea

Introduction

The genus *Bartonella (B.)* includes at least 20 species and subspecies, and several of these are human pathogens [22]. Clinical manifestations of *Bartonella* infection include Carrion's disease, trench fever, cat scratch disease, bacillary angiomatosis, endocarditis, chronic bacteremia, neuroretinitis, and osteomyelitis [13]. Cat scratch disease is zoonotic and primarily caused by *B. henselae* [15]. *B. clarridgeiae* can also cause cat scratch fever. *B. henselae* and *B. clarridgeiae* can also infect dogs [4,7], and both species can function as bacterial reservoirs for infection [5,9,10,18,19]. Cat scratch disease was recently reported in a woman with a pet dog in Korea [5]. However, the prevalence of *Bartonella* spp. from companion animals in Korea has

not been previously investigated. We examined the prevalence of *B. henselae* and *B. clarridgeiae* in dogs and cats in the present study using a recently developed nested PCR method.

Blood, saliva, and nail samples were collected from healthy pet dogs (n = 54) and cats (n = 48) at the Veterinary Medical Teaching Hospital of Seoul National University, Korea. All samples were collected from November 2005 to July 2006. Feral cats (n = 103) were captured in neighborhoods throughout Seoul and were isolated in an animal shelter. *B. henselae* strain Houston-1 (ATCC 49882) and *B. clarridgeiae* strain (ATCC 51734) were obtained from the American Type Culture Collection (USA) and used for positive control samples. Genomic DNA was extracted using Genomic Blood DNA and Genomic Cell/Tissue DNA Extraction Kits (iNtRoN Biotechnology, Korea), per the manufacturer's instructions. Primary PCR was performed with the P-bhenfa (5'-TCTTCGTTTCTCTTCTTCA-3') and P-benr1 (5'-CAAGCGCGCTCTAACC-3') primers which amplified *B. henselae* (186 bp) and *B. clarridgeiae* (168 bp) fragments. Nested PCR amplified *B. henselae* (152 bp) and *B. clarridgeiae* (134 bp) fragments with the N-bhenf1a (5'-GATGATCCCAAGCCTTCTGGC-3') and N-bhenr (5'-AACCAACTGAGCTACAAGCC-3') primers [15]. Primary and nested PCR reactions were performed as previously described [15].

All PCR products were analyzed by sequencing with an automated sequencer ABI 3100 Genetic Analyzer (Bionics, Korea) and results were confirmed to be from *B. henselae* (GeneBank access number DQ000494) and from *B. clarridgeiae* (GeneBank access number: DQ003029).

B. henselae was detected in 14.2% of blood samples (14/98), 3.9% of saliva samples (4/102), and 4.8% of nail samples (5/103) from feral cats. In contrast, only 6.3% (3/48) of blood samples from pet cats were positive for *B. henselae*. *B. henselae* was not detected in pet cat saliva samples (n = 46), pet cat nail samples (n = 44), or in any pet

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dog samples (n = 54).

B. henselae was detected in 41.8% (41/98) of blood samples, 44.1% (45/102) of saliva samples, and 42.7% (44/103) of nail samples from feral cats by nested PCR. In addition, 33.3% of blood samples (16/48), 43.5% of saliva samples (20/46), and 29.5% of nail samples (13/44) from pet cats were *B. henselae* positive. *B. henselae* DNA was also detected in 16.6% (9/54) blood samples, 18.5% (10/54) of saliva samples and 29.6% (16/54) of nail samples from dogs (Table 1).

B. clarridgeiae was detected in 2 feral cat blood samples, a feral cat saliva sample, 3 dog blood samples, a dog saliva sample, and 2 dog nail samples. Additionally, 2 samples (1 dog blood and 1 dog nail) were co-infected with *B. henselae* and *B. clarridgeiae* (Table 2). PCR product and DNA sequencing data are shown in Fig. 1.

Cats are usually the main zoonotic reservoir for *Bartonella* infection [14], although dogs may also serve as zoonotic reservoirs secondary to *B. henselae* and *B. clarridgeiae* infection [4,7]. Cat scratch disease was identified in a case with suspected human:canine transmission in Korea [5,18]. However, there are no current surveys evaluating *Bartonella* spp. prevalence in cats and dogs.

A previous study reported that 39% of cats were *B. henselae* positive among a population of 146 cats in Japan [11], and this result was significantly higher than the previous 7.2% prevalence among cats in Japan. Previous studies conducted in various countries identified higher *Bartonella* bacteremia prevalence in shelter cats than in pet cats [2,3,8,10]. The *Bartonella* prevalence in pet cats in the present study (33.3%) was significantly higher than the prevalence in previous studies, including Germany (13%), France (11%), and the Netherlands (22%) [1,5,17].

Conversely, the prevalence of *B. henselae* in sheltered cats (41.8%) was similar to the prevalence identified in other studies. These findings suggest that pet cats may serve as a reservoir for *B. henselae* infection to their owners. This is particularly relevant to immunocompromised pet owners.

B. henselae prevalence in cats is higher than *B. clarridgeiae* prevalence [16], but this may be dependent on age, sex, and type of breeding [6]. The *B. henselae* prevalence in cats and dogs was greater than *B. clarridgeiae* and was higher in cats than in dogs. These results supported previous studies which suggested that *B. henselae* was the major zoonotic pathogen. A recent survey of *Bartonella* seropositive healthy blood donor in Sweden demonstrated a similar prevalence to

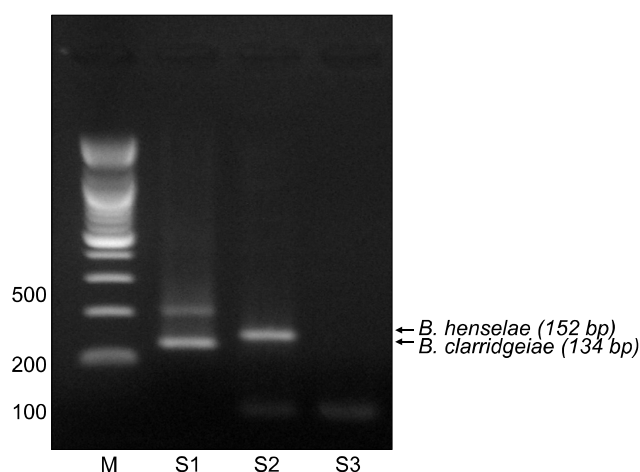


Fig. 1. *Bartonella* (*B.*) *henselae* (S2) and *B. clarridgeiae* (S1) nested PCR amplification bands from 2 cats. The negative control band (S3) is visualized on the right side. M: standard maker.

Table 1. Prevalence of *Bartonella henselae* infection in cat and dog blood, saliva, and nail samples detected by nested PCR

| Samples | Number of positive samples / Number of tested samples (%) | | | |
|---------|---|--------------|------------------------|--------------|
| | Feral cats | Pet cats | Subtotal (Feral + Pet) | Dogs |
| Blood | 41/98 (41.8) | 16/48 (33.3) | 57/146 (39.0) | 9/54 (16.6) |
| Saliva | 45/102 (44.1) | 20/46 (43.5) | 65/148 (43.9) | 10/54 (18.5) |
| Nails | 44/103 (42.7) | 13/44 (29.5) | 57/147 (38.8) | 16/54 (29.6) |

Table 2. Prevalence of *Bartonella clarridgeiae* infection in cat and dog blood, saliva, and nail samples detected by nested PCR

| Samples | Number of positive samples / Number of tested samples (%) | | | |
|---------|---|-------------|------------------------|-------------|
| | Feral cats | Pet cats | Subtotal (Feral + Pet) | Dogs |
| Blood | 2/98 (2.04) | 0/48 (0.00) | 2/146 (1.37) | 3/54 (5.56) |
| Saliva | 2/102 (1.96) | 0/46 (0.00) | 2/148 (1.35) | 1/54 (1.85) |
| Nails | 0/103 (0.00) | 0/44 (0.00) | 0/147 (0.00) | 1/54 (1.85) |

dogs in the present study [12]. Zoonotic diseases have become an increasingly important public health concern [5]. Our results suggest that *B. henselae* and *B. clarridgeiae* are highly prevalent in Korean cats and dogs. Further, cats and dogs may serve as reservoirs for human *Bartonella* infection.

In conclusion, data from the present study suggests that *Bartonella* infection prevalence in Korean shelter cats is similar to those of previously described countries. However, the prevalence of *B. henselae* in Korean pet cats was higher than reported prevalence in other countries. This is the first report examining the prevalence of *B. henselae* and *B. clarridgeiae* infection in domestic cats and dogs in Korea.

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