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 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-bh enfa (5'-TCTTCGTTTCTCTTCTTCTTCA-3') and P-benr1 (5'-CAAGCGCGCGCTCTACC-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'-TCTCTGTTTCTCTTCTCTTA ACC-3') and P-benr1 (5'-CAAGCGCGCGCTCTTA ACC-3') primers which amplified *B. henselae* (186 bp) and *B. clarridgeiae* (168 bp) fragments. 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...
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

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**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) |

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**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.
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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References

1. Bergmans AM, de Jong CM, van Amerongen G, Schot CS, Schols L.M. Prevalence of *Bartonella* species in domestic cats in The Netherlands. J Clin Microbiol 1997, 35, 2256-2261.
2. Branley J, Wolfson C, Waters P, Gottlieb T, Bradbury R. Prevalence of *Bartonella henselae* bacteremia, the causative agent of cat scratch disease, in an Australian cat population. Pathology 1996, 28, 262-265.
3. Chomel BB, Carlos ET, Kasten RW, Yamamoto K, Chang CC, Carlos RS, Abenes MV, Pajares CM. *Bartonella henselae* and *Bartonella clarridgeiae* infection in domestic cats from The Philippines. Am J Trop Med Hyg 1999, 60, 593-597.
4. Chomel BB, Mac Donald KA, Kasten RW, Chang CC, Wey AC, Foley JE, Thomas WP, Kittleson MD. Aortic valve endocarditis in a dog due to *Bartonella clarridgeiae*. J Clin Microbiol 2001, 39, 3548-3554.
5. Chung JY, Han TH, Kim BN, Yoo YS, Lim SJ. Detection of *Bartonella henselae* DNA by polymerase chain reaction in a patient with cat scratch disease: a case report. J Korean Med Sci 2005, 20, 888-891.
6. Engvall EO, Brändström B, Fernér C, Blomqvist G, Englund L. Prevalence of *Bartonella henselae* in young, healthy cats in Sweden. Vet Rec 2003, 152, 366-369.
7. Gundi VA, Bourry O, Davous B, Raoult D, La Scola B. *Bartonella clarridgeiae* and *B. henselae* in dogs, Gabon. Emerg Infect Dis 2004, 10, 2261-2262.
8. Heller R, Artois M, Xemar V, De Briel D, Gehin H, Jaulhac B, Monteil H, Piemont Y. Prevalence of *Bartonella henselae* and *Bartonella clarridgeiae* in stray cats. J Clin Microbiol 1997, 35, 1327-1331.
9. Keret D, Giladi M, Kletter Y, Wientroub S. Cat-scratch disease osteomyelitis from a dog scratch. J Bone Joint Surg Br 1998, 80, 766-767.
10. Marston EL, Finkel B, Regnery RL, Winoto II, Graham RR, Wignal S, Simanjuntak G, Olson JG. Prevalence of *Bartonella henselae* and *Bartonella clarridgeiae* in an urban Indonesian cat population. Clin Diagn Lab Immunol 1999, 6, 41-44.
11. Maruyama S, Nakamura Y, Kabeya H, Tanaka S, Sakai T, Katsube Y. Prevalence of *Bartonella henselae*, *Bartonella clarridgeiae* and the 16S rRNA gene types of *Bartonella* species among pet cats in Japan. J Vet Med Sci 2000, 62, 273-279.
12. McGill S, Wesslén L, Hjelm E, Holmberg M, Auvinen MK, Berggren K, Grandin-Jari B, Johnson U, Wikström S, Friman G. *Bartonella* spp. seroprevalence in healthy Swedish blood donors. Scand J Infect Dis 2005, 37, 723-730.
13. Molia S, Chomel BB, Kasten RW, Leutenegger CM, Steele BR, Marker L, Martenson JS, Keet DF, Bengis RG, Peterson RP, Munson L, O’Brien SJ. Prevalence of *Bartonella* infection in wild African lions (Panthera leo) and cheetahs (Acinonyx jubatus). Vet Microbiol 2004, 100, 31-41.
14. Pons I, Sanfeliu I, Quesada M, Anton E, Sampere M, Font B, Pla J, Segura F. Prevalence of *Bartonella henselae* in cats in Catalonia, Spain. Am J Trop Med Hyg 2005, 72, 453-457.
15. Rampersad JN, Watkins JD, Samalal MS, Deonanan R, Ramsubeik S, Ammons DR. A nested-PCR with an Internal Amplification Control for the detection and differentiation of *Bartonella henselae* and *B. clarridgeiae*: an examination of cats in Trinidad. BMC Infect Dis 2005, 5, 63.
16. Rolain JM, Locatelli C, Chabanne L, Davoust B, Raoult D. Prevalence of *Bartonella clarridgeiae* and *Bartonella henselae* in domestic cats from France and detection of the organisms in erythrocytes by immunofluorescence. Clin Diagn Lab Immunol 2004, 11, 423-425.
17. Sander A, Bühler C, Pelz K, von Cramm E, Bredt W. Detection and identification of two *Bartonella henselae* variants in domestic cats in Germany. J Clin Microbiol 1997, 35, 584-587.
18. Tsukahara M, Tsuneoka H, Iino H, Okuno K, Murano I. *Bartonella henselae* infection from a dog. Lancet 1998, 352, 1682.
19. Windsor JJ. Cat-scratch disease: epidemiology, aetiology and treatment. Br J Biomed Sci 2001, 58, 101-110.
20. Yamamoto K, Chomel BB, Kasten RW, Hew CM, Weber DK, Lee WI, Koehler JE, Pedersen NC. Infection and re-infection of domestic cats with various *Bartonella* species or types: *B. henselae* type I is protective against heterologous challenge with *B. henselae* type II. Vet Microbiol 2003, 92, 73-86.