

PCR detection of *Bartonella* spp. in the dog

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Abstract

Our study aimed at using PCR to identify the incidence of *Bartonella* spp. in blood of dogs. Altogether 286 dogs of 92 breeds aged 3 month to 17 years were tested from October 2008 to December 2009. Healthy dogs as well as dogs with various clinical symptoms of disease were included in the group. Samples were tested by polymerase chain reaction (PCR) specific for the presence of *Bartonella* spp. Following the DNA examination in 286 dogs by PCR and subsequent sequencing, two samples were identified as *Bartonella henselae* (0.7%). Other species of *Bartonella* were not found. It was the first time in the Czech Republic when incidence of *Bartonella* spp. was determined in dogs.

Vector born disease, zoonosis, infection, PCR diagnostics, cat scratch disease, B. henselae

In the last two decades, more than twenty *Bartonella* species or subspecies have been described. At least eleven of them are considered as human pathogens (Boulouis et al. 2005; Maggi et al. 2012b). Out of these eleven, six were identified also in dogs and cats. Cats are a major reservoir of *B. henselae*, *B. clarridgeiae*, *B. koehlerae*, and *B. bovis* (Gurfield et al. 1997; Bermond et al. 2002). Dogs can be infected with *B. vinsonii* subsp. *berkhoffii*, *B. henselae*, *B. clarridgeiae*, *B. washoensis*, *B. elizabethae*, and *B. quintana* (Chomel et al. 2006; Beerlage et al. 2012).

In the last few years, new *Bartonella* species have been isolated also from many wild rodents and ruminants. Some recent studies indicate that these new species could be responsible for certain human and canine infections (Boulouis et al. 2005; Maggi et al. 2012a).

In general, *Bartonella* bacteria induce none or only mild clinical signs in the natural hosts. However, infection of other than natural, i.e. accidental host can result in a serious or life-threatening disease.

In humans, this bacterium causes the so-called cat scratch disease (CSD) which is spread worldwide. In children, this infection is a relatively common cause of fever of unknown origin (Carithers 1985; Boulouis et al. 2005). In immunocompromised patients this infection has a serious or even fatal course.

Bartonella was first isolated in dogs in 1993 (Breitschwerdt et al. 1995). Ticks were identified as the bartonellosis vector in dogs; in Europe the most prominent are *I. ricinus* (Billeter et al. 2008; Cotté et al. 2008). The most important in dogs is *Bartonella vinsonii* subsp. *berkhoffii*. It has been proven that this species plays an important role in cases of endocarditis in dogs and humans (Breitschwerdt et al. 1999). Dogs can be further infected with *B. henselae*, *B. clarridgeiae*, *B. washoensis*, *B. elizabethae*, and *B. quintana* (Mexas et al. 2002; Chomel et al. 2006; Saunders and Monroe 2006; Morales et al. 2007; Diniz et al. 2009).

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There is not enough information about bartonellosis in dogs in various countries. The incidence is higher in warmer climates. There are several studies on seroprevalence of *Bartonella* antibodies in dogs, however, none of these studies reported the detection of DNA of individual *Bartonella* species in dogs (Guptill 2003; Solano-Gallego et al. 2004; Boulouis et al. 2005; Parpaglia et al. 2007; Yabsley et al. 2008). In this study we therefore determined the incidence of *Bartonella* spp. in dogs by PCR.

Materials and Methods

Altogether 286 dogs of 92 breeds were tested from October 2008 to December 2009. The examined dogs were patients of the Small Animal Clinic coming from Brno and the South Moravian Region of the Czech Republic. Crossbreeds prevailed ($n = 35$) followed by Dachshunds ($n = 31$) and German Shepherds ($n = 15$). The group included 124 females, of these 39 neutered; and 162 males, of these 12 castrated. The age of the animals ranged from 3 months to 17 years (mean 6.5 year). Their weight varied from 2 to 62 kg. A total of 51 healthy dogs plus 235 dogs with various clinical symptoms of disease were included in the group. Standard haematological and biochemical blood tests were carried out for most of the dogs. Blood was collected from v. cephalica antebrachii or v. jugularis. From each dog we obtained samples of whole blood. Samples were stored at -18°C until testing.

Whole blood samples were tested by PCR. DNA was extracted from samples using the commercial QIAGEN NucleoSpin Blood kit (Machery-Nagel, Germany). DNA samples were tested with the use of primers: forward 5'-(C/T)CTTCGTTTCTCTTTCTTCA-3' and reverse 5'-AACCAACTGAGCTACAAGCC-3' according to Jensen et al. (2000). The estimated size of the final product in the case of *B. henselae* was 172 bp, *B. clarridgeiae* 154 bp, *B. elizabethae* 241 bp, *B. quintana* 157 bp, and *B. vinsonii* subsp. *berkhoffii* 260 bp. As positive control we used the DNA of *B. henselae*. The PCR reaction was carried out in a total amount of 20 μl , using 2 μl of DNA, 10 μl of Combi PPP Master Mix (Top-Bio), 0.1 μl of each primer diluted to 100 pmol/ μl and 7.8 μl PCR water. Biometra T Personal Thermocycler (Whatman Biometra, Göttingen, Germany) was used for the reaction. The PCR product was visualized by ethidium bromide and 3% agarose gel (70 ml TAE, 2.1 g agarose, 3 μl ethidium bromide – stock solution 10 mg/ml in PCR water) at the voltage of 120 V. The electrophoresis was carried out using the Standard Power Pack P25T and horizontal electrophoresis by Cleaver Scientific Ltd. Transilluminator Vilber Lourmat (Deutschland GmbH, Germany) was used for visualisation.

PCR products from positive samples were purified from the agarose gel using QIAquick Gel Extraction kit (Qiagen, Machery-Nagel, Germany) and then subjected to sequencing (Macrogen, Seoul, Korea).

Results

Following the DNA examination in 286 patients by PCR and subsequent sequencing, two samples were identified as *Bartonella henselae* (0.7%). Other species of *Bartonella* were not found.

The first positive dog

Eight-year-old neutered female Poodle weighing 7 kg was presented at the clinic with signs of lethargy and loss of appetite. The clinical examination revealed slight enlargement of the mandibular and superficial cervical lymph nodes, and a body temperature of 39.6°C . A blood sample was collected in December 2008. Haematological examination revealed mild thrombocytopenia ($183 \times 10^9/\text{l}$); other basic haematological and biochemical indicators were within the physiological range.

The second positive dog

It was a 2-year-old Border Collie dog weighing 21 kg. The dog showed no clinical signs of disease. It was brought for a preventive check. A blood sample was collected in March 2009. Haematological examination revealed only lymphopaenia ($0.783 \times 10^9/\text{l}$) and slightly elevated ALT activity (1.69 $\mu\text{kat/l}$); other indicators were within the physiological range.

Discussion

Bartonella has a potential to cause serious disease in humans as well as in a wide range of domestic and wild animals. Cats are usually without clinical symptoms or they are only

nonspecific. But they are important reservoirs of *B. henselae* causing the cat scratch disease (CSD). Dogs are considered to be the reservoir of *B. vinsonii* subsp. *berkhoffii*. It has been demonstrated, however, that they can be infected with other species including *Bartonella henselae* (Mexas et al. 2002; Chomel et al. 2006; Saunders and Monroe 2006; Morales et al. 2007; Diniz et al. 2009). *Bartonella henselae* has been described in a number of cases of dog diseases (Guptill 2003). These were mostly chronic diseases characterized by apathy, loss of appetite, weight loss, exercise reluctance, ataxia, lymphadenopathy and other, often nonspecific symptoms. Blood is usually tested for thrombocytopaenia. Surprisingly enough, in our group we detected *B. henselae* only in two dogs (0.7%). Our first patient had symptoms corresponding to those of bartonellosis (lethargy, loss of appetite, lymphadenopathy) at the time of examination and the thrombocytopenia finding was appropriate. However, the described clinical and laboratory results may be identical in other infections and diseases and this is why *B. henselae* cannot be identified as the causative agent. The second positive dog was brought to the clinic for preventive check showing no signs of disease.

Bartonellas are transmitted by vectors. In case of cats the vectors are fleas, in particular cat fleas (*Ctenocephalides felis*), ticks and other blood-sucking insects. Ticks were identified as the vector in dogs. The most important tick in Europe is *Ixodes ricinus* described in the study by Cotté et al. (2008) as a competent vector for *B. henselae*. Transmission is possible even through the saliva in dog bites. People can be infected by ticks, cat and dog scratches or bites (Boulouis et al. 2005; Podsiadly et al. 2007; Billeter et al. 2008; Maggi et al. 2012a).

By PCR testing of whole blood we established the presence of *B. henselae* in two dogs (0.7%). A Polish study by Podsiadly et al. (2007) reports that they examined 54 dogs for *Bartonella* spp. DNA in the blood. DNA was not identified in a single case but they detected the presence of specific IgG antibodies against *B. henselae* in 50% of tested dogs. Many authors state that if *Bartonella* DNA is detected in an organism, the serology is negative (Duncan et al. 2007; Morales et al. 2007; Duncan et al. 2008; Cherry et al. 2009; Diniz et al. 2009).

Even though dogs are natural hosts of *B. vinsonii* subsp. *berkhoffii*, in our study we detected only the presence of *B. henselae*. However, this is a serious zoonotic agent and our results show the necessity to consider dogs as a potential source of human infection. The prevention of bartonellosis generally lies in the protection of animals against flea and tick infestation. It is therefore advisable to treat animals with antiparasitics on a regular basis, either in the form of spray, spot on drugs or collars.

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