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#### Bartonellosis

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#### 1 Abstract

2	Bartonellosis is a constellation of clinical conditions affecting human beings and a variety
3	of animals. Many Bartonella infections are zoonotic, with some of the most commonly reported
4	zoonotic manifestations of infection including cat scratch disease, bacillary angiomatosis,
5	endocarditis, and neuroretinitis. Companion animals serve as reservoirs for several zoonotic
6	species of Bartonella, and may also serve as sentinels for zoonotic Bartonella species harbored
7	by wildlife.
8	This article provides an overview of bartonellosis of dogs and cats, and discusses the public
9	health implications of animal bartonellosis.
10	
11	Keywords: Bartonella, bartonellosis, cat, dog, zoonosis
12	
13	INTRODUCTION
14	Bartonella spp. are small, vector-transmitted Gram-negative bacteria. Since the early
15	1990s many Bartonella species have been described, and there are now over 22 named species of
16	Bartonella, including species that originally comprised the genera Rochalimaea and Grahamella
17	(Brenner et al., 1993). The Bartonellas are phylogenetically close to the Rickettsiae and bacteria
18	of the species Brucella, Agrobacterium, and Afipia (Birtles et al., 1995; Brenner et al., 1993;
19	Weisburg et al., 1985). These relationships have provided some insights into the pathogenesis
20	and natural history of <i>Bartonella</i> infections.
21	The first Bartonella species described was B. bacilliformis, named for Peruvian
22	microbiologist Alberto Barton, and endemic to South America. Human bartonellosis caused by
22	<b>B</b> basilliformic (not considered zoonotic) is transmitted by condition and consists of two aligned

conditions; a devastating hemolytic anemia, Oroya fever or Carrión's disease, and verruga
peruana or Peruvian wart, a systemic disorder with verrucous cutaneous lesions that result from
endothelial cell proliferation. Carrión's disease is named for Peruvian medical student Daniel
Carrión, who in 1885 inoculated himself with material from a cutaneous lesion of verruga
peruana, documented his ensuing illness, demonstrating the relationship between the two
conditions, and sadly, succumbed to the disease.

The role of *Bartonella* species as pathogens of human beings and domestic animals is the subject of much ongoing investigation. Establishing true causal associations of *Bartonella* infections with clinical disease has been challenging in many instances. Current investigations are helping to better define the potential virulence factors of some *Bartonella* species, and to contribute to an understanding of the pathogenesis of *Bartonella* infections. This paper focuses on bartonellosis of dogs and cats, and discusses public health implications of animal bartonellosis. ETIOLOGY AND EPIDEMIOLOGY

37 Bartonella species have a worldwide distribution with highest prevalences in areas where 38 conditions are most favorable for arthropod vectors. Feline B. henselae infection was first 39 reported in 1992 (Regnery et al, 1992), and canine B. vinsonii subsp. berkhoffii infection in 1994 40 (Breitschwerdt et al, 1995). Reports of *Bartonella* infections in many other animal species have 41 followed. High prevalences are reported for *B. henselae* in cats (up to 93 percent seroprevalence 42 in some cat colonies), B. bovis and B. chomelii in cattle (42 to 89% prevalence), B. vinsonii 43 subsp. berkhoffii in coyotes (35-71% seroprevalence), B. vinsonii subsp. berkhoffii in dogs (1-44 38% seroprevalence) and *B. henselae* in dogs (10-35% seroprevalence) (Chang et al, 2000a; 45 Chang et al, 2000b; Jameson et al, 1995; Cherry et al, 2009; Nutter et al, 2004; Solano-Gallego 46 et al, 2004; Henn et al, 2005; Suksawat et al, 2001; Maillard et al, 2006). Seroprevalences are

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47	higher in older cats than in younger animals, whereas prevalence of bacteremia is higher in
48	younger animals (Chang et al, 2000a; Maillard et al, 2006; Chomel et al, 1995).
49	Many Bartonella species appear to be well-adapted to extended survival in mammalian
50	reservoir hosts. Prolonged bacteremia in clinically normal reservoir hosts is considered common
51	(Chomel et al., 2003a; Chomel et al., 2009a; Jacomo et al., 2002). Cats appear to be the primary
52	mammalian reservoir for the important zoonotic species, B. henselae, and also for B.
53	clarridgeiae and possibly B. koehlerae (Chomel et al., 2004). Coyotes and dogs may be
54	mammalian reservoirs for <i>B. vinsonii</i> subsp. <i>berkhoffii</i> and dogs and foxes for <i>candidatus</i> B.
55	rochalimae (Henn et al, 2009; Chomel et al 2003a). Other examples of reservoir hosts include
56	cattle, which appear to be the mammalian reservoir for <i>B. bovis</i> , California ground squirrels for
57	candidatus B. washoensis, and human beings for B. quintana and B. bacilliformis (Bermond et
58	al., 2002; Kosoy et al., 2003). The mechanisms that support persistent Bartonella bacteremia in
59	mammals are being investigated. Location within erythrocytes and vascular endothelial cells is
60	believed to protect Bartonellas from antimicrobial agents (Rolain et al., 2003c). Immune system
61	avoidance via intracellular location, frequent genetic rearrangements, and alteration of outer
62	membrane proteins, is also considered important (Berghoff et al., 2007; Chomel et al., 2003a;
63	Chomel et al., 2009a; Jacomo et al., 2002; Mandle et al., 2005; Seubert et al., 2002). Genetically
64	diverse strains of B. henselae and of B. vinsonii subsp. berkhoffii are described (Maggi et al.,
65	2006). For example, there are two recognized 16S rRNA types of <i>B. henselae</i> and there are
66	subgroups within each type (Maruyama et al., 2001; Zeaiter et al., 2002). Other molecular
67	methods also demonstrate remarkable diversity among <i>B. henselae</i> isolates, (Berghoff et al., 2007;
68	Dillon et al., 2002; Iredell et al., 2003; Kyme et al., 2003; Li et al., 2006; Maruyama et al., 2000;
69	Iredell et al., 2002; Monteil et al., 2007). There is evidence of genomic variation in B. henselae

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70	during the course of infection in cats (Berghoff et al., 2007; Kabeya et al., 2002). Such variation
71	likely enhances the ability of B. henselae to persist in infected cats for prolonged periods, and may
72	be useful in furthering the understanding of the pathogenicity of various B. henselae isolates
73	(Arvand et al., 2001; Arvand et al., 2007).
74	Infection of a single host animal with more than one Bartonella species is reported in cats,
75	dogs, cattle, and species of wild mammals, including rodents (Cherry et al., 2009; Diniz et al., 2007;
76	Gurfield et al., 1997; Telfer et al., 2007). Additionally, co-infection with multiple genetic types of
77	B. henselae or B. vinsonii supbspecies berkhoffii is reported for cats and dogs, respectively
78	(Duncan et al., 2007a; Gurfield et al., 1997). Regional differences in prevalence of infection of
79	cats with different rRNA types of <i>B. henselae</i> are reported (Bergmans et al., 1997; Guptill et al.,
80	2004; Gurfield et al., 2001; Heller et al., 1997).
81	At least fourteen Bartonella species or candidate species are considered zoonotic or
82	potentially zoonotic. Of the zoonotic species, several may be transmitted to human beings via
83	companion animals, including B. henselae (cats, possibly dogs), B. vinsonii subsp. berkhoffii (dogs),
84	and B. koehlerae (possibly cats). Bartonella clarridgeiae and candidatus B. rochalimae are
85	considered potentially zoonotic, with transmission potentially via cats (B. clarridgeiae) or dogs
86	(potentially both species) (Avidor et al., 2004; Chomel et al., 2004; Chomel et al., 2009b). Whether
87	B. clarridgeiae has truly been associated with human disease is still debated. Bartonella quintana
88	has long been understood to be a human pathogen with human beings as the reservoir host, and has
89	not previously been considered zoonotic. However, recent reports have noted the presence of <i>B</i> .
90	quintana in cats and dogs, and one report suggested transmission of B. quintana from companion
91	animals to human beings (Breitschwerdt et al., 2007; La et al., 2005; Kelly et al, 2006). Candidatus

92 B. rochalimae may be zoonotic with foxes and dogs as possible reservoir hosts (Chomel et al.,

93 2009b).

94 More is known about *B. henselae* than is known for any of the other zoonotic *Bartonella* 95 species, and *B. henselae* may be the *Bartonella* species most commonly zoonotic. The 96 epidemiology of *Bartonella* infections in companion animals, wild animals, and human beings is 97 still being clarified. It is possible that species of *Bartonella* not previously considered a zoonotic 98 threat may also be transmitted to human beings from animals. New information about existing 99 Bartonella species, and the frequent addition of new Bartonella species to our knowledge base, 100 makes it challenging to remain up-to-date regarding the many possible forms and sources of 101 bartonellosis in animals and in human beings. For example, a recent publication reported that 102 Bartonella DNA of 4 species considered zoonotic or potentially zoonotic (B. grahamii, B. 103 elizabethae, B. washoensis, B. clarridgeiae) was detected in some exotic animals imported into 104 Japan as pets (Inoue et al., 2009). 105 TRANSMISSION AND PATHOGENESIS 106 Bartonella henselae is naturally transmitted among cats by cat fleas (Ctenocephalides 107 felis felis). Chronic recurring bacteremia is common in infected cats, and likely facilitates vector 108 transmission, similar to what is documented for other pathogens that are transmitted by

109 hematophagous arthropod vectors (Barbour and Restrepo, 2000; Breitschwerdt and Kordick,

110 2000; Schulein et al., 2001). B. henselae was experimentally transmitted among cats by

111 transferring fleas fed on naturally infected cats to specific pathogen free (SPF) cats, and by

112 intradermal (ID) inoculation of excrement collected from fleas fed on B. henselae-infected cats

113 (Chomel et al., 1996; Finkelstein et al., 2002; Foil et al., 1999). Transmission via flea saliva has

not been documented (Foil et al., 1999). Cats were also experimentally infected with *B. henselae* 

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115	and B. clarridgeiae through IV or IM inoculation with infected cat blood, and B. henselae was
116	transmitted by intravenous, subcutaneous, ID, or oral routes of inoculation with plate-grown
117	bacteria (Abbott et al., 1997; Guptill et al., 1997; Guptill et al., 1999; Kordick et al., 1999;
118	Kordick and Breitschwerdt, 1997; Mikolajczyk and O'Reilly, 2000; O'Reilly et al., 1999).
119	Bartonella henselae transmission has not occurred when infected cats cohabit with uninfected
120	cats in a flea-free environment, indicating that transmission among cats does not occur through
121	cat bites, scratches, grooming, or sharing of food dishes and litter boxes. In addition, there was
122	no transmission between bacteremic female cats and uninfected males during mating, or to the
123	kittens of infected females either during gestation or in the neonatal period, also in flea-free
124	environments (Abbott et al., 1997; Guptill et al., 1997; Guptill et al., 1998).
125	Ticks have been proposed as vectors for transmission of Bartonella species among cats,
126	human beings, dogs, and other mammalian hosts (Bergmans, 1996; Cotté et al., 2008;
127	Pappalardo et al., 1997; Slater et al., 1990; Welch et al., 1999). Bartonella henselae and other
128	Bartonella spp. were detected by PCR in questing ticks (Chang et al., 2002b; Chang et al., 2001;
129	Sanogo et al., 2003). Transtadial transmission of B. henselae was recently demonstrated in Ixodes
130	ricinus ticks and these ticks are likely competent vectors for B. henselae (Cotté et al., 2008). Deer
131	keds likely transmit B. schoenbuchensis, and human body lice transmit B. quintana (Dehio et al.,
132	2004; Raoult and Roux, 1999; Reeves et al., 2006). Bartonella species have been detected in
133	other biting flies, and in many species of fleas and ticks in addition to those already noted
134	(Chomel et al., 2009a; Marié et al., 2006; Parola et al., 2002; Rolain et al., 2003a).
135	Virulence factors of <i>Bartonella</i> species
136	A comprehensive review of virulence factors of <i>Bartonella</i> species is beyond the scope of

this paper. Investigation of potential virulence factors for most *Bartonella* species is ongoing in

138 several laboratories; the reader is referred to a recent comprehensive review of *Bartonella* 139 virulence factors and persistence mechanisms for detailed information (Minnick and Battisti, 140 2009). Virulence factors characterized in several *Bartonella* species include adhesins, heme 141 acquisition/utilization mechanisms, type IV secretion systems (T4SS), and a low-potency 142 lipopolysaccharide (LPS), among others (Minnick and Battisti, 2009). Interestingly, purified 143 LPS of *B. quintana* was recently shown to inhibit inflammatory changes in a mouse model of 144 immune-mediated arthritis via inhibition of TLR-4 activation (Abdollahi-Roodsaz et al., 2007). 145 Type IV secretion systems, while not identified in the type species *B. bacilliformis*, are important 146 virulence factors for other *Bartonella* species. The T4SS trw has been identified in *B. henselae*, 147 B. quintana, B. birtlesii, B. grahamii, and B. tribocorum, and is believed important in erythrocyte 148 and endothelial cell invasion (Minnick and Battisti, 2009; Dehio, 2008). The VirB/Vir D4 T4SS, 149 investigated in B. tribocorum and B. henselae, among others, has important effects including 150 activation of nuclear factor kappa B (NF-kB), leading to release of pro-inflammatory or pro-151 angiogenic cytokines, and inhibition of apoptosis of colonized endothelial cells (Dehio, 2008; 152 Minnick and Battisti, 2009). Further investigation of virulence and persistence mechanisms of 153 the Bartonella species will advance our understanding of the remarkable ability of these 154 organisms to persist in vertebrate hosts for prolonged periods.

#### 155 Relapsing bacteremia and immune protection

Experimentally infected cats maintained relapsing *B. henselae* or *B. clarridgeiae* bacteremia for as long as 454 days (Kordick et al., 1999). Relapsing bacteremia reported in some naturally infected cats appeared to result from reinfection via fleas, indicating a lack of immune protection following *Bartonella* infection (Arvand et al., 2008; Kordick et al., 1995).

160 Experimentally infected cats became immune to challenge infection with homologous 161 strains of *Bartonella*, but there was a lack of protection against re-infection with heterologous 162 Bartonella isolates (Greene et al, 1996; Yamamoto et al., 1997). The level of bacteremia and 163 degree of susceptibility to reinfection following challenge inoculation varied with the strain and 164 species of Bartonella. Cats infected with B. henselae 16S rRNA type I or II were susceptible to 165 challenge infection with B. clarridgeiae, and cats infected with B. koehlerae or B. clarridgeiae 166 were susceptible to challenge infection with B. henselae 16s rRNA type I or type II. Cats 167 previously infected with B. henselae 16S rRNA type II were resistant to re-infection with the 168 same B. henselae strain, but were susceptible to infection with B. henselae 16S rRNA type I 169 (Yamamoto et al., 1997). In contrast, cats infected with B. henselae 16S rRNA type I were, as 170 expected, protected from re-infection with the same strain, but were also partially or completely 171 protected against challenge infection with a *B. henselae* 16S rRNA type II isolate (Yamamoto et 172 al., 2003). The lack of or incomplete cross-protection among *Bartonella* isolates, likely due to 173 genetic variation, has made vaccine development difficult.

174 Relapsing bacteremia is also described in rats infected with *B. tribocorum*, and relapsing 175 illness in human beings is reported in conjunction with infection with several *Bartonella* species 176 (Dehio, 2001; Maguiña et al., 2009; Schülein et al., 2001). Relapsing bacteremia may be 177 facilitated by antigenic variation, or by localization to so-called "privileged" sites such as within 178 erythrocytes or other host cells, as has been demonstrated for *B. tribocorum* and other *Bartonella* 179 species.

180 The exact localization of *Bartonella* species in mammalian hosts has not been completely 181 determined, and may differ among *Bartonella*-host pairs. Evidence indicates that individual 182 *Bartonella* species have been associated with reservoir hosts for hundreds to thousands of years,

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183 and this long period of "co-evolution" has likely led to numerous means of immune evasion and 184 prolonged coexistence of *Bartonella* spp. and various reservoir hosts (Drancourt et al., 2005; La 185 et al., 2004). Bartonella sp. other than B. bacilliformis have been detected within erythrocytes of 186 naturally infected cats and human beings (Mehock et al., 1998; Rolain et al., 2002). Bartonella 187 may also localize and persist within vascular endothelial cells, as suggested by an in vivo model 188 of B. tribocorum infection of rats (Dehio, 2001). Studies in a rat model of B. tribocorum 189 infection suggest that *Bartonella* may, following initial inoculation, colonize vascular endothelial 190 cells as a primary niche. In this rat model, *Bartonella* were shown to periodically 191 (approximately every 5 days) leave a primary niche and invade circulating erythrocytes, in which 192 protected environment they then remained for the normal lifespan of the erythrocytes (Schülein 193 et al., 2001). The details of host/pathogen interactions in other *Bartonella*/reservoir host pairs 194 have not yet been reported.

195 CLINICAL FINDINGS IN CATS AND DOGS

196 <u>Cats</u>. Existing data indicate that few cats naturally infected with *Bartonella* have notable 197 clinical signs. Given the long association of *Bartonella* and domestic cats, there have likely been 198 adaptations of the host and parasite to facilitate co-existence and minimize pathogenic effects on 199 the mammalian host. Occurrence and severity of clinical signs likely varies with the strain of 200 Bartonella infecting the cat (Mikolajczyk and O'Reilly, 2000; O'Reilly et al., 1999). Naturally 201 infected cats may have mild clinical signs that owners do not observe. Some naturally infected 202 cats developed fever following elective surgical procedures (Breitschwerdt, 1997). Uveitis may 203 be another manifestation of natural Bartonella infection (Lappin and Black, 1999). Bartonella 204 henselae type I was associated with fatal blood culture-negative vegetative aortic valve 205 endocarditis in two naturally infected cats (Chomel et al., 2003c; Chomel et al., 2009c). Whether

206	Bartonella spp. are present in lymph nodes of some cats with persistent lymphadenomegaly or some
207	cases of peliosis hepatis in cats is unknown (Brown et al., 1994; Kirkpatrick et al., 1989).
208	The majority of information regarding clinical signs is available from experimental studies
209	in which cats were infected with <i>B. henselae</i> . A few cats were experimentally infected with <i>B</i> .
210	clarridgeiae and B. henselae together, and clinical findings in these cats paralleled those reported
211	for cats infected with <i>B. henselae</i> alone. Cats experimentally infected with <i>B. koehlerae</i> or <i>B</i> .
212	rochalimae exhibited no clinical signs (Chomel et al., 2009b; Yamamoto et al., 2002).
213	Clinical signs in cats experimentally infected with B. henselae were usually mild though
214	varied with the strain of <i>B. henselae</i> used for inoculation. Cats inoculated intradermally
215	developed areas of induration or abscesses at inoculation sites, between approximately 2 days
216	and 3 to 4 weeks after inoculation, from some of which pure cultures of <i>B. henselae</i> were
217	obtained (Greene et al., 1996; Guptill et al., 1997; Kordick et al., 1999; Mikolajczyk and
218	O'Reilly, 2000; O'Reilly et al., 1999). Other transient clinical findings in many experimentally
219	infected cats included generalized or localized peripheral lymphadenomegaly (lasting for about 6
220	weeks following inoculation), and short periods of fever (>39.4°C [103°F]) during the first 48 to
221	96 hours following inoculation and again at approximately 2 weeks following inoculation. There
222	were instances of mild neurologic signs (nystagmus, whole body tremors, mild focal motor
223	seizures, either decreased or exaggerated responses to external stimuli, behavior changes), and
224	epaxial muscle pain in a few cats. Some cats were lethargic and anorexic during febrile periods
225	(Guptill et al., 1997; Kordick et al., 1999; Kordick and Breitschwerdt, 1997; Mikolajczyk and
226	O'Reilly, 2000; O'Reilly et al., 1999). Reproductive failure occurred in some cats experimentally
227	infected via intradermal inoculation with B. henselae (Guptill et al., 1998; Abbott et al., 1997).

Some cats infected via fleas exhibited no clinical signs while in one study two cats developed
cardiac disease (Bradbury and Lappin, 2009; Chomel et al., 1996).

230 A potential causative role of *Bartonella* spp. in chronic diseases of cats has been 231 proposed because Bartonella bacteremia is common and prolonged. For example, results of a 232 study in Japan suggested that cats seropositive for both *B. henselae* and feline immunodeficiency 233 virus were more likely to have gingivitis or lymphadenomegaly than cats seropositive for either 234 agent alone (Ueno et al., 1996). Results of a Swiss study suggested possible associations 235 between B. henselae seropositivity and stomatitis or unspecified urinary tract disorders (Glaus et 236 al., 1997). However, because of the high prevalence of Bartonella exposure in the domestic cat 237 population, extensive, carefully controlled epidemiologic investigations are needed to determine 238 whether particular clinical conditions are truly associated with *B. henselae* infections in cats. 239 Some small epidemiologic studies have evaluated the association between *Bartonella* exposure 240 and clinical conditions such as some neurologic conditions, stomatitis, uveitis, and fever. To 241 date, there has not been statistically significant association of documented active Bartonella 242 infection with the aforementioned conditions in naturally infected cats (Fontenelle et al., 2008; 243 Lappin et al., 2008; Pearce et al., 2006; Quimby et al., 2008). Clinical conditions that have been 244 proposed as attributable to feline bartonellosis may also result from other etiologies, and 245 determining in which cats *Bartonella* infection in fact causes clinical signs is difficult. 246 **Dogs**: There are case reports of dogs with varying clinical conditions from which *Bartonella* 

247 DNA has been detected by polymerase chain reaction amplification of various gene fragments,

or less commonly, *Bartonella* has been isolated (Cross et al., 2008; Leibovitz et al., 2008; Mellor

et al., 2006; Saunders and Monroe, 2006). There are fewer data available for dogs than for cats

250 experimentally infected with *Bartonella*. Dogs experimentally infected with *B. vinsonii* subsp.

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251	berkhoffii exhibited no clinical signs other than transient fever in a few dogs. Infected dogs had
252	some evidence of mild immunosuppression including either temporary cyclic decreases or mild
253	sustained decreases in peripheral blood CD8+ lymphocyte numbers (Pappalardo et al., 2000a;
254	Pappalardo et al., 2001). Two dogs experimentally infected with <i>B. rochalimae</i> exhibited no
255	clinical abnormalities other than inflammation at the inoculation site (Chomel et al., 2009b).
256	Some conditions with which Bartonella spp. have been associated, as a result of detection of
257	Bartonella in tissues of clinically ill dogs, include endocarditis (B. vinsonii subsp. berkhoffii, B.
258	clarridgeiae, B. quintana, candidatus B. washoensis, B. rochalimae), cardiac arrhythmias (B.
259	vinsonii subsp. berkhoffii), granulomatous rhinitis (B. vinsonii subsp. berkhoffii), peliosis hepatis
260	(B. henselae), meningoradiculoneuritis with dermatitis or panniculitis (B. vinsonii subsp.
261	berkhoffii), and granulomatous lymphadenitis (B. henselae) (Breitschwerdt et al., 1999; Chomel
262	et al., 2003b; Chomel et al., 2001; Cross et al., 2008; Henn et al., 2009; Kitchell et al., 2000;
263	Leibovitz et al., 2008; MacDonald et al., 2004; Mellor et al., 2006; Morales et al., 2007;
264	Pappalardo et al., 2000b). The majority of these reports involve only one or just a few dogs, with
265	endocarditis the exception. Endocarditis appears to be the clinical condition having the strongest
266	association with Bartonella infection in dogs, and most cases of canine endocarditis attributable
267	to Bartonella are caused by B. vinsonii subsp. berkhoffii. A recent study found no association
268	between Bartonella and chronic rhinosinusitis in dogs (Hawkins et al., 2008).

Whether *Bartonellae* are primary pathogens of dogs, or opportunistic pathogens, is not completely clear. For dogs and cats, the potential for co-infection with other vector-transmitted pathogens, and the likelihood that clinical conditions may be a result of co-infection rather than attributed to one potential pathogen, must always be considered when evaluating ill animals.

273 DIAGNOSIS

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Cats: The high prevalence of *Bartonella* bacteremia in cats makes establishing a causative
association between *Bartonella* infection and illness difficult. In addition to the potential need to
test for *Bartonella* infections in ill cats, veterinarians will be asked to test healthy pet cats
belonging to clients with *Bartonella*-related illnesses, or to screen healthy cats that are being
considered as pets, particularly for people considered most susceptible to *Bartonella* infections
(see Public Health).
Clinical Laboratory Findings and Pathologic Findings: Most experimentally infected cats had no

abnormalities on complete blood counts, serum biochemical tests, or urine analysis. Some cats
had transient anemia early in the course of infection, and some had persistent eosinophilia
(Kordick et al., 1999). Mature neutrophilia occurred in some cats, especially during periods of
skin inflammation (Guptill et al., 1997; Guptill et al., 1999). Acutely and chronically infected
cats had hyperplasia of lymphoid organs, and small foci of lymphocytic, pyogranulomatous, or
neutrophilic inflammation in multiple tissues (lung, liver, spleen, kidney, heart) (Guptill et al.,
1997; Kordick et al., 1999).

Finding *B. henselae* in erythrocytes of infected cats has not been an effective means of detection using conventional staining methods. Confocal microscopy or special fluorescence staining has been used to document intraerythrocytic location of *B. henselae*, *B. clarridgeiae* and *B. koehlerae* (Mehock et al., 1998; Rolain et al., 2003b; Rolain et al., 2004).

292 <u>Bacterial Isolation</u>: A positive blood culture result, or culture of other tissue, is the most reliable

test for definitive diagnosis of active *Bartonella* infection. However, because of the relapsing

294 nature of *Bartonella* bacteremia in cats, culture is not always a sensitive diagnostic tool. Blood

295 culture is indicated for sick cats whose history and clinical presentation suggest possible

296 Bartonella infection, or when a client's physician requires such testing of pet cats. Blood for

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297 culture should be obtained using sterile technique, and the blood placed in lysis centrifugation 298 blood culture tubes (Isolator tubes, Wampole, Cranbury, NJ) or EDTA-containing tubes. If 299 blood is collected into EDTA tubes the blood should be chilled or frozen during shipment, and 300 ideally, plastic EDTA tubes should be used. Culture of blood collected into EDTA and frozen 301 prior to culture was more sensitive than use of lysis centrifugation tubes for *Bartonella* culture 302 (Agan and Dolan, 2002; Brenner et al., 1997). Blood should be sent to laboratories familiar with 303 the culture of these fastidious organisms, and the laboratories should be contacted for specific 304 instructions regarding sample collection and submission. Special media (such as fresh chocolate 305 agar, or brain-heart infusion agar enriched with blood) and culture conditions (5%  $CO_2$ , 306 temperatures of 35 to 37 degrees Celsius), are used to enhance the likelihood of isolating 307 Bartonella from blood and other tissues. Extended culture times of up to 6-8 weeks are often 308 required (La Scola and Raoult, 1999). 309 Serologic Testing: Measuring serum antibodies has limited value for determining whether an ill 310 cat has an active Bartonella infection. Serum IgG antibodies persist in experimentally infected 311 cats for prolonged periods. How long antibodies persist following clearance of infection is not 312 known. Indirect fluorescent antibody (IFA), enzyme immunoassay (EIA), and Western blot tests 313 are available. Because of the genetic diversity of *Bartonella* organisms, infections with some 314 strains or species of *Bartonella* may be missed using any method, depending on the antigen 315 preparations used (Giladi et al., 2001; McCool et al., 2008). The positive predictive value (PPV) 316 of IFA or EIA (IgG) serologic tests for bacteremia in cats is only 39% to 46%; false positive tests 317 appear to be common. The utility of a negative serologic result is greater; false negatives are 318 uncommon and the negative predictive value of serologic tests for bacteremia or presence of 319 DNA in cat blood is high at 87% to 97% (Bergmans et al., 1997; Chomel et al., 1995; Guptill et

al., 2004; Gurfield et al., 2001; Lappin et al., 2008). No cut-off values allowing use of serologic
testing to determine whether a cat is currently infected with *Bartonella* have been established
(Yamamoto et al., 2003).

323 The use of Western blot tests has been advocated for serodiagnosis of feline B. henselae 324 infections, but the diagnostic accuracy of Western blot tests awaits further investigation. In 325 human medicine, variability in Western blot testing is still reported, and serologic responses of 326 people as evaluated by immunoblot vary (McCool et al., 2008; Sander et al., 2001). Results of one study in cats indicated no differences in Western blot patterns of cats evaluated over the 327 328 course of infection (Kordick et al., 1999). Results of another study indicated that antibodies in 329 sera of infected cats reacted with an increasing number of bands of polyacrylamide gel-separated 330 proteins over the course of infection (Freeland et al., 1999). In a recent report of naturally 331 infected cats, the PPV of a Western blot test for presence of Bartonella DNA in cat blood was reported to be only 18.8% (Lappin et al., 2008). 332

333 Nucleic Acid Detection: Standard PCR testing for the presence of Bartonella DNA may be no 334 more sensitive than blood culture for detection of active Bartonella infection, and detecting DNA 335 does not always equate to detection of living organisms. Nested PCR testing may increase 336 sensitivity for detection of *Bartonella* DNA in cat blood, although the risks of contamination 337 with this procedure are recognized, and laboratory precautions to minimize this risk are essential 338 (Roy et al., 2001). Real-time PCR testing also improves sensitivity with less likelihood of 339 contamination. Additional benefits of PCR testing are that the species and/or strain of 340 Bartonella may be identified by sequencing of the reaction product, and results of PCR testing 341 are available more quickly than those of blood culture (Fenollar and Raoult, 2004). Blood 342 samples for PCR testing should be obtained using sterile technique. Care must be taken in

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343	collection and sample processing to avoid sample contamination and DNA degradation.
344	Laboratories experienced in molecular diagnostics should be used, and should be contacted for
345	submission guidelines. Numerous primer pairs are utilized for PCR testing; these vary by
346	laboratory, and primer pairs utilized by commercial laboratories may be proprietary. Broad
347	range primers are useful for detecting the presence of any Bartonella DNA in clinical samples,
348	and more species-specific sets of primers may be utilized to determine Bartonella species.
349	Multiple approaches, including multiocus sequence typing (MLST) and multiple locus variable
350	number tandem repeat analysis (MLVA) have been utilized for B. henselae species typing
351	(Iredell et al., 2003; Monteil et al., 2007; Bouchouicha et al, 2009a).
352	<b>Dogs</b> : Many clinical conditions that have been proposed to be attributable to canine
353	bartonellosis may also result from other etiologies. Bartonella infection should always be
354	considered when working with dogs with endocarditis, and for some instances of granulomatous
355	disease, however, for other clinical entities, determining for which dogs clinical signs are likely
356	to be caused by Bartonella infection may be difficult. Clinical conditions may be a result of co-
357	infection with multiple pathogens, particularly multiple arthropod-transmitted pathogens.
358	Bacterial isolation: Culture appears to be insensitive for diagnosis of canine bartonellosis.
359	Blood cultures of experimentally infected dogs remained positive only through 42 days PI in
360	most dogs, whereas DNA was detected in peripheral lymph nodes by PCR at up to 142 days PI
361	and in the liver at 247 days PI (Pappalardo et al., 2000a; Pappalardo et al., 2001). It is possible
362	that DNA detected by PCR was that of organisms that were not viable, but this is considered
363	unlikely. The use of an insect growth medium enriched with multiple micronutrients and
364	defibrinated sheep blood for pre-enrichment culture of blood and potentially other tissues,
365	followed by PCR testing of the liquid culture and sub-inoculation of the liquid culture onto agar

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366 plates, may enhance the sensitivity of culture for diagnosis of *Bartonella* infections of dogs 367 (Duncan et al., 2007b; Maggi et al., 2005a).

368 Serologic testing: Serologic testing is available for several *Bartonella* species. Dogs may not

369 remain serologically positive for long periods. Experimentally infected dogs produced serum

370 IgG antibodies to *B. vinsonii* subsp. *berkhoffii* that peaked at reciprocal titers as high as 4096

371 within the first 1-4 weeks post-inoculation (PI). Antibody titers then declined but remained

372 increased above a reciprocal titer of 64 in some dogs through at least 243 days PI in one study

373 and 132 PI days in another study (Pappalardo et al., 2000a; Pappalardo et al., 2001).

374 There is growing evidence that dogs naturally infected with *Bartonella* (as determined by PCR

testing) may be seronegative (Diniz et al., 2007; Duncan et al., 2008). How widespread this 375

phenomenon is has yet to be determined, but it suggests that serology is not as sensitive a test for 376 377

dogs as it is for cats.

378 Nucleic Acid Detection: As noted for cats, PCR testing may be more sensitive than serology or 379 bacterial culture for detecting Bartonella DNA in dog tissues. Guidelines for molecular 380 diagnostic testing as noted for cats apply. A combination of use of an enriched culture medium 381 with PCR testing may be the most sensitive means of detecting *Bartonella* infection in dogs

382 (Duncan et al., 2007b).

383 THERAPY

384 Documenting clearance of Bartonella infections through antibiotic treatment is difficult 385 due to the prolonged relapsing bacteremia of most infected cats, and the potential difficulty in 386 detecting Bartonella in dogs. No regimen of antibiotic treatment has been proven effective, in 387 controlled studies with long-term follow-up, for definitively eliminating Bartonella infections in 388 cats or dogs (Bergmans et al., 1996; Greene et al., 1996; Kordick et al., 1997; Regnery et al.,

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389	1996). Enrofloxacin (5.4–7.6 mg/kg, given PO every 12 hours) for 14 or 28 days appeared to
390	clear B. henselae or B. clarridgeiae infection in 4 of 6 or 5 of 7 treated cats, respectively, that
391	were followed for 12 weeks after treatment (Kordick et al., 1997). However, enrofloxacin causes
392	retinal degeneration in cats, and use of doses of greater than 5 mg/kg per day is contraindicated
393	(Wiebe, 2002). Doxycycline (4-12 mg/kg, given PO, every 12 hours) cleared bacteremia in only
394	1 of 6 cats treated for 14 days and 1 of 2 cats treated for 28 days (Kordick et al., 1997).
395	Antibiotics tested in another study, including erythromycin 11 to 22 mg/kg every 8 hours),
396	amoxicillin (11 to 22 mg/kg every 8 hours), and tetracycline hydrochloride (13.75 mg/kg every 6
397	hours), rapidly decreased the level of bacteremia in infected cats. Treated and untreated cats
398	became blood-culture negative after the same period of time, however, making proof of efficacy
399	of antibiotics difficult (Regnery et al., 1996). In some studies, cats were not followed for more
400	than 8 weeks after treatment, thereby making it difficult to assess drug efficacy due to the
401	possibility of chronic relapsing bacteremia in those cats (Greene et al., 1996; Regnery et al.,
402	1996). Azithromycin (various doses), a macrolide antibiotic with good intracellular penetration, has
403	been recommended for treatment of infected dogs and cats, but data from controlled efficacy studies
404	with long-term follow-up are lacking. Azithromycin appears to have important immunomodulatory
405	and anti-inflammatory properties in addition to a broad antimicrobial spectrum of activity (Culic et
406	al., 2002; Labro, 2000; Labro and Abdelghaffar, 2001; Ortega et al., 2004). These properties may
407	make it difficult to determine whether beneficial effects reported following azithromycin treatment
408	of cats are a result of anti-Bartonella activity, or instead are a result of the antimicrobial action of
409	azithromycin on other bacteria, of azithromycin's other properties, or of some combination of all of
410	these.

411	Routine treatment with antibiotics may induce resistant strains of any bacterial species,
412	therefore treatment should be reserved for use only in Bartonella-positive animals showing clinical
413	signs. Client education regarding the uncertainty of treatment efficacy, the importance of prolonged
414	follow-up, and the possibility of re-infection following treatment, is necessary. The importance of
415	flea control and other means of preventing transmission (see Prevention) should be emphasized.
416	The American Association of Feline Practitioners Panel Report on feline Bartonella infections is a
417	good resource for questions regarding feline bartonellosis diagnosis, treatment and prevention
418	(Brunt et al., 2006).
419	Dogs showing clinical signs of disease may benefit from treatment for Bartonella infection.
420	The optimal antibiotic treatment regimen is not well-documented. Treatment with doxycycline,
421	azithromycin, enrofloxacin and other antibiotics has been reported (Breitschwerdt and Chomel,
422	2005). There are no published randomized controlled clinical trials of antibiotic treatment for
423	bartonellosis in dogs.
424	PREVENTION
425	Prevention of Bartonella infections is best accomplished by avoiding exposure to
426	infected animals and arthropod vectors. Vector control measures are effective for interrupting
427	transmission of <i>B. henselae</i> among cats (Bradbury and Lappin, 2009). Bartonella henselae and
428	B. clarridgeiae have been transmitted through inoculation of infected cat blood, therefore cats
429	should not receive blood transfusions from cats of unknown Bartonella status or from cats that
430	are culture or seropositive for <i>Bartonella</i> (Kordick and Breitschwerdt, 1997; Wardrop et al.,
431	2005). It is not known how long Bartonella may survive in stored blood; there is one report of
432	survival of <i>B. henselae</i> in stored human red blood cells for up to 35 days (Magalhães et al.,

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433 2008). Similar precautions may apply to dogs. A vaccine to prevent *Bartonella* infection in cats434 or dogs is not available.

435 BARTONELLOSIS IN OTHER ANIMAL SPECIES

Bartonella infections are reported in horses and some domestic and wild ruminants. Bartonella
infection is common in many species of small, wild rodents, and has also been reported in marine
mammals, bats, and other species. There are reports indicating transmission of Bartonella
infection from rodents to human beings (Iralu et al., 2006; Kosoy et al., 2003; Welch et al.,

440 1999).

441 Bartonella bovis bacteremia appears to be common in dairy and beef cattle, with prevalences of greater than 50 percent in some dairy herds and beef herds (Breitschwerdt et al., 442 443 2001; Cherry et al., 2009; Maillard et al., 2006; Martini et al., 2008). As discussed for cats, it 444 may be difficult to attribute clinical conditions in cattle to Bartonella infection due to the high 445 prevalence of *Bartonella* bacteremia in this species. Endocarditis attributable to *B. bovis* was 446 reported in two French cows (Maillard et al., 2007). Results of another study indicated that 447 cattle, a reservoir host for *B. bovis*, rarely show clinical signs of infection. Interestingly, cattle 448 with *B. bovis* bacteremia had a statistically significantly shorter interval from calving to 449 breeding, and fewer retained placentas, than did cattle that were not bacteremic (Maillard et al., 450 2006). Other Bartonella species detected in healthy cattle include B. chomelii and B. henselae 451 (Cherry et al., 2009; Maillard et al., 2004a). Bartonellas have been detected in biting flies, and 452 these arthropods may be vectors for transmission of *Bartonella* species among cattle (Chung et 453 al., 2004; Halos et al., 2004).

454 A recently described *Bartonella*, *candidatus* B. melophagi, was isolated from commercial 455 sheep blood and has been isolated from sheep keds also. It is likely that domestic sheep are

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456 reservoir hosts for this *Bartonella* species, and that a primary vector is the sheep ked. To date, 457 there are no reports of clinical disease in sheep associated with *Bartonella* species, and no 458 information regarding prevalence of *Bartonella* infections in sheep (Bemis and Kania, 2007). 459 *Candidatus* B. melophagi was recently isolated from the blood of two ill human beings, 460 however, the clinical importance of this organism in the illnesses of those individuals has not 461 been established (Maggi et al., 2009). Bartonella are also reported from wild elk and deer 462 (Bermond et al., 2002; Chang et al., 2000b; Dehio et al., 2001; Maillard et al., 2004b). 463 Bartonella henselae has been reported in 3 horses, one with vasculitis, one with chronic 464 arthropathy, and one aborted fetus (Johnson et al., 2009; Jones et al., 2008). Whether B. henselae or other Bartonella species have a role as pathogens in horses or sheep remains to be 465 466 determined.

It was recently suggested that *Bartonella* may be a cause of endocarditis in sea otters
(Chomel et al., 2009c). *Bartonella* species have also been detected in whales and porpoises
(Harms et al., 2008; Maggi et al., 2005b).

470 PUBLIC HEALTH

471 Several *Bartonella* species or subspecies are considered zoonotic or possibly zoonotic,
472 including but not limited to, *B. henselae*, *B. clarridgeiae*, *B. koehlerae*, *B. vinsonii* subsp
473 *berkhoffii*, *B. vinsonii* subsp. *arupensis*, *B. grahamii*, *B. elizabethae*, *B. tamiae*, *B. alsatica*,
474 *candidatus* B. washoensis, and *candidatus* B. rochalimae (Breitschwerdt and Chomel, 2005;
475 Chomel et al., 2009c; Lin et al., 2008).
476 *Bartonella* spp. cause a wide variety of clinical syndromes in people, for example, CSD

477 (typical and atypical forms, including encephalopathies in children); bacillary angiomatosis and

478 peliosis; relapsing fever with bacteremia; endocarditis; optic neuritis; pulmonary, hepatic, and

479 splenic granulomas; osteomyelitis and many others (Florin et al., 2008; Massei et al., 2005; 480 Massei et al., 2000). Immunocompetent individuals may have more localized infections, 481 whereas infections in immunocompromised individuals may more often be systemic and can be 482 fatal. As previously noted, there is more information available about zoonotic *B. henselae* 483 infections than for other zoonotic Bartonella infections. In one study, a high prevalence of 484 antibodies to B. henselae was found among Italian children without evidence of obvious illness 485 (Massei et al., 2003). This suggests that exposure and asymptomatic infection may have 486 occurred early in life with spontaneous resolution, although the possibility of long-term 487 asymptomatic infection must be considered. 488 Bartonella henselae isolates from people and their animal contacts have been shown to 489 be related, thus indicating the animals as sources of human infection (Bouchouicha et al., 2009b; 490 Breitschwerdt et al., 2007; Chang et al., 2002a). Different B. henselae types may induce varying 491 pathologic features in infected people, and certain types have been proposed to be more likely 492 associated with human disease than others (Arvand et al., 2007; Bouchouicha et al., 2009b; 493 Woestyn et al., 2004). Some studies do not support these conclusions, however, and further 494 work is required to discern what Bartonella henselae types are associated, for example, with 495 human disease or feline colonization (Lindroos et al., 2006). 496 Cats are the primary reservoir and vector for transmission of *B. henselae* and probably 497 also B. clarridgeiae to human beings. The role of arthropods in direct transmission of Bartonella 498 to human beings is not certain. Transmission of *B. henselae* from cats to human beings is 499 believed to occur most commonly through contamination of scratches with flea excrement 500 (Chomel et al, 1996). Transmission may also occur through cat bites if cat blood or flea

501 excrement contaminate the bite site (Chomel et al., 1996; Finkelstein et al., 2002). It has been

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502 suggested that B. henselae and B. vinsonii subsp. berkhoffii may be transmitted by dog bites or 503 scratches, and *Bartonella* species have been detected by PCR in oral swabs collected from dogs 504 (Duncan et al., 2007a; Keret et al., 1998; Rolain et al., 2009). Bartonella vinsonii subspecies 505 arupensis, B. elizabethae, B. grahamii, and B. washoensis are believed to be transmitted via 506 contact with rodents (Iralu et al., 2006; Kosoy et al., 2003). The role of animals in the 507 epidemiology of *B. quintana* infections is not clear. 508 Pet owners should be informed of the current understanding regarding how cats and dogs 509 acquire *Bartonella* infections, and how these infections may be transmitted to people, including 510 the association of flea infestation or ticks with transmission. Given the number of other animal 511 species in which *B. henselae* has been detected, other means of transmission (including other

arthropod vectors) should be considered, and caution exercised when handling animals of anyspecies.

514 Common sense precautions for avoiding transmission of *Bartonella* spp. from animals to human beings include excellent flea and tick control, avoiding interactions that result in scratches 515 516 or bites, thoroughly washing bite or scratch wounds, seeking medical attention when necessary, 517 and acquiring new pets of known good health status that are and have been ectoparasite free. 518 Stray or impounded cats less than 1 year old are most likely to be bacteremic with B. henselae 519 (Chomel et al., 1995). The United States Centers for Disease Control, the National Institutes of 520 Health, and the Infectious Diseases Society of America Guidelines for Prevention and Treatment 521 of Opportunistic Infections in HIV-Infected Adults and Adolescents, and HIV-exposed and HIV-522 Infected Children recommend the following when acquiring a new cat: adopt a cat older than 1 523 year of age that is in good health and flea-free, avoid rough play with cats, maintain flea control, 524 wash any cat-associated wounds promptly, and do not allow a cat to lick wounds or cuts. The

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- 525 Guidelines note that there is no evidence to indicate any benefit to cats or their owners from
- 526 routine culture or serologic testing of cats for *Bartonella* infection (Kaplan et al., 2009;
- 527 Mofenson et al., 2009). There are not specific recommendations in that document regarding
- 528 Bartonella and other companion animal species.

529 SUMMARY

- 530 A great deal has been learned about *Bartonella* infections recently, and much more discovery lies
- ahead. A large proportion of domestic and non-domestic animals harbor persistent *Bartonella*
- 532 spp. infections. Ongoing and future studies will help to ascertain what the possible ramifications
- 533 of chronic *Bartonella* bacteremia may be for animal health. Veterinarians should be aware of the
- 534 high prevalence of feline *Bartonella* bacteremia, and carefully consider their approach to
- 535 evaluating cats for possible Bartonella-related clinical conditions. Based on current knowledge,
- healthy cats and dogs should not be routinely tested for *Bartonella*, and antibiotic treatment for
- 537 Bartonella should not be routinely recommended for healthy cats and dogs. When ill animals are
- treated, treatment should always be accompanied by comprehensive vector control measures, and
- 539 client education. Client education should emphasize the role of arthropod vectors in *Bartonella*
- 540 transmission, other potential avenues for transmission, and also address the potential risks
- 541 associated with antibiotic treatment, the likelihood of re-infection of an animal with *Bartonella*
- 542 following antibiotic treatment, and the zoonotic potential of *Bartonella*.
- 543 **Conflict of Interest**: None
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