Characterization of Clinical Isolates of Bartonella henselae Strains, South Korea

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Bartonella henselae, a gram-negative bacterium, is a common causative agent of zoonotic infections. We report 5 cultureproven cases of *B. henselae* infection in South Korea. By alignment of the 16S rRNA sequences and multilocus sequencing typing analysis, we identified all isolates as *B. henselae* Houston-1 strain, which belongs to sequence type 1.

The genus *Bartonella* includes infectious, gram-negative, facultative intracellular bacteria of numerous species. Among the *Bartonella* species, *B. henselae* is known as one of the most noteworthy pathogens (1). *B. henselae* causes cat-scratch disease, which is a common zoonosis and manifests various clinical symptoms (2).

A case of *B. henselae* infection in South Korea was confirmed in 2005 by PCR (3). Although a few more studies have been published after this case of *B. henselae*, only 2 cases were culture-proven: 1 from blood and 1 from bone marrow (4,5). Because of difficulties in cultivation and isolation, studies of the isolation of *B. henselae* from clinical specimens remain scarce. In this study, we analyzed the characteristics of the isolated *B. henselae* strains in South Korea and compared the clinical features of the patients.

The Study

We conducted the study among patients who visited Inha University Hospital, a tertiary hospital in Incheon, South Korea, during 2009–2016. From these patients, we isolated 5 cases in which *B. henselae* was identified from cultures of blood or bone marrow (Table 1).

Case-patient 1 (IIBC1301) was a 22-year-old man hospitalized for left inguinal lymphadenopathy that had started 10 days earlier. His body temperature was 38.5° C, and he had rashes that started on the palms and soles and subsequently spread to his entire body. *B. henselae* was isolated from the blood that was cultured on the second day of hospitalization.

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Case-patient 2 (IIBC1302) was a 40-year-old woman hospitalized for fever and myalgia, symptoms that had lasted for 1 month. The patient had an erythematous papular rash on her face and extremities and tenderness in her abdomen. Computed tomography (CT) of the abdomen showed chronic cholecystitis; therefore, levofloxacin and metronidazole were prescribed (online Technical Appendix Figure, panel A, https://wwwnc.cdc.gov/EID/article/24/5/17-1497-Techapp1.pdf). B. henselae was identified from cultures of blood obtained on the first day of the hospitalization. The patient had not raised any animals. After discharge, the patient experienced continuous fever, poor oral intake, and weight loss. Reevaluation showed centrilobular ground-glass opacity in both lung fields on chest CT and growth of Mycobacterium tuberculosis on sputum acid-fast bacilli culture (online Technical Appendix Figure, panel B). A pulmonary tuberculosis infection was diagnosed and treated with antituberculosis medication.

Case-patient 3 (IIBC1303) was a 52-year-old woman hospitalized for fever and left flank pain; her symptoms had persisted for 1 month. She also reported right-side neck swelling and pain at neck levels II, III, and VA. *B. henselae* was isolated from cultures of blood collected on the 16th day of hospitalization. She had no contact with animals.

Case-patient 4 (IIBC1304) was a 42-year-old man we previously reported (5) whose main complaints were fever, rash, and arthralgia. *B. henselae* was isolated from a bone marrow sample. The patient had no contact with or experience in raising pets.

Case-patient 5 (IIBC1305), also previously published (4), was a 73-year-old woman who had *B. henselae* isolated from her blood. She also did not have any contact with animals.

Bartonella species can be grown by blood agar–based culture systems. However, it is difficult to culture them this way because the growth of bacterial cells is slow, and obtaining colonies on the agar plate takes a long time. On the other hand, *Bartonella* species grow more rapidly with cell culture–based systems (6). For testing of these patients, we grew ECV304 cells in M199 media containing 10% heat-inactivated fetal bovine serum and inoculated 1 mL of whole blood or other samples from the patients onto the cells. After 24 hours, we washed the cells with Dulbecco's phosphate-buffered saline and maintained them in M199 media. We performed an immunofluorescence assay

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Bartonella henselae, Soul	li Kulea									
Characteristic	Case-patient 1	Case-patient 2	Case-patient 3	Case-patient 4 (5)	Case-patient 5 (4)					
Age, y/sex	22/M	40/F	52/F	42/M	73/F					
Clinical symptoms	Inguinal LAP, rash	Fever, myalgia	Febrile sense, left	Rash, fever,	Fever, general					
			flank pain	myalgia	weakness					
Lymphadenopathy	External iliac chain,	Left neck level IV, V	Right neck II, III, VA	Right	None					
	inguinal area,			supraclavicular						
	supraclavicular area			area						
Leukocytes, cells/µL	10,920	6,130	8,180	19,260	5,120					
AST/ALT, IU/dL	132/270	107/51	30/16	212/246	47/56					
ESR, mm/h/CRP, mg/dL	21/3.93	44/5.5	4/0.14	25/12.9	22/13.16					
Treatment	Third-generation	Levofloxacin,	Third-generation	Doxycycline,	Third-generation					
	cephalosporin,	metronidazole, third-	cephalosporin,	changed to	cephalosporin,					
	doxycycline	generation	minocycline,	minocycline	doxycycline					
		cephalosporin and	metronidazole	-						
		doxycycline								
B. henselae IgG titer	1:160	1:640	1:160	1:1,280	1:160					
Pets	None	None	None	None	None					
Co-occurring conditions	None	Pulmonary	None	None	None					
		tuberculosis								
*ALT, alanine aminotransferase; AST, aspartate aminotransferase; CRP, C-reactive protein; ESR, erythrocyte sedimentation rate.										

Table 1. Demographic and clinical characteristics of 5 case-patients whose serum sample cultures revealed the presence of Bartonella henselae, South Korea*

(IFA) with the patient's own serum (1:40 diluted) every week after the inoculation. When the growth of bacteria was observed, we scraped all cultured cells from the T25 flask. We then reinoculated 1 mL of infected ECV304 cells onto uninfected ECV304 cells in a T75 flask for expansion of bacterial cells.



Figure. Phylogenetic tree of 5 *Bartonella henselae* clinical isolates from patients in South Korea (black dots) and closely related species based on 16S rRNA gene sequences. Database accession numbers are provided for reference sequences. Scale bar indicates nucleotide substitutions per site.

		Allele at the 8 loci								
Isolate	Specimen type	16S	batR	ftsZ	gltA	groEL	nlpD	ribC	rpoB	Sequence type
IIBC1301	Blood	1	1	1	1	1	1	1	1	1
IIBC1302	Blood	1	1	1	1	1	1	1	1	1
IIBC1303	Blood	1	1	1	1	1	1	1	1	1
IIBC1304	Bone marrow	1	1	1	1	1	1	1	1	1
IIBC1305	Blood	1	1	1	1	1	1	1	1	1

Table 2. Characteristics of clinical Bartonella henselae isolates from 5 case-patients, South Korea*

To identify the bacterial isolates, we amplified and sequenced the 16S rRNA gene (7). The pathogens cultured from the specimens showed the highest sequence similarities with *B. henselae* Houston-1 strain (GenBank accession nos. KY773227, KY773228, KY773229, KY773290, and KY885188). The similarity was >99% (Figure) (8,9). IFA results using a commercial *Bartonella* IFA IgG kit (FOCUS Diagnostics, DiaSorin Molecular, Cypress, CA, USA) also showed positive results for all patients' serum samples; titers ranged from 1:40 to 1:1,280 (Table 1). We also performed multilocus sequence typing to determine the genotypes of *B. henselae* isolates (*10*) and found that all isolates belonged to sequence type 1 (Table 2).

Conclusions

We cultured B. henselae isolates from clinical samples and compared characteristics of 5 patients: 3 new cases and 2 previously reported cases from which B. henselae was isolated (Table 2). Because of the diverse manifestations of B. henselae infection, the symptoms were similar to those of other bacterial infections. B. henselae infections in 3 patients were initially misdiagnosed as other diseases: sexually transmitted disease (case-patient 1), enteric fever-like syndrome (case-patient 2), and acute pyelonephritis (casepatient 3). The diagnosis of B. henselae infection was made even more difficult because none of these 5 patients reported a history of raising cats. However, the absence of contact with animals should not preclude infection; even though B. henselae infection is usually related to cat scratches or bites, it may also occur without animal contacts (5). It is also noteworthy that the patient described in case 2 was coinfected with pulmonary tuberculosis. Co-infection with B. henselae and Mycoplasma spp. has also been reported in previous studies (11,12). Co-infection with other bacteria suggests that infection with Bartonella species may weaken the host's immune system, leaving the host vulnerable to secondary infections. In addition, these co-infections may cause difficulty in diagnosing Bartonella infection.

Multilocus sequence typing indicated that all isolates from this study belonged to *B. henselae* sequence type 1. This result is consistent with previous studies, which showed relatively less diversity among human strains than among the feline reservoir (10,13).

In summary, the clinical features of *B. henselae* infection are diverse and nonspecific, which could initially lead to misdiagnosis as other diseases. Physicians and patients

should consider that *Bartonella* infection presents various clinical symptoms and might be a common cause of fever of unknown origin, irrespective of exposure to cats. Once *Bartonella* infection is suspected, cell culture should be considered to confirm the diagnosis.

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Dr. Kwon is a medical doctor at Inha University Hospital in Incheon, South Korea. Her research interests include infectious diseases, especially focusing on intracellular bacteria and epidemiology. Dr. Park is a research fellow at Inha University in Incheon. Her primary research interests include antimicrobial agents and vaccines for treatment of infectious diseases.

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Bartonella henselae

[bär'' tə-nel'ə henz' ə-lā]

Bartonella is a genus of gram-negative bacteria named after Peruvian scientist Alberto Leonardo Barton. He identified a unique bacterium in 1905 during an outbreak among workers building a railway between Lima and La Oroya, a mining town in the Andes. The illness, usually fatal, was characterized by fever and severe anemia. Many of the sick were brought to Guadalupe Hospital in Lima, where Dr. Barton isolated the etiologic agent (which had been transmitted by sandflies) in patients' blood cells. It was later called *Bartonella bacilliformis*.

The species *B. henselae* was named after Diane Hensel, a technologist in the clinical microbiology laboratory, University Hospitals, Oklahoma City, who in 1985 observed a *Campylobacter*-like organism in blood cultures of HIV-infected patients. The organism was first named *Rochalimaea henselae* and then *B. henselae*, when sequencing showed identity with that genus.

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