Case Report: Polymerase Chain Reaction Testing of Tick Bite Site Samples for the Diagnosis of Human Granulocytic Anaplasmosis

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Abstract. Human granulocytic anaplasmosis (HGA) is a tick-borne infectious disease caused by Anaplasma phagocytophilum, an obligate intracellular bacterium. Until now, the utility of tick-bite site samples for HGA diagnosis has not been reported. Using a patient's buffy coat and tick-bite site crust samples, we performed polymerase chain reaction (PCR) testing using *Ehrlichia*- or *Anaplasma*-specific primers. PCR with buffy coat and crust samples obtained before doxycycline administration was positive. Six days after doxycycline administration, PCR with the buffy coat sample was negative but PCR with a crust tissue sample from the tick-bite site remained positive. This is the first case to suggest that crust tissue at the tick-bite site may be useful for early HGA diagnosis in patients who have already been treated with antibiotics such as doxycycline.

INTRODUCTION

Human granulocytic anaplasmosis (HGA) is a tick-borne infectious disease caused by *Anaplasma phagocytophilum*, an obligate intracellular bacterium that was first identified as a human pathogen in 1994.¹ In Korea, the first case of HGA was reported in 2013 as an emerging infectious disease.² HGA is mainly transmitted by ticks of the *lxodes* species, which are common vectors of Lyme disease and babesiosis.³ As patients with HGA develop nonspecific symptoms such as fever, malaise, headache, and chills, the diagnosis may be challenging. As such, differential diagnoses typically include rickettsial diseases, infectious mononucleosis, Lyme disease, and West Nile virus infection.⁴

Isolation of *A. phagocytophilum* in culture is difficult and usually takes longer than 30 days.⁵ Therefore, the use of polymerase chain reaction (PCR) testing for the rapid diagnosis of HGA is important. Herein, we aimed to evaluate the usefulness of a crust tissue at the tick-bite site for the diagnosis of HGA via PCR testing.

CASE REPORT

A 79-year-old man was admitted for fever, which started 6 days prior to his admission. Four days prior to admission, the patient experienced myalgia accompanied by chills. The patient was monitored for continuous symptoms and eventually admitted to hospital due to worsening of his condition. He was a farmer who had worked in rice and pepper fields, and, aside from right ureter stones, he had no other medical conditions.

The physical examination conducted on admission revealed that the patient had a fever of 37.8°C, a pulse rate of

102 beats/minute, and a respiratory rate of 22 breaths/minute. On admission, he had hyperemic conjunctiva and a lesion that resembled a tick-bite site was found on his left flank area (Figure 1).

The patient's white blood cell count was 3,310/mm³ (91.9% polymorphonuclear cells and 2.2% lymphocytes), his hemoglobin level was 13.2 g/dL (reference values: 11-17 g/dL), and his platelet count was 24,000/mm³ (reference values: $150,000-400,000/\mu$ L). Other laboratory values were as follows: aspartate aminotransferase level, 93.5 U/L (reference values: 0-35 U/L); alanine aminotransferase level, 36.3 U/L (reference values: 0-35 U/L); lactate dehydrogenase level, 619 U/L (reference values: 100-190 U/L); and creatine phosphokinase level, 309 U/L (reference values: 0-150 U/L).

For a febrile illness, 2 g ceftriaxone was administered at admission. On the second day of admission (seventh day since the onset of symptoms), a crust lesion and blood samples were collected and sent to the laboratory for testing (Table 1); we collected the tick-bite site crust sample by simply detaching the crust with a scalpel (no. 11 blade) and tweezers.

HGA was suspected, but morulae were not observed on peripheral blood smear. PCR was performed with genomic DNA from the patient's buffy coat and crust samples. Conventional PCR or nested PCR were performed using *Ehrlichia*-or *Anaplasma*-specific primers targeting the GroEL heat-shock protein operon gene (*groEL*), 16S ribosomal RNA gene (16S rRNA), ankyrin-repeat protein AnkA gene (*ankA*), and major surface protein 2 (*msp2*) gene.^{6–9} The primer sequences and PCR conditions used are summarized in Table 2. The amplified PCR products were cloned into a PCR 2.1 vector from the TA cloning kit (Invitrogen, Carlsbad, CA) and positive recombinant plasmids were sequenced. The sequencing result was checked using BlastN from the National Center for Biotechnology Information (Bethesda MD) to identify the bacterium.

A serological test (IgG and IgM) for the detection of *A. phagocytophilum* was performed using an indirect immunofluorescent antibody assay (IFA) from a commercial kit (Fuller Laboratories, Fullerton, CA) according to the

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FIGURE 1. A crust lesion suspected to be a tick-bite site on the left flank area of a 79-year-old man. This figure appears in color at www.ajtmh.org.

manufacturer's protocol.¹⁰ The serological positive cutoff value of the kit is 1:80 for the IgG titer and 1:16 for the IgM titer.

The PCR to detect *A. phagocytophilum* and sequencing were performed using the buffy coat and crust lesion samples obtained 7 days from symptom onset (Table 1). All of the sequencing results confirmed the presence of *A. phagocytophilum*.

Twelve days from the onset of symptoms, ceftriaxone therapy was switched to 100 mg doxycycline BID administration. A nested PCR for *groEL* was performed using the buffy coat and crust lesion specimens at each time point after the symptoms had begun (Table 1). On day 6 of doxycycline administration, which was 18 days after the onset of symptoms, PCR for the buffy coat samples was negative. However, PCR for the crust lesion sample remained positive (Supplemental Figure 1). On admission, using the buffy coat and crust lesion samples, PCR for 16S rRNA and *groEL* genes were performed, and their respective 926

base pair (bp) and 362 bp amplicons were cloned. The sequencing results showed 2 bp differences for 16S rRNA and 1 bp difference for *groEL*.^{2,11} Phylogenetic trees were constructed based on sequences of 16S rRNA and *groEL* genes from GenBank using the neighbor-joining method on the ClustalX software program (Dublin, Ireland; Figure 2). In a homology test, *A. phagocytophilum* was 99.9% homologous to the *A. phagocytophilum* strain HZ detected in the United States. It was also 99.9% homologous to the *A. phagocytophilum* strain confirmed in Gwangwon, Korea (accession no. KF805344), as well as to the Russian strain (accession no. HM366590) and Chinese strain (accession no. JN990105).

Seven days from the initiation of symptoms, the IFA results were negative for both IgM and IgG, whereas 10 days from the initiation of symptoms, the IgG titer increased to 1:80. However, IgM remained negative even at 108 days from the onset of symptoms.

Doxycycline administration continued for 10 days and the patient was discharged without complications.

DISCUSSION

The laboratory diagnosis of HGA is based on Anaplasma culture, a PCR test, or antibody titer increase using specimens obtained from the acute phase of the disease and during recovery.¹² However, serological tests are challenging and may take several weeks to detect a 4-fold increase in antibodies, which is required to confirm the diagnosis. In this case, which is the second reported HGA case from Korea, it took longer than a week to detect the antibodies. Even afterward, a relatively low IgG titer was observed and IgM was not detected on day 108 from symptom onset. In some cases, delays in the administration of effective antibiotics may cause fatal complications.^{13,14} In addition, it is technically challenging and lengthy process to isolate Anaplasma in culture. For these reasons, HGA may be difficult to diagnose without molecular technology. Therefore, a rapid and accurate diagnosis is necessary to ensure appropriate treatment. PCR is a useful tool for diagnosing infectious diseases caused by microbes that are difficult to culture or slow growing.¹⁵ For the diagnosis of rickettsial diseases, the usefulness of PCR testing using a buffy coat sample is well known.¹⁶ In addition, PCR conducted with

Diagnostic test results of a 79-year-old patient with human granulocytic anaplasmosis						
Collection date	Time from symptom onset to specimen collection (days)	IFA	Specimen used	groEL nested PCR result		
April 24, 2015	7	IgM (–)	Buffy coat	Positive		
		lgG (–)	Crust	Positive		
April 27, 2015	10	IgM (–)	Buffy coat	Positive		
•		IgG 1:80				
April 29, 2015	12 (doxycycline start)	IgM (–)	Buffy coat	Positive		
•		IgG 1:80	-			
May 5, 2015	18	IgM (–)	Buffy coat	Negative		
		IgG 1:80	Crust	Positive		
May 20, 2015	33	IgM (–)	Buffy coat	Negative		
•		lgG 1:80	Crust	Negative		
June 22, 2015	66	IgM (–)	Buffy coat	Negative		
		IgG 1:320	-	0		
August 3, 2015	108	IgM (–)	Buffy coat	Negative		
0		IgG 1:160	,	5		

TABLE 1 Diagnostic test results of a 79-year-old patient with human granulocytic anaplasmosis

groEL = GroEL heat-shock protein operon gene; IFA = immunofluorescent antibody assay; PCR = polymerase chain reaction.

Target gene*	Primer name (sequences)	Annealing temperature (°C)	Product size (base pairs)	References
groEL nested PCR for Anaplasmataceae	GR0607F (5'-GAAGATGCWGTWGGWTGTACKGC-3')	57	688	6
(external primers)	GR01294R (5'-AGMGCTTCWCCTTCWACRTCYTC-3')		445	0
groEL nested PCR for Anaplasmataceae (internal primers)	GR0677F (5'-ATTACTCAGAGTGCTTCTCARTG-3') GR01121R (5'-TGCATACCRTCAGTYTTTCAAC-3')	57	445	6
16S rRNA nested PCR for Anaplasma and	AE1-F (5'-AAGCTTAACACATGCAAGTCGAA-3')	59	1.406	7
<i>Ehrlichia</i> species (external primers)	AE1-R (5'-AGTCACTGA CCCAACCTTAAATG-3')	00	1,100	•
16S rRNA nested PCR for E. chaffeensis	EC-F (5'-CAATTGCTTATAACCTTTTGGTTATAAAT-3')	56	390	7
(internal primers)	EC-R (5'-TATAGGTACCGTCATTATCTTCCCTAT-3')			
16S rRNA nested PCR for A.	AP-F (5'-GTCGAACGGATTATTCTTTATAGCTTGC-3')	56	926	7
phagocytophilum (internal primers)	AP-R (5'-CCCTTCCGTTAAGAAGGATCTAATCTCC-3')			
ankA nested PCR for A. phagocytophilum	ANK-F1 (5'-GAAGAAATTACAACTCCTGAAG-3')	53	705	8
(external primers)	ANK-R1 (5'-CAGCCAGATGCAGTAACGTG-3')			_
ankA nested PCR for A. phagocytophilum	ANK-F2 (5'-TTGACCGCTGAAGCACTAAC-3')	54	664	8
(internal primers)	ANK-R2 (5'-ACCATTTGCTTCTTGAGGAG-3')			
msp2 conventional PCR for A.	MSP3F (5'-CCA GCG TTT AGC AAG ATA AGA G-3')	54	334	9
phagocytophilum	MSP3R (5'-GCC CAG TAA CAT CAT AAG C-3')			

TABLE 2 PCR conditions and oligonucleotide primers used in this study

* ankA = ankyrin-repeat protein gene; groEL = GroEL heat-shock protein operon gene; msp2 = major surface protein 2 gene; 16S rRNA = 16S ribosomal RNA gene.

eschar skin biopsy specimens detected *Rickettsia rick-ettsii*, *Rickettsia parkeri*, and *Rickettsia akari* in patients with rickettsial diseases such as scrub typhus or spotted fever.¹⁷ However, *Anaplasma* does not form typical eschars on the tick-bite site, and as such, a diagnosis using PCR testing of a tick-bite site crust sample has not been reported.

In a study that examined a tick-feeding site in sheep that were infected by human *A. phagocytophilum* isolates, biopsy of the feeding site showed signs of necrosis, and higher numbers of eosinophils and neutrophils were observed. *Anaplasma phagocytophilum*–infected granulocytes were also observed in the skin biopsy immunohistochemistry study.¹⁸ In another study, skin biopsies were collected from tick feeding lesions on 38 grazing lambs. PCR analysis of blood samples showed 33 lambs (86.8%) were positive for *A. phagocytophilum*. However, 37 lambs (97.4%) had one or more skin biopsies and 70 of 76 biopsies from tick attachment sites (92.1%) were PCR positive.¹⁹

Therefore, like eschars from the scrub typhus or spotted fever groups, in patients with HGA, tick-bite sites were hypothesized to have a large inoculum around the site in which the bacteria had multiplied and spread.^{20,21}

From both blood and crust samples obtained before doxycycline administration, the *Anaplasma* PCR tests were positive. Six days after administering doxycycline, PCR for the buffy coat sample was negative; however, PCR for the tick-bite site crust sample remained positive. Thus, bacterial DNA from the crust appears to last longer than that from the blood. The positive PCR result of *A. phagocytophilum* in a crust after the onset of treatment with doxycycline might be the result of high inoculum or theoretically less than optimal diffusion (access) of doxycycline into the crust tissue. We did not perform immunohistochemical staining of crust tissue. Further study is required.

In conclusion, this study argues for the usefulness of PCR testing using a tick-bite site crust sample, similar to

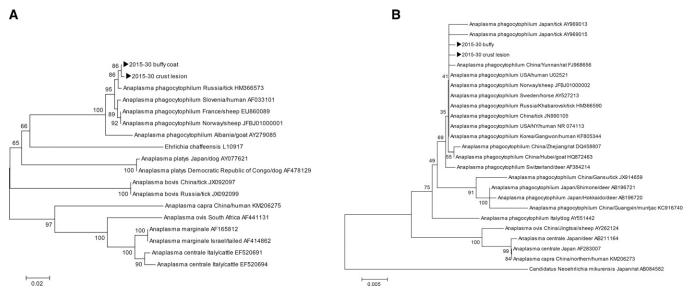


FIGURE 2. A phylogenetic tree based on (A) groEL (362 base pair [bp]) and (B) 16S rRNA (810 bp) sequences from GenBank and buffy coat and crust lesion specimens collected in this study (►). Scale bars indicate (A) 0.02 and (B) 0.005 base substitutions per site.

the use of the buffy coat sample, in the early and acute diagnosis of HGA. This study is the first to suggest that the crust tissue at the tick-bite site may be a useful specimen for the diagnosis of HGA, even in patients who received antibiotics such as doxycycline, which are active against HGA. More samples are required to verify whether PCR positivity on crust tissue at the tick-bite site is useful for diagnosing the disease even after antibiotic administration.

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