Transmission of Bordetella holmesii during Pertussis Outbreak, Japan


We describe the epidemiology of a pertussis outbreak in Japan in 2010–2011 and Bordetella holmesii transmission. Six patients were infected; 4 patients were students and a teacher at the same junior high school. Epidemiologic links were found between 5 patients. B. holmesii may have been transmitted from person to person.

Bordetella holmesii, a small gram-negative coccoid bacillus that was first reported in 1995 (1), was originally identified as a member of the Centers for Disease Control and Prevention nonoxidizer group 2. The organism is associated with bacteremia, endocarditis, and pneumonia, usually in patients with underlying disorders such as asplenia or sickle cell anemia, and has been isolated from blood and sputum samples (1–5). B. holmesii may also be responsible for causing symptoms similar to those of pertussis (whooping cough) in otherwise healthy patients (6). Large surveillance studies in the United States and Canada have shown that the organism was detected in nasopharyngeal swab (NPS) specimens of patients with pertussis-like symptoms (7,8). Although humans may be infected with B. holmesii, transmission of B. holmesii between humans has not yet been fully elucidated.

Pertussis is a highly contagious disease caused by the bacterium B. pertussis. The organism is known to be transmitted from person to person by airborne droplets (9). During a recent pertussis outbreak, we conducted a laboratory-based active surveillance study and detected 76 suspected cases of pertussis. Among these cases, we identified not only B. pertussis infection but also B. holmesii infection. The purpose of this study was to describe the epidemiology of the pertussis outbreak and to determine the epidemiologic relatedness of B. holmesii transmission.

The Study

During 2010–2011, a pertussis outbreak occurred in a suburban town (town A) in Nobeoka City in Miyazaki Prefecture, Japan. Town A has a population of 4,227 persons and an elementary school and junior high school. The first pertussis case (in a 17-year-old boy) was reported in September 2010. From that time, we conducted a laboratory-based active surveillance study in the area until April 2011, in cooperation with clinics, hospitals, and local health departments. Pertussis-suspected cases were defined as an illness with cough lasting for ≥2 weeks, and pertussis-confirmed cases were defined as the presence of 1 of the following laboratory findings: a culture-positive result for Bordetella species from NPS specimens, or a positive result for molecular testing for Bordetella species.

For molecular testing, we conducted conventional PCR specific for insertion sequence IS481, which is known to detect B. pertussis and B. holmesii, and B. pertussis–specific loop-mediated isothermal amplification assays (10–12). To further confirm B. holmesii infection, B. holmesii–specific real-time PCR specific for the recA gene was also performed as described (8). For confirmed cases of B. holmesii infection, we collected the general information for patients (clinical symptoms, treatment, contact information, and outcome) by face-to-face interview or questionnaire.

During the pertussis outbreak, we identified 76 suspected pertussis cases. Among these suspected cases, 35 cases were confirmed by laboratory testing and involved persons 2–63 years of age (median age 13 years); 1 case occurred in a 2-year-old child, 14 in 6- to 12-year-old children, 14 in 13–15-year-old children, and 6 in persons >16 years of age. Despite pertussis vaccination rates in childhood of 82.3%–92.6%, most (80%) patients were students 6–12 and 13–15 years of age. Among the 35 confirmed cases, 29 and 6 patients showed B. pertussis and B. holmesii infection, respectively. Confirmed cases of B. holmesii infection were observed within the last half of the epidemic curve, i.e., weeks 1–12 of 2011 (Figure 1). There were no cases of co-infection with B. pertussis and B. holmesii.

All NPS specimens from patients with B. holmesii infection showed a negative result for the B. pertussis loop-mediated isothermal amplification, but showed positive results in the IS481 PCR and B. holmesii recA real-time PCR (Table 1). B. holmesii-like organisms were obtained...
Transmission of *Bordetella holmesii* from 5 patients, and these were identified as *B. holmesii* by *recA* gene sequencing. Patient 6 had been treated with antimicrobial drugs before the culture test, which probably resulted in a culture-negative test result in this patient. Real-time PCR confirmed that patient 6 had a low *B. holmesii* DNA load (threshold cycle 36.6) in her NPS specimen. All patients experienced paroxysms of coughing without posttussive vomiting, especially at night, and 3 also experienced a whoop (Table 2). Five of 6 patients had a persistent cough lasting >2 weeks. No patients experienced any complications, and they were treated mainly with azithromycin, resulting in complete recovery. None of the patients were immunocompromised.

Our surveillance study showed epidemiologic linkage between 5 patients (patients 2–6), but not for patient 1 (Figure 2). Patient 1, a student in high school A, had the first case of *B. holmesii* infection (probable index case). He lived in a dormitory outside Nobeoka City. However, his family home was in town A in Nobeoka City, and he returned to his home at the end of 2010 and remained there in 2011. Patients 2, 3, and 4 were students at the same junior high school (B) in town A and were close friends. Patient 5 was a teacher at junior high school B and was in charge of patient 4. Patient 6 was a medical practitioner at clinic C, which is 1 of 2 clinics in town A. Patients 1–5 visited clinic C in early January, late February, late February, early March, and late March, 2011, respectively. The duration of illness in patient 1 did not overlap with that of the other patients, whereas that of patients 2–6 clearly overlapped.

Conclusions

In the past 16 years, ≈70 *B. holmesii* clinical strains have been isolated from human patients in several countries, mainly the United States. All reported cases of *B. holmesii* infection have been sporadic occurrences. Thus, the reservoir of *B. holmesii* is currently unknown. Moreover, whether *B. holmesii* is transmitted between humans is not known. In this report, we have demonstrated that 5 patients infected with *B. holmesii* showed epidemiologic linkage. In particular, the fact that 4 of these patients attended the same junior high school suggests that *B. holmesii* may be transmitted from person to person.

From Figure 1, it is evident that the numbers of suspected and laboratory-confirmed cases of pertussis were higher during the period September 2010–April 2011, compared to the previous years. The peak of cases occurred during the early spring, with the highest numbers of cases seen in late February and early March.

Table 1. Characteristics of *Bordetella holmesii* infection in 6 patients during pertussis outbreak, Japan, September 2010–April 2011*

<table>
<thead>
<tr>
<th>Patient no.</th>
<th>Age, y/sex</th>
<th>Duration of cough, d†</th>
<th><em>B. holmesii</em> test results</th>
</tr>
</thead>
<tbody>
<tr>
<td>1</td>
<td>17/M</td>
<td>5</td>
<td>+ (28.7)</td>
</tr>
<tr>
<td>2</td>
<td>15/F</td>
<td>4</td>
<td>+ (23.4)</td>
</tr>
<tr>
<td>3</td>
<td>15/F</td>
<td>&gt;14</td>
<td>+ (21.6)</td>
</tr>
<tr>
<td>4</td>
<td>14/F</td>
<td>8</td>
<td>+ (25.1)</td>
</tr>
<tr>
<td>5</td>
<td>40/M</td>
<td>8</td>
<td>+ (27.0)</td>
</tr>
<tr>
<td>6</td>
<td>45/F</td>
<td>15</td>
<td>+ (36.6)</td>
</tr>
</tbody>
</table>

*All patients had negative results for *B. pertussis* loop-mediated isothermal amplification and positive results for IS481 PCR. Ct, cycle threshold; –, negative; +, positive.
†At time of specimen collection.
‡Detection limit was a Ct value of 38.7, corresponding to 100 fg of DNA of *B. holmesii* ATCC51541.
§All strains were isolated from nasopharyngeal swab specimens.
A previous report suggested that *B. holmesii* and *B. pertussis* may co-circulate in young adults (7). However, the relationship between pertussis epidemics and *B. holmesii* infection is not fully understood. Our active surveillance study showed that *B. holmesii* infection spread concurrently with the *B. pertussis* epidemic, but that there was no co-infection of *B. holmesii* and *B. pertussis*. Our observations demonstrate that accurate diagnosis is needed to discriminate between *B. holmesii* and *B. pertussis* infections during a pertussis outbreak because symptoms associated with these 2 diseases are similar.

In 2012, a patient with *B. holmesii* infectious pericarditis was reported in Japan (13). This is probably the first case report of *B. holmesii* infection in Asia. Previous surveillance studies conducted in the United States and Canada have shown low rates (0.1%–0.3%) of *B. holmesii* infection in patients with cough (7,8). However, in a recent study, *B. holmesii* DNA was detected in 20% of NPS specimens collected from patients in France who had been given a diagnosis of *B. pertussis* infection (14). These surveillance data indicate that *B. holmesii* infection is present in adolescents and adults, and that the organism is associated with pertussis-like symptoms. However, other causes of viral or bacterial respiratory infection cannot be excluded. Because of lack of specific diagnostic tools to detect bordetellae, *B. holmesii* infection may have been underestimated. Accurate diagnosis and further studies are required to fully elucidate the nature of *B. holmesii* infection.

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### References


2. Lindquist SW, Weber DJ, Mangum ME, Hollis DG, Jordan J. *Bor


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**Rabies**

[ra’bēz]

From the Latin *rabere* (to rage), which may have roots in the Sanskrit *rabhas* (to do violence). Acute progressive fatal encephalomyelitis caused by neurotropic viruses in the genus Lyssavirus—from the Greek *lyssa* (frenzy or madness). In Greek mythology, Lyssa was the goddess of rage, fury, and rabies, known for driving mad the dogs of the hunter Acteon and causing them to kill their master.

Democritus (460–370) described rabies, and Hippocrates is believed to refer to the disease when he said that “persons in a frenzy drink very little, are disturbed and frightened, tremble at the least noise, or are seized with convulsions.” According to Aristotle, “Dogs suffer from the madness. This causes them to become irritable and all animals they bite to become diseased.” The disease in humans was characterized by hydrophobia, in which the sick person was simultaneously tormented with thirst and fear of water. The Roman writer Cardanus described the saliva from a rabid dog as a *virus*, the Latin word for poison.

Canine rabies has been eliminated in the continental United States. However, dog bites remain a concern for travelers to areas where the disease is enzootic.

**Sources**


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