SHORT REPORT

Bordetella holmesii in children suspected of pertussis in Argentina


1 Laboratorio VacSal, Instituto de Biotecnología y Biología Molecular IBBM – FCE – UNLP, CONICET, La Plata, Buenos Aires, Argentina
2 Meningitis and Vaccine Preventable Diseases, Division of Bacterial Diseases, National Center for Immunization and Respiratory Diseases, Office of Infectious Diseases, US Centers for Disease Control and Prevention, Atlanta GA, USA
3 Servicio de Bacteriología Clínica-INEI-ANLIS ‘Dr. Carlos G. Malbrán’, Av. Vélez Sarsfield, Ciudad Autónoma de Buenos Aires, Argentina
4 Laboratorio Central-Subsecretaría de Salud de Neuquén, Neuquén, Argentina
5 CEMAR – Dirección de Servicios de Laboratorios y Análisis Clínicos de la secretaría de Salud Pública de la Municipalidad de Rosario, Santa Fe, Argentina
6 Municipalidad de Rosario Hospital, De Niños Zona Norte – Rosario – Santa Fe, Argentina
7 Dirección Provincial de Farmacia, Bioquímica y Droguería Central del Ministerio de Salud de la Provincia de Santa Fe, Argentina

Received 10 January 2012; Final revision 26 May 2012; Accepted 29 May 2012; first published online 5 July 2012

SUMMARY

We describe nine patients (eight aged < 1 year) clinically diagnosed with pertussis yet laboratory-confirmed with Bordetella holmesii infections, a human pathogen normally isolated from blood. Most patients reported cough and cold symptoms. No death was reported. We report B. holmesii isolation in infants with respiratory symptoms in Argentina.

Key words: Bordetella pertussis, emerging infections, pertussis (whooping cough).

In Argentina, as in other countries, the number of reported pertussis cases has been increasing for over a decade [1]. Heightened public and provider awareness, improved diagnostics and surveillance, waning immunity, and bacterial evolution may explain the changing epidemiology of pertussis [1–4]. An additional explanation may be an increase in the circulation of other Bordetella spp. that produce similar syndromes. To detect these different species, several PCR methodologies with different target sequences were designed. Some of these sequences are included in Supplementary Table S1 (available online). Although the IS481 PCR assay is very sensitive, it lacks specificity, as the B. holmesii and B. bronchiseptica genomes contain copies of IS481 [4, 5]. Countries employing only the IS481 PCR assay for pertussis diagnosis lack the ability to distinguish B. pertussis from other Bordetella spp. Thus, a proportion of cases clinically and laboratory
diagnosed as pertussis may be caused by species other than *B. pertussis*, such as *B. holmesii*.

A multi-target real-time PCR assay in combination with a single-target real-time PCR assay for *ptx*/*SI* has been developed for detection of *B. pertussis*, *B. parapertussis*, and *B. holmesii* [4]. In 2010, the Argentine national reference laboratories began using this assay through collaboration with the US Centers for Disease Control and Prevention and the Latin American Pertussis Project†.

Cases included in this study were identified through the Sistema Nacional de Vigilancia de la Salud (National Health Surveillance System) between 1 July and 31 December 2010. Immediate reporting of suspected pertussis cases is mandatory in Argentina. A standard epidemiological reporting form collects clinical symptoms, demographic characteristics, vaccination history, and laboratory results for all suspected pertussis cases. This information is generally obtained when the patient is clinically suspected of pertussis. A nasopharyngeal aspirate or Dacron® swab specimen should be obtained from all suspected pertussis patients and tested by PCR or culture to attempt to confirm the diagnosis. The Argentine national laboratory network consists of 18 provincial and two national reference laboratories. All specimens are tested with conventional PCR based on IS481, *ptx* promoter and IS1001, which can distinguish between *B. pertussis* and *B. parapertussis* [6]. Specimens that are inconclusive or that are IS481 positive and *ptx* promoter or IS1001 negative are then tested at the national reference laboratories by the multi-target real-time PCR assay, which can additionally detect *B. holmesii* using the hIS1001-like primer probe set [4]. Specimens are also tested by culture using charcoal agar plates (Oxoid, France) or Bordet-Gengou agar, both supplemented with defibrinated sheep blood. Suspected *Bordetella* colonies are presumptively identified by their phenotypic characteristics and then tested biochemically and by agglutination with antiserum for *B. pertussis* and *B. parapertussis* (Murex Diagnostic, UK).

Symptoms and vaccination history of PCR-confirmed *B. pertussis* case-patients identified at the National Reference Laboratory at La Plata during 1 July–31 December 2010, were compared with the same data from confirmed *B. holmesii* case-patients for the same period. We used Epi Info™ version 3.5.3 (CDC, USA) for all analyses.

During the study period, 1475 suspected case-patients were tested for pertussis at the National Reference Laboratory at La Plata, 343 (23.9%) of which were positive for *B. pertussis* and nine (0.6%) for *B. holmesii*. The age distribution of pertussis cases was: <1 year (290 cases, 85.3%), 1–10 years (39 cases, 11.5%), and ≥11 years (11 cases, 3.2%). Three case-patients’ ages were unknown. *B. holmesii* infection was identified by real-time PCR in all nine patients, two of which were also confirmed by culture. Six case-patients lived in the province of Santa Fe, and one each lived in Chubut, Entre Rios, and Neuquén provinces, representing a geographically dispersed population. The cases were temporally spread across the study period. Seven (78%) of the case-patients were aged <6 months, one was aged 7 months, and one was aged 13 years. Five (56%) of the case-patients were female. Only one case-patient was reported to have contact with a previously suspected case of *Bordetella* infection.

For all nine *B. holmesii* case-patients, catarrhal symptoms and non-paroxysmal cough were the most frequent (Table 1). Paroxysmal cough and fever were each reported by 1/3 of case-patients. Post-tussive vomiting occurred in one case. No *B. holmesii* case-patients reported apnoea, cyanosis, or inspiratory ‘whoop’. Mean duration of cough was 7.2 days (s.d. = 11.8, data not shown). No death was reported.

We observed that *B. holmesii* case-patients presented more frequent catarrhal symptoms (*P* = 0.046) and less frequent cyanosis (*P* = 0.026) and paroxysmal cough (*P* = 0.094). The two groups presented similar cough duration (*μ* = 8.4 for *B. holmesii* vs. 7.2 for *B. pertussis*; *P* = 0.8, data not shown). A comparison

<table>
<thead>
<tr>
<th>Table 1. Comparison of case-patient symptoms, Argentina, July–December 2010</th>
</tr>
</thead>
<tbody>
<tr>
<td>Symptom</td>
</tr>
<tr>
<td>-------------------</td>
</tr>
<tr>
<td>Cough</td>
</tr>
<tr>
<td>Fever</td>
</tr>
<tr>
<td>Paroxysms</td>
</tr>
<tr>
<td>Apnoea</td>
</tr>
<tr>
<td>Inspiratory whoop</td>
</tr>
</tbody>
</table>

† The Latin American Pertussis Project is a collaboration between the US Centers for Disease Control and Prevention, the Pan American Health Organization, the Sabin Vaccine Institute, and selected ministries of health to strengthen pertussis surveillance in Latin America.
of vaccination status between the two groups did not produce a statistically significant difference (Fisher’s exact $P$ value $= 0.547$).

*B. holmesii* has been reported to cause respiratory symptoms, although it is associated primarily with bacteraemia in patients with underlying conditions [7, 8]. Most data on *B. holmesii* infection, however, are from individual case reports. We are aware of two published reports of *B. holmesii* cases in patients suspected of pertussis that were identified through surveillance [9, 10]. The majority of the *B. holmesii* case-patients identified in the Massachusetts study [10] was of adolescent (83%) or young-adult age (9%) with mild respiratory symptoms. Eleven case-patients in the Chilean report [9] positive for *B. holmesii* were spread across age groups, with three aged <12 months. In agreement with these last findings, we also detected *B. holmesii* in infants clinically diagnosed as pertussis. Although *B. holmesii* was detected in these patients, other possible pathogens were not tested for and therefore the data are suggestive, but not conclusive, that *B. holmesii* was the responsible pathogen. Frequencies of catarrhal symptoms and cyanosis in these *B. holmesii* case-patients differed significantly from those of Argentinean patients with laboratory-confirmed *B. pertussis*. However, clinical presentations of *B. holmesii* cases were similar enough for clinicians to suspect pertussis and order a pertussis laboratory test.

We did not identify reports of secondary cases, and case-patients were from four dispersed provinces distributed across time, suggesting that these cases were not part of a localized outbreak. Increased awareness in the health system – due to the steady increase in cases of pertussis – and in laboratory capacity, such as incorporating multi-target real-time PCR methodology in 2010, increased the ability to identify this pathogen. Thus, *B. holmesii* may have been circulating unidentified in Argentina previously.

Regardless, detection of nine cases (eight aged <1 year) in a 6-month period is noteworthy in light of two retrospective studies’ inability to detect *B. holmesii* in over 11000 Finnish and Dutch specimens collected between 1992 and 2003 or in 119 specimens positive for IS481 obtained from French patients aged <9 years between 2009 and 2010 [5, 11]. Our findings, which are similar to those of the Chilean report [9], may differ from those of the European countries because of a sustained increase in reported *B. pertussis* cases in Latin America over the past decade. Continued use of the multi-target, real-time PCR assay that can distinguish *Bordetella* spp. can assist in further monitoring the epidemiology and clinical picture of *B. holmesii* in Argentina.

**SUPPLEMENTARY MATERIAL**

For supplementary material accompanying this paper visit http://dx.doi.org/10.1017/S095026881200132X.

**ACKNOWLEDGEMENTS**

This work was supported by Agencia Nacional de Promoción Científica y Tecnológica – ANCPyT and ANLIS – Dr. Carlos Malbran grants to D.F.H. and M.G. D.F.H. is a member of the Scientific Career of CICBA. D.B. and M.E.G. are in receipt of fellowships from Consejo Nacional de Investigaciones Científicas y Tecnológicas – CONICET.

**DECLARATION OF INTEREST**

None.

**REFERENCES**


