More Complications Occur in Macrolide-Resistant than in Macrolide-Sensitive \textit{Mycoplasma pneumoniae} Pneumonia

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We sought to understand the situation of macrolide-resistant genotypes of \textit{Mycoplasma pneumoniae}, and analyze the relationship between macrolide-resistant genotypes and clinical manifestations of \textit{Mycoplasma pneumoniae} pneumonia (MPP). Full-length sequencing of the 23S rRNA gene of \textit{M. pneumoniae} was performed in 235 nasopharyngeal aspirates (NPAs) from children with MPP. We also retrospectively compared the clinical characteristics of macrolide-resistant (MR) \textit{M. pneumoniae} infections and macrolide-sensitive (MS) \textit{M. pneumoniae} infections. A total of 206 patients had point mutations in the \textit{M. pneumoniae} 23S rRNA gene, and these patients are referred to as MR patients. The remaining 29 patients without point mutations are referred to as MS patients. Among 206 MR patients, 199 (96.6%) had A2063G mutations, 6 had A2063T mutations, and the remaining patients had an A2064G mutation. Among the clinical manifestations, we found that the median fever durations were 8 days (range, 0 to 42 days) and 6 days (0 to 14 days) \((P < 0.01)\), the median hospitalization durations were 8 days (2 to 45 days) and 6 days (3 to 16 days) \((P < 0.01)\), and the median fever durations after macrolide therapy were 5 days (0 to 42 days) and 3 days (0 to 10 days) \((P < 0.01)\), respectively, in the MR and MS groups. We also found that the incidence of extrapulmonary complications in the MR group was significantly higher than that in the MS group \((P < 0.05)\). Moreover, the radiological findings were more serious in the MR group than in the MS group \((P < 0.05)\). The increasing prevalence of MR \textit{M. pneumoniae} has become a significant clinical issue in the pediatric patients, which may lead to more extrapulmonary complications and severe clinical features and radiological manifestations.

\textit{Mycoplasma pneumoniae} is one of the most prevalent pathogens causing community-acquired respiratory tract infections in children and young adults \((1, 2)\). \textit{M. pneumoniae} pneumonia (MPP) is usually a benign self-limited disease. However, sometimes it may cause various extrapulmonary complications and progress to a severe life-threatening pneumonia \((3–8)\). These cases might show clinical and radiological deterioration despite of macrolide antibiotic therapy for 7 days or longer \((9)\).

Severe \textit{M. pneumoniae} infections may be related to the occurrence of macrolide-resistant (MR) \textit{M. pneumoniae}. For children, macrolides are the first-choice agents for \textit{M. pneumoniae} infections. However, in recent years, many isolates of \textit{M. pneumoniae} from clinical samples showed resistance to macrolides with a high prevalence of >90% in China \((10)\). The main mechanism of resistance has been shown to be due to mutations in the domain V of the 23S rRNA of \textit{M. pneumoniae} \((10–13)\). The mutations that induce a high-level of macrolide resistance are an A-to-G transition at position 2063 and an A-to-G transition at position 2064, whereas low-level resistance is induced by an A-to-G transition at position 2617 and A-to-T transition at position 2063 \((13)\). Several studies have indicated that macrolide resistance in \textit{M. pneumoniae} may have clinical significance in terms of diminishing response to treatment with drugs \((14–16)\). The presence of MR \textit{M. pneumoniae} has been reported to be mainly associated with the persistent clinical symptoms such as fever, causing prolonged hospital stay and elevated antibiotic change rate; to our knowledge, however, no increase in the incidence of complications has been reported.

In the present study, we characterized macrolide resistance directly using \textit{M. pneumoniae}-positive nasopharyngeal aspirates (NPAs) taken from hospitalized children with MPP in our hospital by analyzing the DNA sequence of domain V of 23S rRNA. Furthermore, we retrospectively compared the different clinical characteristics of MR \textit{M. pneumoniae} infections and MS \textit{M. pneumoniae} infections, including symptoms, extrapulmonary complications, and radiological findings.

\textbf{MATERIALS AND METHODS}

\textbf{Study population and sample collection.} A total of 235 NPAs were collected from children with MPP hospitalized in Children’s Hospital, Zhejiang University School of Medicine, Hangzhou, China, from 1 April 2009 to 31 March 2010. All patients had signs and symptoms indicative of pneumonia on admission, including fever, cough, abnormal lung auscultation, and a new infiltrate on chest radiograph \((17)\). The \textit{M. pneumoniae} infection was confirmed by serologic testing (detecting \textit{M. pneumoniae} IgM by enzyme-linked immunosorbent assay) and/or \textit{M. pneumoniae} PCR tests of the NPA. All 235 children had positive results for the above-mentioned \textit{M. pneumoniae} tests. Furthermore, other microbiologic tests were performed to exclude other respiratory tract infections and tuberculosis, including the protein purified derivative test, blood cultures, nasopharyngeal aspirate/swab for common respiratory tract virus antigens (respiratory syncytial virus, influenza virus, adenovirus, and parainfluenza virus), and serology for \textit{Chlamydia pneumoniae} and \textit{Legionella pneumophila}. No other pathogens were detected by these tests.

NPAs were obtained from the patients on admission. An aliquot of DNA was extracted from NPAs and stored at \(-80^\circ\text{C}\) for determination.

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DNA was extracted using the TIANcombi DNA Lyse&Amp PCR kit (Tiangen, Hangzhou, China) in accordance with the manufacturer’s instructions.

**Amplification of MR genes.** A PCR assay, followed by direct amplicon sequencing, was developed to detect point mutations conferring resistance to macrolides in the *M. pneumoniae* 23S rRNA gene. Domain V of the 23S rRNA gene was amplified using the primers 5’-CCTAGTCGGGT AAATTCCGT-3’ and 5’-CCTAGTCGGGTAATTCCGT-3’ (TaKaRa, Shanghai, China). The primers were used to specifically amplify a 244-bp region of *M. pneumoniae*, including positions 2063 and 2064.

The PCR was carried out in a volume of 50 μl containing 1 μl of total DNA, 5 μl of 10X buffer, 3 μl of 10 mM deoxynucleoside triphosphate, 1 μl of 10 μM concentration of each primer, 38.7 μl of double-distilled H2O, and 0.3 μl of Taq DNA polymerase (Invitrogen, Milan, Italy). The reaction mixture then underwent denaturation for 5 min at 94°C and 35 PCR cycles, each consisting of 1 min at 94°C, 1 min at 53°C, and 1 min at 72°C, followed by a final extension step of 5 min at 72°C. The amplified products were analyzed by electrophoresis on a 2% agarose gel (Invitrogen) and visualized by using a Bio-Rad gel-imaging analysis system.

**DNA sequencing.** All PCR products of 244 bp were sequenced (Yingwewjii Co., Ltd., Shanghai, China), and the DNA sequences were compared to the sequence of an *M. pneumoniae* reference strain M129 (GenBank accession no. X68422) by using BLAST.

**Evaluation of clinical characteristics.** Clinical information was retrospectively collected from the medical records of the patients. Complete information about the antimicrobial agents prescribed, the clinical symptoms, and the radiologic findings was reviewed. The patient demographics, clinical symptoms, the incidence of extrapulmonary complications, and the radiologic findings were compared between MR MPP patients and MS MPP patients when discharged.

In our study, all patients were treated with 10 mg of azithromycin/kg/day for the first 3 days, and then treatment ceased for 4 days; this was followed by another 3-day treatment of azithromycin or by treatment with 15 to 30 mg of erythromycin/kg/day for 7 to 14 days. Temperature and respiratory signs and symptoms were examined at study entry and every 8 h thereafter. A febrile day was defined as a day during which the body temperature exceeded 38.0°C at least once. Total febrile days, every 8 h thereafter. A febrile day was defined as a day during which the body temperature exceeded 38.0°C at least once.

**RESULTS**

**Clinical characteristics.** Upon admission, all patients had symptoms and signs indicative of pneumonia, including fever (>38°C per axilla), cough, and abnormal breath sounds on auscultation. No patients were transferred to the intensive care unit or received mechanical ventilation during hospitalization. All patients were PCR positive, and for 199 patients there were positive serologic results. The median age of the 235 children was 4 years (range, 0 to 14 years). A total of 149 patients were males.

**Analysis of MR genotypes.** The 244-bp bands were detected in all samples. Of the 235 patients with MPP, 206 (87.7%) showing point mutations in domain V of 23S rRNA were defined as MR patients. The remaining 29 patients (12.3%), who showed no point mutations, were defined as MS patients. Of the MR patients, 199 (96.6%) were positive for an A-to-G transition mutation at position 2063 in domain V of 23S rRNA gene (A2063G), 6 showed an A-to-T transition mutation at position 2063 (A2063C), and the remaining one had an A-to-G transition mutation at position 2064 (A2064G). In general, the mutation rate at position 2063 reached 99.5% (205/206).

**Comparison of clinical characteristics between MR and MS patients.** Patients were categorized in one of two groups, MR or MS, on the basis of presence or absence of 23S rRNA gene mutations. A total of 206 patients were in the MR group (132 males, 74 females), with a median age of 4 years (range, 0 to 14 years). Another 29 patients were in the MS group (17 males, 12 females), with a median age of 5 years (range, 0 to 11 years). There were no significant differences between the two groups with respect to age and sex distribution (Table 1).

**Clinical characteristics in the MR and MS patients, including incidence of complications, radiologic findings, and clinical course, are summarized in Table 1. Of the 206 MR patients, extrapulmonary complications were found in 61 cases (29.6%), including liver function abnormalities in 27, myocarditis in 15, rash in 11, encephalitis in 4, proteinuria in 2, hemolytic anemia in 1, and arthritis in 1. In the MS group, 3 patients (10.3%) who had liver function abnormalities had extrapulmonary complications. There was a significant difference in the incidence of extrapulmonary complications between the two groups (*P* < 0.05). We also evaluated the severity of MPP in all patients. The incidences of severe MPP were 18.4% in the MR group and 3.4% in the MS group, with a significant difference (*P* < 0.05). In addition to clinical symptoms, the radiologic findings were more severe in the MR group than in the MS group, with 61.7% of the patients in the MR group showing large lesions versus 41.4% in the MS group (*P* < 0.05).

Concerning clinical course, we found that the median durations of fever were 8 days (0 to 42 days) in the MR group and 6 days (0 to 14 days) in the MS group (*P* < 0.01). The median hospitalization durations were 8 days (2 to 45 days) and 6 days (3 to 16 days) (*P* < 0.01), respectively. The median fever duration after...
macrolide therapy was significantly longer in the MR patients than in the MS patients (median of 5 days versus 3 days, \(P < 0.01\)).

**DISCUSSION**

Although *M. pneumoniae* infection was traditionally thought to be a self-limited process, more and more severe cases even fatal cases of *M. pneumoniae* infections were reported in recent years (3–8). The reasons why more severe *M. pneumoniae* infections occurred remain unclear, but resistance to macrolides does have a close association with the emergence and increase of MR strains. Since 2000, the emergence of macrolide resistance has been reported mainly in Asia (21). In China, prevalence of MR *M. pneumoniae* isolated in pediatric cases has increased rapidly. In 2009, Xin et al. (22) reported that 46/50 (92%) *M. pneumoniae* isolates from pediatric patients were resistant to macrolides. In 2010, Liu et al. (10) reported that 90/100 (90%) *M. pneumoniae* isolates were resistant to macrolides. In our study, we found that 206 (87.7%) NPAs from patients with MPP showed point mutations in domain V of 23S rRNA, which were considered MR. The high prevalence of MR was similar to the data reported previously.

MR genotypes are defined by specific point mutations in the V domain of the *M. pneumoniae* 23S rRNA gene. Several potential transition mutations in the complete sequence of the 23S rRNA were found: A2063G, A2064G, and A2063T. A2063G and A2064G mutations are responsible for the high-level macrolide resistance in *M. pneumoniae* (23). Morozumi et al. (24) found that among 380 *M. pneumoniae* isolates from 3,678 pediatric patients with community-acquired pneumonia, 50 MR strains had an A2063G transition in domain V of the 23S rRNA, whereas 5 had an A2064G transition. In our study, we amplified the full length of the 23S rRNA fragment from 235 NPAs of patients with MPP and compared it to the sequence of M129. We found that among the 206 MR patients 199 (96.6%) possessed an A2063G transition in domain V of the 23S rRNA, 6 cases showed A2063T mutation, and 1 showed an A2064G mutation. Furthermore, the point mutations at position 2063 in domain V of the 23S rRNA gene accounted for 99.5% (205 cases) of all mutations, which was consistent with that reported in the literature (25, 26). The other less frequent point mutation, at position 2617, was also detected, but no positive results were found (data not shown).

As for the current situation of macrolide resistance of *M. pneumoniae*, it is important to evaluate the clinical significance of MR *M. pneumoniae*. Suzuki et al. (16) found the total febrile days and the numbers of febrile days during macrolide administration to be longer in MR patients than in MS patients. Ensuing clinical studies were reported with similar results. In our study, MR *M. pneumoniae*-infected patients showed much a longer median duration of fever, a longer median hospitalization duration, and a longer median fever duration after macrolide therapy than were observed for patients with MS *M. pneumoniae*. These results implied that MR *M. pneumoniae* was refractory to treatment and could result in a prolonged clinical course.

More interestingly, we found that the incidence of extrapulmonary complications in the MR patients was higher than in the MS patients. A total of 29.6% (61/206) of the MR patients had extrapulmonary complications versus 10.3% in the MS group. Moreover, only one patient (3.4%) demonstrated severe pneumonia in the MS group, and 38 of 206 MR patients (18.4%) were diagnosed as severe having MPP. In addition to the clinical symptoms, 61.7% of patients in the MR group had large lesions observed radiologically versus 41.4% in the MS group. To our knowledge, this is the first report showing severe clinical course and more extrapulmonary complications in the MR patients.

Several reasons may explain the difference between previous reports and the findings of our study. First, the children with MPP enrolled in our study reflected a natural clinical course. Because of potential side effects, fluoroquinolones are contraindicated in all children, and tetracyclines, such as doxycycline and minocycline, can only be used in children ≥8 years old; there are almost no alternatives for treating MR *M. pneumoniae* infection in children, especially pre-school-aged children. Therefore, all of the patients in our study received only macrolides. Local *M. pneumoniae* loads at 48 h after macrolide therapy in the MR patients were reported to be much higher than those in the MS patients (14). The higher and more persistent *M. pneumoniae* stimulation may induce a much stronger host response, especially interleukin-8 (IL-8)- and IL-18-
Investigated.

on clinical outcome for children with MPP needs to be further sequenced. The 23S rRNA gene from 235 NPAs and compared it to macrolide resistance. The mutations at positions 2063 and 2064 is usually associated with patients. However, it is widely acknowledged that the presence of M. pneumoniae measured, because in MR patients compared to MS patients. Our study was the first to demonstrate a higher incidence of extrapulmonary complications in MR patients compared to MS patients.

One limitation of our study is that the MIC values were not measured, because M. pneumoniae was not isolated from all these patients. However, it is widely acknowledged that the presence of mutations at positions 2063 and 2064 is usually associated with macrolide resistance.

In conclusion, we amplified and analyzed the full-length sequence of the 23S rRNA gene from 235 NPAIs and compared it to that of the M. pneumoniae reference strain M129. We have shown that the increasing prevalence of MR M. pneumoniae has become a significant clinical issue in pediatric patients, and it may lead to more extrapulmonary complications, severe clinical features, and radiological manifestations. The impact of macrolide resistance on clinical outcome for children with MPP needs to be further investigated.

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