

Characterization and Dynamics of Middle Ear Fluid and Nasopharyngeal Isolates of *Streptococcus pneumoniae* from 12 Children Treated with Levofloxacin[∇]

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Children who had acute otitis media and were treated with levofloxacin were assessed for the emergence of fluoroquinolone-resistant *Streptococcus pneumoniae*. Nasopharynx cultures were obtained from patients at the entry to and during levofloxacin therapy. All nasopharynx isolates ($n = 59$) from 12 children were levofloxacin susceptible without *parC/E* or *gyrA/B* mutations. Pneumococcal nasopharynx persistence was not associated with levofloxacin resistance.

Acute otitis media (AOM) is the leading cause for the prescribed use of antibiotics in children (12). The primary pathogen in AOM is *Streptococcus pneumoniae* (12). The high prevalence of penicillin resistance and multidrug resistance in *S. pneumoniae* isolates has prompted the use of additional agents for the treatment of recurrent/persistent AOM in patients who fail therapy with first- and second-line agents such as β -lactams, macrolides, or sulfonamides. Fluoroquinolones, which have been used widely as an effective therapeutic agent for respiratory tract infections in adults, were studied recently in children for their safety and efficacy in the treatment of recurrent/persistent AOM (1, 2, 8).

An important consideration for the treatment of AOM is how the prescribed antibiotic affects the colonization of the nasopharynx by *S. pneumoniae*. Previous studies have shown that the use of β -lactam and macrolide agents can result in a higher incidence of posttreatment drug-resistant pneumococcal isolates in the nasopharynx than that of pretreatment isolates (4, 6, 11). These resistant pneumococci then may cause a superinfection of the middle ear fluid (MEF) (5). In this study, we examined *S. pneumoniae* isolates from MEF and nasopharynx cultures from children with recurrent/persistent AOM who were treated with levofloxacin to determine if resistance developed during treatment.

A complete description of this trial, conducted as part of a multicenter, nonrandomized, open-label study conducted in the United States, Argentina, Costa Rica, and Israel, has been published previously (1). The current study focused on isolates collected at the Israeli site following approval by the investigational review board at each institution. A data safety monitoring committee was established to monitor for any potential safety risks for children participating in this trial.

Patients (≥ 6 months old to < 5 years old) were enrolled if (i) they had clinical signs and symptoms of AOM and (ii) they were diagnosed with either persistent disease (AOM persisting despite > 72 h of appropriate antibiotic therapy) or recurrent disease (≥ 3 episodes of AOM in 6 months or ≥ 4 episodes of AOM within the previous year). Patients received 10 mg/kg of body weight of levofloxacin suspension orally twice daily for 10 days. Before dosing (pretreatment, day 1), MEF was collected for bacterial culture. Patients were clinically evaluated at three ensuing visits: 4 to 6 days after receiving the first dose of levofloxacin (on therapy [OT]), 2 to 5 days after receiving the last dose (end of therapy [EOT]), and 10 to 17 days after the last dose was received (end of study [EOS]). MEF was collected on day 1 from both ears if bilateral bulging was observed, and a second tap was mandatory at the OT visit if a positive baseline MEF culture was present. Nasopharyngeal swabs were taken at each visit to assess the presence of pneumococci. Of the 83 children enrolled at this site, 24 had pneumococci isolated from the baseline MEF, and 18 of these children met the criteria for microbiologic evaluability (i.e., they returned for a second evaluation and completed their protocol-defined procedures). Of these 18 children, 12 had *S. pneumoniae* organisms isolated from MEF cultures upon entry into the trial and from nasopharynx cultures on at least one occasion during therapy. These isolates were analyzed in this study. It is important to note that none of the children received prior immunization with the 7-valent conjugate pneumococcal vaccine as it was not yet licensed in Israel at the time of this study.

MICs were determined by the broth microdilution test using panels manufactured by Trek Diagnostic Systems (Cleveland, OH) according to CLSI recommendations (3). Pyrosequencing technology (Biotage, Westborough, MA) was used as described previously (10) to detect mutations in the quinolone resistance-determining region (QRDR) of *parC*, *gyrA*, *parE*, and *gyrB*, the targets of the fluoroquinolones (15). Pulsed-field gel electrophoresis (PFGE) was performed as described previously (9), and PFGE patterns were compared as described by Tenover et al. (16). Serotyping was performed by the standard

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TABLE 1. *S. pneumoniae* isolation dates, levofloxacin MICs, and genotype

| Patient | Day of study | Visit ^b | MEF culture ^a | | | Nasopharynx culture | | |
|---------|--------------|--------------------|--------------------------|------------------------------------|------------|--------------------------|-----------------|-----------------|
| | | | Levofloxacin MIC (μg/ml) | PFGE type | Serotype | Levofloxacin MIC (μg/ml) | PFGE type | Serotype |
| 1 | 1 | PT | 0.5 | A | 6A | 0.5 | A | 6A |
| | 5 | OT | NA ^c | — ^d | | 1 | A | 6A |
| | 24 | EOS | NA | — | | 0.5 | A | 6A |
| | 39 | UNS | NA | — | | 0.5 | A | 6A |
| 2 | 1 | PT | 1 | W | 19A | 0.5 | B | NT ^e |
| | 5 | OT | NA | — | | 0.5 | C | NT |
| | 15 | EOT | NA | — | | 0.5 | B | NT |
| | 19 | EOS | NA | — | | 1 | D | 9V |
| 3 | 1 | PT | 0.5 | E | 23F | 0.5 | E | 23F |
| | 4 | OT | NA | — | | 0.5 | E | 23F |
| | 15 | EOT | NA | — | | 0.5 | E | 23F |
| | 25 | EOS | NA | — | | 0.5 | E | 23F |
| 4 | 1 | PT | 1 | F | 19F | 1 | F | 19F |
| | 6 | OT | 1 | F | 19F | 1 | F | 19F |
| | 13 | EOT | NA | — | | 1 | F | 19F |
| | 21 | EOS | NA | — | | 0.5 | G | NT |
| | 35 | UNS | NA | — | | 1 | F | 19F |
| 5 | 1 | PT | 0.5 | H | 6B | NA | — | |
| | 5 | OT | NA | — | | NA | — | |
| | 25 | EOS | NA | — | | 0.5 | H | 6B |
| | 37 | UNS | 0.5 | H | 6B | NA | — | |
| 6 | 1 | PT | 1 (right) 1 (left) | I I | 14 14 | 1 NA | J — | 19F — |
| | 13 | EOT | NA | — | | 1 | I | 14 |
| | 18 | EOS | NA | — | | 0.5 | K | 6B |
| 7 | 1 | PT | 0.5 | L | 6A | 0.5 | L | 6A |
| | 23 | EOS | NA | — | | 1 | M | NT |
| 8 | 1 | PT | 0.5 | N | 23F | 1 | N | 23F |
| | 5 | OT | NA | — | | 1 | N | 23F |
| | 12 | EOT | NA | — | | 1 | N | 23F |
| 9 | 1 | PT | 1 (right) 1 (left) | O O | 15A 15A | 0.5 NA | O — | 15A — |
| | 1 | OT | NA | — | | 1 | O | 15A |
| | 5 | EOT | NA | — | | 1 | O | 15A |
| | 12 | EOS | NA | — | | 1 | O | 15A |
| | 23 | EOS | NA | — | | 1 | O | 15A |
| | 40 | UNS | NA | — | | 1 | O | 15A |
| 10 | 1 | PT | 0.5 (right) 1 (left) | O1 ^f O2 ^f | 19F 19F | 1 | O2 ^e | 19F |
| | 13 | EOT | NA | — | | 1 | P | 11A |
| | 21 | EOS | NA | — | | 1 | O1 ^f | 19F |
| | 42 | UNS | NA | — | | 0.5 | O1 ^f | 19F |
| 11 | 1 | PT | 1 | Q | 17F | 1 | Q | 17F |
| | 14 | EOT | NA | — | | 1 | R | 15B |
| | 21 | EOS | NA | — | | 0.5 | S | 10A |
| 12 | 1 | PT | 1 | T | 19F | 1 | T1 | 19F |
| | 12 | EOT | 0.5 | U | 4 | 1 | U | 4 |
| | 41 | UNS | NA | — | | 0.5 | V | 15B |

^a Some patients had bilateral AOM, and MEF was removed from both left and right ears.

^b PT, pretreatment visit; UNS, unscheduled visit.

^c NA, not applicable since no specimen was obtained.

^d —, no PFGE data were available because *S. pneumoniae* was not isolated during that visit.

^e NT, nontypeable as *S. pneumoniae* because cultures did not react to Omni-sera.

^f PFGE types O1 and O2 are variants (1- to 2-band difference) of PFGE type O.

Quellung method with sera from the Statens Seruminstitut (Copenhagen, Denmark).

All *S. pneumoniae* MEF or nasopharyngeal cultures were susceptible to levofloxacin, with MICs ranging from 0.5 to 1 $\mu\text{g/ml}$ (Table 1). Cultures isolated during or after levofloxacin treatment had the same levofloxacin MIC (or were within one doubling dilution) as the pretreatment isolate(s). None of the isolates had *parC*, *parE*, *gyrA*, or *gyrB* QRDR mutations known to contribute to reduced fluoroquinolone susceptibilities.

Three patients (patients 6, 9, and 10) had *S. pneumoniae* cultured from MEF from both ears at the pretreatment visit, and these isolates were the same or closely related, as determined by PFGE. Three other patients (patients 4, 5, and 12) had *S. pneumoniae* isolated from MEF obtained at some other visit in addition to that from the pretreatment visit. In two of these patients (no. 4 and 5), all MEF *S. pneumoniae* isolates were the same, as determined by PFGE, and in the other patient (no. 12), the MEF *S. pneumoniae* isolates were different (Table 1).

In nine patients (patients 1, 3, 4, and 7 to 12), the pretreatment MEF and nasopharyngeal pneumococcal cultures were identical or closely related, as shown by PFGE (Table 1). All of the nasopharyngeal cultures obtained subsequently from four (patients 1, 3, 8, and 9) of these nine patients (at the OT, EOT, or EOS visit) matched the PFGE patterns of their pretreatment MEF cultures. Subsequent nasopharyngeal cultures from six patients (patients 4, 6, 7, 10, 11, and 12) did not have the same PFGE patterns as those from their pretreatment MEF culture, indicating colonization by a new strain. Only one patient (no. 2) did not have any nasopharyngeal cultures that had PFGE patterns similar to the pretreatment MEF culture, demonstrating the simultaneous presence of two different strains in the ear and the nasopharynx.

Among the 12 children in whom persistent colonization with *S. pneumoniae* was documented, there was no phenotypic evidence of the emergence of levofloxacin resistance, as levofloxacin MICs remained between 0.5 and 1 $\mu\text{g/ml}$ during treatment. *S. pneumoniae* resistance to fluoroquinolones occurs in a stepwise fashion, with the first mutation occurring typically in the QRDR of either *parC* or *gyrA* (depending on the selecting fluoroquinolone), leading to decreased fluoroquinolone susceptibility (15). A second mutation in the paralog gene usually causes a strain to become fluoroquinolone resistant (15). *S. pneumoniae* isolates with a single *parC* QRDR mutation generally have levofloxacin MICs of 1 or 2 $\mu\text{g/ml}$ (10). Susceptibility testing alone may not detect isolates that gain a single QRDR mutation while the child is being treated with levofloxacin. However, genetic analysis of these isolates showed that there were no first-step mutants since QRDR mutations in the target genes were not present. Similarly, Leibovitz et al. reported that all pneumococcal isolates collected from children treated with gatifloxacin for recurrent/nonresponsive AOM were gatifloxacin susceptible (13). However, it is unknown if any of the isolates were first-step mutants, since screening for QRDR mutations was not performed.

Bacterial eradication from MEF during therapy is associated with a higher clinical success rate (7). In this study, the initial *S. pneumoniae* MEF culture was eradicated in 11 of the 12 children by the OT visit; however, six of the children did have another occurrence of otitis media after the EOT or the EOS

visit. Libson et al. reported that the failure to eradicate *S. pneumoniae* from the nasopharynx by the completion of antibiotic therapy predisposed the patient to a recurrence of AOM (14). The 12 patients from our study were chosen for examination because *S. pneumoniae* organisms were isolated from the nasopharynx throughout their treatment. In the Libson et al. study recurrent AOM was usually caused by a pneumococcal isolate in the MEF that was found in the nasopharynx at the EOT visit (14). We observed a similar outcome for one of the six children (no. 5), who had another occurrence of AOM after the EOS visit and had a pneumococcal isolate cultured from the MEF that matched the nasopharynx isolate at the EOS visit. In the other five children, the MEF cultures were negative.

This small experience does not establish the fact that levofloxacin resistance may not emerge in the clinical treatment of children with recurrent/persistent AOM. However, these results clearly indicate that for the children in this study, persistent colonization of the nasopharynx was not associated with levofloxacin resistance. It is uncertain if these findings could be generalized to other patients.

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