A Prospective Study of Etiology of Childhood Acute Bacterial Meningitis, Turkey

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Determination of the etiology of bacterial meningitis and estimating cost of disease are important in guiding vaccination policies. To determine the incidence and etiology of meningitis in Turkey, cerebrospinal fluid (CSF) samples were obtained prospectively from children (1 month–17 years of age) with a clinical diagnosis of acute bacterial meningitis. Multiplex PCR was used to detect DNA evidence of Streptococcus pneumoniae, Haemophilus influenzae type b (Hib), and Neisseria meningitidis. In total, 408 CSF samples were collected, and bacterial etiology was determined in 243 cases; N. meningitidis was detected in 56.5%, S. pneumoniae in 22.5%, and Hib in 20.5% of the PCR-positive samples. Among N. meningitidis– positive CSF samples, 42.7%, 31.1%, 2.2%, and 0.7% belonged to serogroups W-135, B, Y, and A, respectively. This study highlights the emergence of serogroup W-135 disease in Turkey and concludes that vaccines to prevent meningococcal disease in this region must provide reliable protection against this serogroup.

Acute bacterial meningitis is one of the most severe infectious diseases, causing neurologic sequelae and accounting for an estimated 171,000 deaths worldwide per

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group; currently, serogroups A, B, and C account for >90% of meningococcal disease worldwide (12). However, the epidemiologic landscape is constantly changing, and with increasing international travel and cross-border migration, the epidemiology of this disease will remain dynamic. Currently, serogroups A and C predominate throughout Asia and Africa, whereas serogroups B and C are responsible for most cases in Europe and North America (11,13–18). In several countries, including the United States, the proportion of disease caused by serogroup Y has increased over the past decade, where it now accounts for approximately one third of meningococcal cases (19). Serogroup W-135 has also recently emerged in some parts of the world, primarily in the Middle East and Africa, in some instances causing large epidemics (20).

The annual Hajj pilgrimage to Mecca is a major international event; ≈2 million people from around the world gather in one place, where the extreme crowding provides an ideal environment for transmission of meningococcal carriage. On several occasions, meningococcal disease outbreaks have subsequently spread worldwide by returning pilgrims. A major serogroup A meningococcal disease epidemic occurred in the 1980s, affecting Muslim pilgrims initially, followed by populations in other Middle Eastern and African countries (21). After this epidemic, Hajj pilgrims were vaccinated with a bivalent (A and C) meningococcal polysaccharide vaccine before entering Saudi Arabia. With the emergence of serogroup W-135 meningococcal disease among Hajj pilgrims in the Middle East during 2000 and 2001 (20), vaccine recommendations for pilgrims were changed to quadrivalent (A, C, W-135, and Y) meningococcal polysaccharide vaccine in 2002 (22).

Global surveillance of confirmed meningococcal cases, including surveillance of the diversity of causative strains, is essential to managing disease and developing vaccines. This study was undertaken to determine the current etiology of bacterial meningitis in Turkey, with particular emphasis on serogroup distribution of meningococci. Turkey is a predominantly Muslim country, and as such epidemics originating at the Hajj may have an effect on the national epidemiology. Although limited epidemiologic studies are available, cases of invasive meningococcal disease as well as carriage of serogroup W-135 have been reported in Turkey (23–25). This finding is in contrast to Western Europe, where the incidence of W-135 disease remains low. Turkey has no surveillance system for bacterial meningitis, and exact rates of meningococcal disease and serogroup distribution are unknown. Reliable surveillance data from countries such as Turkey are vital to understand, and better anticipate, the constantly changing landscape of bacterial meningitis and meningococcal disease.

Materials and Methods

Study Design

From February 16, 2005, through February 15, 2006, active surveillance of acute bacterial meningitis among children admitted to 12 participating hospitals was undertaken. Turkey is divided into 7 geographic areas (Figure 1). Twelve health centers in 9 cities located in all of these 7 geographic regions were selected to represent the population characteristics of the country. Two centers from each of the 3 biggest cities and 1 center from each of the other cities were included. Each health center served as a referral center for its region in the field of pediatric diseases. The centers serve ≈32% of the entire pediatric population of Turkey. Approval was obtained from the ethical committees of the participating centers and Ministry of Health.

In each hospital, suspected cases of acute bacterial meningitis were identified by a pediatrician, based on the following criteria: any sign of meningitis (fever [axillary measurement ≥38°C], vomiting [≥3 episodes in 24 h], headache, meningeal irritation signs [bulging fontanel, Kernig or Brudzinski signs, or neck stiffness]) in children >1 year of age; fever without any documented source; impaired consciousness (Blantyre Coma Scale <4 if <9 months of age and <5 if ≥9 months of age) (26); prostration (inability to sit unassisted if ≥9 months of age or breastfeed if <9 months of age) in those <1 year of age; and seizures (other than those regarded as simple febrile seizures with full recovery within 1 h). For each suspected case, demographic data, predominant clinical signs and symptoms, prior history of use of antimicrobial agent, and laboratory results were recorded by using a standardized case report form.

Cerebrospinal fluid (CSF) samples were obtained from all patients <17 years of age with clinical suspected meningitis. Patient samples were included in further analyses.
if the CSF had 1) >10 leukocytes/mm³ in the CSF, and/or 2) higher CSF protein levels than normal for the patient’s age, and/or 3) lower CSF glucose levels than normal for the patient’s age. In addition to these patients, all who had a positive CSF culture, PCR, Gram stain, or antigen detection test result were also included in the study. No neonates (<1 month of age) were included in the study since the pathogens of neonatal meningitis were expected to be different than those of the older case-patients (3).

CSF Cultures and Bacterial Isolates

CSF cultures, Gram stain, and latex agglutination tests (Wellcogen Bacterial Antigen Kit, Lenexa, KS, USA) were performed in the local hospitals. A CSF sample (minimum 0.5 mL) from each patient was kept at −20°C until transportation to the Central Laboratory (Hacettepe University, Ankara, Turkey) for PCR analysis. CSF samples and, if available, bacterial isolates were sent to the Central Laboratory, where all isolates were recultured on chocolate and blood agars and grown at 37°C in 5% CO₂. Suspected meningococcal colonies were characterized by Gram stain, oxidase test, and rapid carbohydrate utilization test (Gallerie Pasteur, Pasteur Merieux, Lyon, France). The microbiology laboratory records were cross-checked in each hospital for missing data. The phenotypic determination, based on the antigenic formula (serogroup:serosubtype) of meningococcal isolates, was performed by standard methods in the Meningococcal Reference Unit, Health Protection Agency, Manchester, United Kingdom (18,20–25,27).

DNA Isolation

All samples collected in the Central Laboratory were kept at −80°C and were thawed immediately before each test. Bacterial colonies were suspended in 500 μL sterile double-distilled water and vortexed. Bacterial suspensions and CSF were boiled for 3 min at 100°C, then centrifuged for 5 min at 10,000 × g, and the supernatant was retained. DNA concentration was estimated spectrophotometrically, and 15 μL (≈50 ng) was used in each final reaction mixture.

PCR Amplification

For the simultaneous identification of bacterial agents, single tube, multiplex PCR assay was performed. The specific gene targets were ctrA, bex, and ply for N. meningitidis, Hib, and S. pneumoniae, respectively (28). In each assay, the final reaction mixture of 50 μL contained 15 μL (≈50 ng) DNA, 1× PCR buffer, 3 mmol/L MgCl₂, 200 μmol/L of each dNTP (AB Gene, Epsom, UK), 0.6 μmol/L of each corresponding oligonucleotide primer (Sigma Aldrich, Seelze, Germany) as described (28) and 1 U of Taq polymerase (AB Gene). The PCR was performed by using a DNA thermal cycler (Perkin-Elmer Cetus, Emeryville, CA, USA model 9600) under the following conditions: a first cycle of denaturation at 95°C for 5 min followed by 35 cycles of 95°C for 25 s, 57°C for 40 s, and 72°C for 60 s.

Among the samples positive for N. meningitidis, serogroup prediction (A, B, C, W-135, and Y) was based on the oligonucleotides in the siaD gene for serogroups B, C, W-135, and Y and in orf2 of a gene cassette required for serogroup A (28). For serogroup determination, amplification reactions (50 μL) contained 15 μL of DNA, 60 mmol/L Tris-HCl (pH 8.8), 17 mmol/L (NH₄)₂SO₄, 5 mmol/L MgCl₂, 0.5 mmol/L of each dNTP, 0.3 μmol/L corresponding oligonucleotides, and 1 U of Taq polymerase. The PCR conditions were as follows: denaturation at 94°C for 3 min, followed by 35 cycles of 92°C for 40 s, 55°C for 30 s, and 72°C for 20 s in a DNA thermal cycler. A final extension cycle at 72°C for 10 min was then performed (29).

All amplicons were analyzed by electrophoresis on standard 3% agarose gels and visualized by using UV fluorescence. A negative control consisting of distilled water and a positive control consisting of a reference strain (S. pneumoniae ATCC 49613, Hib ATCC 10211, N. meningitidis serogroup C L94 5016 also known as C11 [C:16:P1.7–1,1], serogroup A M99 243594 [A:4,21:P1.20,9], serogroup Y M05 240122 [Y:NT:P1.5], serogroup W135 M05 240125 [W135:2a:NST], serogroup B M05 240120 [B:NT:NST]) were analyzed simultaneously.

Statistical Analysis

Continuous variables were compared by the Student t test and categorical variables with χ² or Fisher exact tests. A 2-tailed p value <0.05 was considered significant. All statistical analysis was performed with SPSS version 11.5 (SPSS Inc, Chicago, IL, USA).

Results

Meningitis Cases

In total, 408 children were hospitalized with a clinical diagnosis of meningitis during the study period (Figure 1), and a CSF sample from each patient was obtained. The distribution of these suspected cases according to the geographic regions was as follows: 109 (26.7%) in Southern Anatolia, 74 (18.1%) in Aegean region, 79 (19.6%) in Central Anatolia, 53 (13.0%) in Marmara region, 43 (8.7%) in Eastern Anatolia, 24 (5.9%) in Black Sea region, and 26 (6.4%) in Mediterranean region. The mean age of the 408 children was 4.8 years (standard deviation 4.1 years), and the boy-to-girl ratio was 1.5:1. Of 408 patients diagnosed with acute bacterial meningitis, 20 (4.9%) died and 14 (5.7%) of these deaths were among patients with laboratory-confirmed cases.
Laboratory-Confirmed Meningitis Cases and Etiology

Of the 408 cases, bacterial meningitis was confirmed by PCR, culture, or latex agglutination test in 243 (59.6%) patients. Regional incidence rates of laboratory-confirmed meningitis were estimated as ranging from 1/100,000 population in the Black Sea region to 10.9/100,000 population in the Southeast Anatolia region (Figure 1). Nationwide, the highest incidence was in children 1–12 months of age and was slightly more common in boys. The boy-to-girl ratio of the confirmed cases was 1.3:1, and the age distribution was as shown in Figure 2.

Overall, the diagnosis of acute bacterial meningitis was confirmed with CSF culture in 41 (17%) of 243 cases, with latex agglutination test in 56 (23%), and with PCR in 243 (100%) (Table). Latex agglutination test was positive in 37 cases for \textit{N. meningitidis}, in 10 cases for \textit{Hib}, and in 9 cases for \textit{S. pneumoniae}.

Where data were available, 7 (17%) of 41 cases with positive CSF culture and 111 (54.9%) of 202 cases with negative CSF culture had a history of use of antimicrobial agent(s) before lumbar puncture, which may account for the relatively low diagnosis rate by using this technique. \textit{N. meningitidis} was reported in 23 cases, \textit{S. pneumoniae} was reported in 12, and \textit{Hib} was reported in 6 cases as positive in CSF culture.

Blood culture was positive in 12 (4.9%) of 243 cases—4 each of \textit{N. meningitidis}, \textit{Hib}, and \textit{S. pneumoniae}. Phenotyping of 21 available isolates indicated W135:2a:P1.5,2 (5 cases), A:21:NT:P1.10 (1 case), B:NT:P1.12,4 (2 cases), B:22:NT:NT (2 cases), B:NT:NT:P1.14 (2 cases), B:15:P1.7,16 (3 cases), B:14:NT:P1.13 (2 cases), B:15:NT:P1.16 (2 cases), X:NT:P1.7,1 (1 case), Y:NT:P1.5:NT (1 case).

PCR analysis was by far the most reliable method of confirming bacterial meningitis, accounting for all confirmed cases with 243 positive results. In these PCR-positive samples, 138 (56.5%) were attributable to \textit{N. meningitidis}, 55 (22.5%) to \textit{S. pneumoniae}, and 50 (20.5%) to \textit{Hib} (Figure 3). Of the 408 patients, 118 (48.5%) of 243 cases with positive PCR and 96 (58.2%) of 165 cases with negative PCR had received antibacterial drugs in the week before CSF sampling.

In the evaluation of the bacterial agents among the 7 different geographic regions of the country, \textit{N. meningitidis} was the most common cause of acute bacterial meningitis in all regions except the Mediterranean region, located on the southern coast of Turkey. Here \textit{S. pneumoniae} were the prominent bacteria, and \textit{N. meningitidis} were detected in only 2 cases.

Comparison of the incidence of \textit{N. meningitidis}, \textit{S. pneumoniae}, and \textit{Hib} among different age groups demonstrated that \textit{N. meningitidis} was the prominent bacterial agent causing acute bacterial meningitis, especially in children <7 years of age (Figure 2). The highest incidence was detected during the first year of life for all 3 bacteria.

CSF findings were recorded for 368 (90.2%) of 408 CSF samples sent to the Central Laboratory. As a mean, in PCR-negative samples, CSF protein level was significantly lower (70.2 vs. 130.4 mg/dL; \( p = 0.003 \)) and glucose level was significantly higher (55.6 vs. 41.3 mg/dL; \( p = 0.01 \)) than in PCR-positive samples. Total cell counts were not significantly different in PCR-negative and -positive samples (157.9 and 211.1/mm\(^3\); \( p>0.05 \)), but polymorphonuclear cell count was significantly higher in PCR-positive samples (8,284.3 vs. 38.5/mm\(^3\); \( p = 0.001 \)). Among the PCR-positive samples, total cell and polymorphonuclear cell counts were not significantly different between samples positive for \textit{N. meningitidis}, \textit{S. pneumoniae}, and \textit{Hib} (\( p>0.05 \)).

Meningococcal Epidemiology

Among the samples that were positive for \textit{N. meningitidis} following PCR analysis, serogroup W-135 was the cause of most infections; 59 (42.7%) cases were serogroup W-135, 43 (31.1%) were serogroup B, 3 (2.2%) were serogroup Y, and 1 (0.7%) was serogroup A. There were no cases with a positive result for serogroup C (this was also

![Figure 2. Distribution of bacteria causing childhood acute bacterial meningitis in different age groups. \textit{Neisseria meningitidis} was the most common cause of meningitis, and the highest estimated incidence was in children <1 year of age for all 3 bacteria.](image_url)

**Table. Laboratory confirmation of bacterial meningitis**

<table>
<thead>
<tr>
<th>Category</th>
<th>PCR positive (n = 243)</th>
<th>PCR negative (n = 165)</th>
<th>Total (n = 408)</th>
</tr>
</thead>
<tbody>
<tr>
<td>CSF culture</td>
<td></td>
<td></td>
<td></td>
</tr>
<tr>
<td>Positive</td>
<td>41</td>
<td>0</td>
<td>41</td>
</tr>
<tr>
<td>Negative</td>
<td>202</td>
<td>165</td>
<td>367</td>
</tr>
<tr>
<td>Latex agglutination test</td>
<td></td>
<td></td>
<td></td>
</tr>
<tr>
<td>Positive</td>
<td>53</td>
<td>3</td>
<td>56</td>
</tr>
<tr>
<td>Negative</td>
<td>190</td>
<td>162</td>
<td>352</td>
</tr>
</tbody>
</table>

*Specimens positive for individual assays and combinations of assays. CSF, cerebrospinal fluid.
the case following analysis of CSF culture-positive samples) and in 32 (23.2%) *N. meningitidis*–positive samples the serogroup could not be determined by the PCR assay.

Analysis by age reveals the greatest meningococcal disease incidence is in children <3 years of age, particularly infants <1 year of age. The numbers of cases caused by the 2 most common *N. meningitidis* serogroups (serogroups W-135 and B) were similar in the most vulnerable age groups (<3 years of age), but W-135 was more common in children 4–16 years of age (Figure 4).

Etiologic and meningococcal serogroup distribution among the different geographic regions is illustrated in Figure 5. *N. meningitidis* serogroup W-135 was more prominent than the other meningococcal serogroups in the Southeast Anatolia, Aegean (Western Turkey), Eastern Anatolia, and Black Sea regions. *N. meningitidis* serogroup B was much more common in the Marmara region (northwestern Turkey), and in the Central region; the numbers of serogroup B and serogroup W-135 cases were similar. The Mediterranean region had 2 *N. meningitidis*–positive samples; both were nongroupable by PCR analysis.

### Discussion

In this study, 243 cases of laboratory-confirmed acute bacterial meningitis were recorded. Because our study centers provide service to 32% of the population of Turkey, extrapolation from the number of cases recorded suggests that 759 acute bacterial meningitis cases (excluding neonatal cases) per year occur in the whole country. The population of children 1 month through 16 years of age was calculated as 21.6 million. Therefore, the annual incidence of acute bacterial meningitis was estimated as 3.5 cases/100,000/year. Although similar to incidence rates reported from other countries without routine vaccination against *N. meningitidis*, *S. pneumoniae*, and Hib (3,4,10,30), this value likely represents a lower limit estimate of the true disease incidence, given the inherent limitations of hospital-based surveillance. Furthermore, the specific role of these 3 most common bacterial causes of acute bacterial meningitis varies between regions.

An accurate laboratory confirmation of the etiology in acute bacterial meningitis is essential to provide optimal patient therapy, appropriate case contact management, and reasoned public health actions. Prospectively, it also provides information upon which to base decisions regarding immunization programs, especially for countries without routine vaccination against the main acute bacterial meningitis pathogens (28,31). Although bacterial culture is considered to be the standard method, the negative effect of prior antimicrobial drug use on its sensitivity necessitates nonculture techniques for diagnosis. Among nonculture diagnostic tests, PCR is the most accurate and reliable method, especially among patients with a history of antimicrobial drug use before spinal tap (32). This finding was evident in the present study, in which PCR analysis was the most sensitive method, confirming 243 cases (59.6%) among 408 children meeting the case definition for bacterial meningitis (100% of all cases that were confirmed by any method). Using other methods that are more sensitive may increase the rate of laboratory confirmation.

Several reports review the rates of bacterial causes of acute bacterial meningitis from many different countries, based on CSF cultures. Some factors, such as previous antimicrobial drug treatment, interfere with the recovery of microorganisms from CSF. In our study, bacterial isolation was only possible in 41(16.8%) of 243 of confirmed cases. However, the most important factor for this low positive

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Although this vaccine generates a robust immune response against serogroup W-135, in contrast to what has been demonstrated for serogroup C meningococcal conjugate vaccines (37), meningococcal polysaccharide vaccines are not thought to reliably prevent asymptomatic carriage. A study from the United States reported that 0.8% of returning vaccinated pilgrims in 2001 were carrying W-135, whereas no pilgrims carried this serogroup upon departure from the United States (38). Therefore, the rapid rise in the proportion of cases caused by serogroup W-135 may be attributable to transmission from pilgrims returning from the Hajj carrying this particular serogroup. Although not definitive, this conclusion is further supported by the finding that all serogroup W-135 isolates available for phenotypic characterization were identical to the Hajj-associated clone, W135:2a:P1.5,2 (20). Because strains with this serologic profile have not been uniquely associated with the Hajj outbreak (39), additional typing data (e.g., multilocus sequence typing or multilocus enzyme electrophoresis), and epidemiologic investigations will be required to support this hypothesis. Therefore, it remains speculative that the increased W-135 disease in Turkey may be caused by spread of Hajj epidemic strain through carriage and transmission by pilgrims.

This study demonstrates the need for good quality, continued surveillance of bacterial meningitis cases, as well as the etiology and epidemiology of the causative bacteria. Only by accurately monitoring meningococcal epidemiology will effective vaccination policies be developed. The bacterial meningitis epidemiologic landscape is not static, and the causative agents change with time and across regions of the world. This study has demonstrated that the relative contribution of serogroup W-135 to the meningococcal disease incidence in Turkey is increasing, which is in contrast to the rest of Europe. Turkey may remain isolated in terms of W-135 disease incidence or it may represent the beginning of epidemiologic change in Eastern Europe. This possibility should be investigated in greater depth and monitored prospectively. Introduction of vaccines can dramatically reduce the meningitis disease incidence, but these vaccines must be targeted against the correct bacteria and, where relevant, the correct bacterial serogroup. The choice of meningococcal conjugate vaccines in Turkey will need to include coverage for serogroup W-135; introduction of such a vaccine would be helpful in protecting the Turkish population from this invasive bacterial meningitis. Moreover, it may also be prudent to switch the meningococcal vaccine used for pilgrims to conjugated vaccine to prevent the carriage of the microorganism by pilgrims.

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Dr Ceyhan is professor of pediatrics and pediatric infectious diseases at Hacettepe University, Ankara. His areas of interest

Figure 5. Etiology of confirmed cases of bacterial meningitis in different geographic regions. W-135 was the most prominent Neisseria meningitidis serogroup in the Southeast Anatolia, Aegean, Eastern Anatolia, and Black Sea regions. The percentages of cases caused by Streptococcus pneumoniae and Haemophilus influenzae type b (Hib) are also shown.
References


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