

High Prevalence of Antimicrobial Resistance among *Shigella* Isolates in the United States Tested by the National Antimicrobial Resistance Monitoring System from 1999 to 2002

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Shigella spp. infect approximately 450,000 persons annually in the United States, resulting in over 6,000 hospitalizations. Since 1999, the National Antimicrobial Resistance Monitoring System (NARMS) for Enteric Bacteria has tested every 10th *Shigella* isolate from 16 state or local public health laboratories for susceptibility to 15 antimicrobial agents. From 1999 to 2002, NARMS tested 1,604 isolates. Among 1,598 isolates identified to species level, 1,278 (80%) were *Shigella sonnei*, 295 (18%) were *Shigella flexneri*, 18 (1%) were *Shigella boydii*, and 7 (0.4%) were *Shigella dysenteriae*. Overall, 1,251 (78%) were resistant to ampicillin and 744 (46%) were resistant to trimethoprim-sulfamethoxazole (TMP-SMX). Prevalence of TMP-SMX- or ampicillin- and TMP-SMX-resistant *Shigella sonnei* isolates varied by geographic region, with lower rates in the South and Midwest regions (TMP-SMX resistance, 27% and 30%, respectively; ampicillin and TMP-SMX resistance, 25% and 22%, respectively) and higher rates in the East and West regions (TMP-SMX resistance, 66% and 80%, respectively; ampicillin and TMP-SMX resistance, 54% and 65%, respectively). Nineteen isolates (1%) were resistant to nalidixic acid (1% of *S. sonnei* and 2% of *S. flexneri* isolates); 12 (63%) of these isolates had decreased susceptibility to ciprofloxacin. One *S. flexneri* isolate was resistant to ciprofloxacin. All isolates were susceptible to ceftriaxone. Since 1986, resistance to ampicillin and TMP-SMX has dramatically increased. *Shigella* isolates in the United States remain susceptible to ciprofloxacin and ceftriaxone.

Shigellosis is an important cause of gastroenteritis, resulting in an estimated 450,000 cases in the United States each year (27). *Shigella* infections can lead to illness ranging from mild, self-limited diarrhea to severe dysentery with frequent passage of blood and mucus, high fever, cramps, tenesmus, and in rare cases, bacteremia. Complications of shigellosis are seen most frequently in children, the elderly, and the immunocompromised. Prompt treatment with effective antimicrobial agents may shorten the duration of clinical symptoms and carriage and reduce the spread of infection (36).

Antimicrobial resistance has complicated the selection of empirical agents for treatment of shigellosis, particularly in children. When the prevalence of resistance to ampicillin among *Shigella* isolates increased in the 1970s, trimethoprim-sulfamethoxazole (TMP-SMX) became the alternative (31). In 1986, national laboratory-based surveillance for antimicrobial susceptibility of *Shigella* isolates revealed that 32% were resistant to ampicillin and 7% were resistant to TMP-SMX (46). In 1995, laboratory-based surveillance demonstrated resistance to ampicillin in 67% and resistance to TMP-SMX in 35% of *Shigella* isolates (15). More recent data from Oregon revealed high rates of TMP-SMX (63%), ampicillin (59%), and multidrug (13%) resistance among *Shigella* isolates from this state (35).

National surveillance for antimicrobial resistance among *Shigella* isolates began in 1999 as part of the National Antimicrobial Resistance Monitoring System (NARMS) for Enteric Bacteria (10). To describe trends in antimicrobial resistance among *Shigella* isolates from 1999 through 2002, we analyzed NARMS data for the first 4 years of surveillance. Our findings indicate an increase in the rates of resistance of *Shigella* isolates to ampicillin and TMP-SMX and evidence of emerging resistance to nalidixic acid.

MATERIALS AND METHODS

Surveillance and antimicrobial susceptibility testing. NARMS was established in 1996 to monitor antimicrobial resistance of human nontyphoid *Salmonella* and *Escherichia coli* O157 isolates (10). *Shigella* was added to NARMS surveillance in 1999, and annual reports are available electronically (10). Public health laboratories in 15 states (Colorado, Connecticut, Florida, Georgia, Kansas, Maryland, Massachusetts, Minnesota, New Jersey, New York, Oregon, Tennessee, Washington, and West Virginia) and two local health departments (Los Angeles County and New York City) participated in *Shigella* surveillance in NARMS. In all sites, participating public health laboratories were requested to send every 10th *Shigella* isolate received from clinical laboratories to the Centers for Disease Control and Prevention (CDC) for antimicrobial susceptibility testing. Duplicate isolates from the same patient are not routinely excluded by laboratories; however, duplicate submission of *Shigella* isolates is rarely identified at the CDC. Participating public health laboratories reported the patient's age, date of birth, sex, and county of residence, the specimen source (stool, blood, or other), the date of isolate collection, and the date of isolate receipt at the state public health laboratories. For our analysis, we combined isolates from New York City and New York State as New York, for a total of 16 sites. Postcensus estimates for the population in these 16 sites in 2000 was 105 million persons, or 37% of the U.S. population in 2000 (48).

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Shigella isolates were tested at the CDC with a semiautomated system (Sensititre; Trek Diagnostics, Westlake, OH) to determine the MIC ranges of 15 antimicrobial agents: amikacin, ampicillin, amoxicillin-clavulanic acid, ceftiofur, ceftriaxone, cephalothin, chloramphenicol, ciprofloxacin, gentamicin, kanamycin, nalidixic acid, streptomycin, sulfamethoxazole, tetracycline, and trimethoprim-sulfamethoxazole. Ceftiofur is the only expanded-spectrum cephalosporin approved for systemic use in food animals in the United States and is included in NARMS because the use of this antimicrobial in food animals has been implicated as a factor responsible for the emergence of ceftriaxone-resistant enteric pathogens, such as *Salmonella* (16). This has not been documented for *Shigella*. MIC results were dichotomized (isolates with intermediate susceptibility were categorized as sensitive), and CLSI (formerly NCCLS) criteria were used when applicable (14). Ceftiofur resistance was defined as a MIC of ≥ 8 $\mu\text{g/ml}$ based on population distributions of MICs for gram-negative isolates. Decreased susceptibility to ciprofloxacin was defined as a MIC of ≥ 0.125 $\mu\text{g/ml}$.

Statistical analysis. Statistical analysis focused on the relationships between antimicrobial resistance patterns and *Shigella* serotype, geographic location, and age and sex of the patient. We defined a pansensitive isolate as one that was sensitive to all antimicrobial agents included in the analysis. We defined multidrug resistance as resistance to two or more classes of antimicrobial agents (tested separately) among all classes included in the analysis. To calculate incidence rates by geographic region, we grouped participating sites into four regions: West (Los Angeles County, Oregon, and Washington), Midwest (Colorado, Kansas, and Minnesota), South (Florida, Georgia, Maryland, Tennessee, and West Virginia), and Northeast (Connecticut, Massachusetts, New York, and New Jersey). Region-specific rates of isolation per 100,000 population were determined for the study period with 2000 census data from the U.S. Bureau of the Census (48).

We compared the number of *Shigella* isolates submitted to NARMS from 1999 to 2002 to the number of laboratory-confirmed *Shigella* infections reported to the CDC via the Public Health Laboratory Information System (PHLIS) (11). Unlike the NARMS surveillance system, PHLIS is used by public health laboratory directors and state and territorial epidemiologists from every state in the United States to report all laboratory-confirmed *Shigella* isolates. Therefore, the comparison with PHLIS provides information about the generalizability of the data gathered through NARMS. In addition, we compared the proportion of *Shigella* isolates resistant to TMP-SMX and ampicillin found in our study to two previous national laboratory-based surveillance studies to examine trends in resistance to these agents (15, 46).

Statistical analyses were conducted using SAS version 8 (SAS Institute Inc., Cary, NC) statistical software. We used χ^2 tests to compare proportions and Fisher's exact test when appropriate. Continuous variables were compared by the Wilcoxon rank-sum test. *P* values are based on two-tailed test results, and *P* values of <0.05 were considered statistically significant.

RESULTS

During the 4-year period 1999 to 2002, the CDC received and tested 1,604 *Shigella* isolates from participating NARMS sites included in this analysis. This represents approximately 9% of 17,432 culture-confirmed cases reported to PHLIS by these sites. Among the 1,604 isolates, 275 (17%) were from New York (of which 192 [70%] were from New York City), 197 (12%) were from New Jersey, 177 (11%) were from Georgia, 166 (10%) were from Minnesota, 147 (9%) were from Massachusetts, and 642 (40%) were from other sites. Among 1,598 isolates identified to species level, 1,278 (80%) were *Shigella sonnei* (compared with approximately 82% reported through PHLIS for this time period and from these sites), 295 (18%) were *Shigella flexneri* (versus 16%), 18 (1%) were *Shigella boydii* (versus 0.7%), and 7 (0.4%) were *Shigella dysenteriae* (versus 0.4%). Most *Shigella* isolates were isolated from specimens collected from stool (97%).

Geographic distribution. During the 4-year study period, the number of *Shigella* isolates received per 100,000 population was 1.75 in the Northeast, 1.51 in the South, 2.30 in the Midwest, and 1.10 in the West. *S. sonnei* accounted for 76% of all *Shigella* isolates from the Northeast, 89% from the South, 87% from the Midwest, and 62% from the West. *S. flexneri* accounted for 22%

of all *Shigella* isolates from the Northeast, 10% from the South, 12% from the Midwest, and 35% from the West. *S. boydii* accounted for 1% of isolates from the Northeast, 1% from the South, 0.7% from the Midwest, and 2% from the West. *S. dysenteriae* accounted for 1% of isolates from the Northeast and 0% from the South, Midwest, and West.

Age and gender. Information on patient age was available for 1,442 (90%) of 1,598 isolates identified to species level that were received during the study period. The median age was 8 years (range, <1 year to 99 years), and 51% were male. Infants and children from 1 to 4 years of age accounted for the highest proportion of *S. sonnei* (32%) and *S. flexneri* (29%) isolates submitted to NARMS. In the PHLIS data reported in 2002, infants and children aged 1 to 4 years accounted for a similarly high proportion of infections with *S. sonnei* (33%) and *S. flexneri* (26%) (11). Consistent with PHLIS data, persons from whom *S. sonnei* isolates were submitted to NARMS were younger than those with *S. flexneri* (median age, 8 years versus 12 years; $P = 0.03$). Overall, more *S. flexneri* isolates in NARMS were from males than females (20% versus 17%; $P = 0.09$), particularly among persons 20 to 49 years of age (31% versus 16%; $P < 0.01$).

Seasonality. Among the 1,278 *S. sonnei* isolates, 233 (18%) were isolated in the winter quarter (December, January, and February), 282 (22%) in the spring (March, April, and May), 412 (32%) in the summer (June, July, and August), and 351 (27%) in the fall (September, October, and November). Among *S. flexneri* isolates, similar proportions of isolates (24% to 27%) were isolated in each quarter.

Antimicrobial resistance. Among the 1,604 *Shigella* isolates tested, 115 (7%) were pansusceptible (97 [8%] of 1,278 *S. sonnei*, 14 [5%] of 295 *S. flexneri*, 3 [17%] of 18 *S. boydii*, and 1 [14%] of 7 *S. dysenteriae* isolates). Large proportions of isolates were resistant to ampicillin (1,251 [78%]), streptomycin (891 [56%]), sulfamethoxazole (757 [47%]), and TMP-SMX (744 [46%]). Resistance to both ampicillin and TMP-SMX was seen in 613 (38%) of isolates (Table 1).

Similarly high proportions of *S. sonnei* and *S. flexneri* isolates were resistant to ampicillin (79% and 76%; $P = 0.18$), but resistance to TMP-SMX was more common among *S. sonnei* than among *S. flexneri* isolates (48% versus 40%; $P = 0.02$) (Table 1). Resistance to both ampicillin and TMP-SMX was similar among *S. sonnei* and *S. flexneri* isolates (39% and 35%; $P = 0.19$). However, resistance to chloramphenicol was common among *S. flexneri* isolates and extremely rare among *S. sonnei* isolates (70% versus 1%; $P < 0.01$).

None of the *Shigella* isolates tested were resistant or had decreased susceptibility to ceftriaxone (MIC ≥ 4 $\mu\text{g/ml}$). Only one isolate (*S. flexneri*) was resistant to ciprofloxacin; it was obtained from a child who recently traveled to China (7). In addition, 19 (1%) isolates (13 *S. sonnei*, 5 *S. flexneri*, and 1 *S. boydii* isolate) were resistant to nalidixic acid; 12 (63%) of these isolates also had decreased susceptibility (MIC > 0.125 $\mu\text{g/ml}$) to ciprofloxacin.

Among the 1,604 isolates tested, 1,031 (64%) were resistant to two or more agents (multidrug resistant). Common resistance patterns included the combination of ampicillin, streptomycin, sulfamethoxazole, and tetracycline resistance (322 [31%]), ampicillin, streptomycin, and sulfamethoxazole resistance (149 [14%]), and ampicillin and streptomycin resistance

TABLE 1. Antimicrobial resistance of *Shigella* isolates by species, United States, 1999 to 2002

Antimicrobial agent or characteristic	Breakpoint(s) (R)	% Resistant					
		All isolates (n = 1,604)	<i>S. sonnei</i> (n = 1,278)	<i>S. flexneri</i> (n = 295)	<i>S. boydii</i> (n = 18)	<i>S. dysenteriae</i> (n = 7)	Unknown (n = 6)
Amikacin	≥64	0.06	0.08	0	0	0	0
Amoxicillin-clavulanic acid	≥32/16	2	2	5	0	0	0
Ampicillin	≥32	78	79	76	0	71	100
Ceftiofur ^a	≥8	0.06	0	0.3	0	0	0
Ceftriaxone	≥64	0	0	0	0	0	0
Cephalothin	≥32	6	7	3	0	0	0
Chloramphenicol	≥32	14	1	70	0	43	67
Ciprofloxacin	≥4	0.06	0	0.3	0	0	0
Gentamicin	≥16	0.2	0.2	0.3	0	0	0
Kanamycin	≥64	0.9	0.8	1	0	0	0
Nalidixic acid	≥32	1	1	2	6	0	0
Streptomycin ^a	≥64	56	55	55	72	71	83
Sulfamethoxazole	≥512	47	45	54	78	71	50
Tetracycline	≥16	45	33	92	50	71	83
TMP-SMX	≥4/76	46	48	40	39	57	50
Ampicillin and TMP-SMX	≥32 and ≥4/76	38	39	35	0	43	50
% Pansusceptible		7	8	5	17	14	0
% Multiresistant ^b		64	59	85	83	71	100

^a No CLSI interpretive standards for this antimicrobial agent. The MIC for ceftiofur was based on population distributions of MICs for gram-negative isolates. The MIC of streptomycin is based on the Canadian Integrated Program for Antimicrobial Resistance (19).

^b Defined as resistant to two or more antimicrobial agents.

(108 [10%]). Resistance to the combination of ampicillin, chloramphenicol, streptomycin, sulfamethoxazole, and tetracycline was seen in 85 (8%) isolates.

The prevalence of TMP-SMX- or ampicillin- and TMP-SMX-resistant *Shigella sonnei* isolates varied by geographic region (Fig. 1). Among *S. sonnei* isolates, 66% of 493 in the Northeast, 27% of 421 in the South, 30% of 238 in the Midwest, and 80% of 126 in the West were resistant to TMP-SMX. Among *S. sonnei* isolates, 54% of 493 in the Northeast, 25% of 421 in the South, 22% of 238 in the Midwest, and 65% of 126 in the West were resistant to both ampicillin and TMP-SMX. No geographic variability was found among *S. flexneri* isolates; 47% of 142 in the Northeast, 37% of 46 in the South, 44% of 34 in the Midwest, and 29% of 73 in the West were resistant to TMP-SMX. Among *S. flexneri* isolates, 41% of 142 in the Northeast, 35% of 46 in the South, 38% of 34 in the Midwest, and 23% of 73 in the West were resistant to both ampicillin and TMP-SMX.

Resistance rates among *S. sonnei* isolates differed between sexes. Specifically, *S. sonnei* isolates from males were more likely to be resistant to TMP-SMX or to both ampicillin and TMP-SMX than *S. sonnei* isolates from females (53% versus 45% resistance to TMP-SMX, *P* = 0.01, and 45% versus 36% resistance to ampicillin and TMP-SMX, *P* < 0.01, respectively). There were no significant sex differences in infection with TMP-SMX-resistant *S. flexneri* in males and females (42% and 39%; *P* = 0.58) or with ampicillin- and TMP-SMX-resistant *S. flexneri* in males and females (38% and 33%; *P* = 0.43). Persons infected with TMP-SMX-resistant *S. sonnei* were significantly older than persons infected with TMP-SMX-susceptible strains (9 years versus 7 years; *P* < 0.01). Otherwise, there were no statistically significant differences in median age between resistant and susceptible strains of *S. sonnei* or *S. flexneri*. The relatively greater prevalence of antimicrobial-resistant *S. sonnei* in males was observed in each of the four regions.

Overall, the proportion of *Shigella* isolates resistant to am-

picillin, TMP-SMX, and both agents increased since national surveys were conducted in 1986 and 1995. Over this 16-year period, the proportion resistant to ampicillin increased from 32% to 67% to 78%, the proportion resistant to TMP-SMX

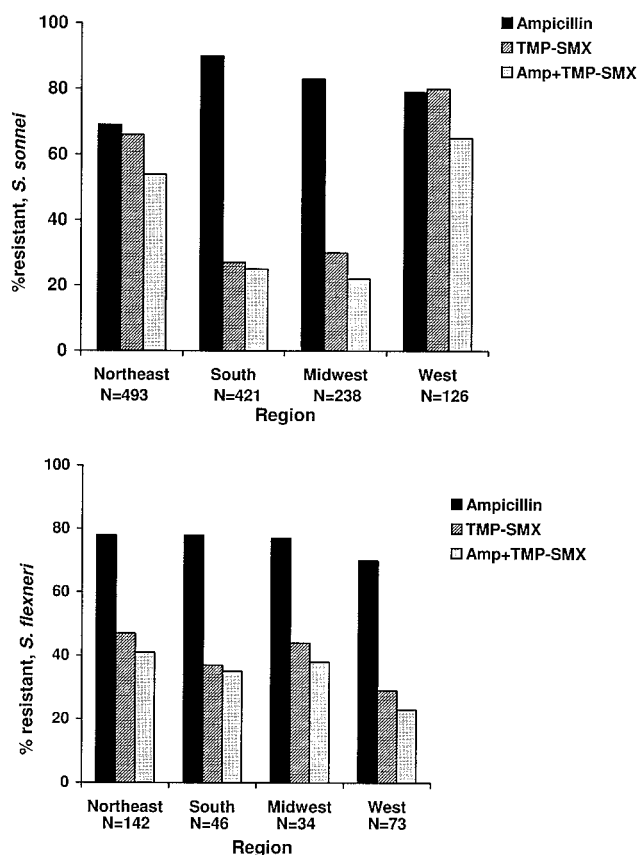


FIG. 1. Percentage of *S. sonnei* and *S. flexneri* isolates resistant to TMP-SMX, ampicillin (Amp), or both by region from 1999 to 2002.

increased from 7% to 35% to 46%, and the proportion resistant to both agents increased from 6% to 19% to 38% (15, 46).

DISCUSSION

Antimicrobial resistance among *Shigella* spp. is common in the United States, and between 1999 and 2002, only 7% of isolates reported through NARMS were pansensitive. Resistance to TMP-SMX or to both ampicillin and TMP-SMX was more common among *S. sonnei* than *S. flexneri* isolates, while resistance to chloramphenicol was more common among *S. flexneri* isolates, findings consistent with previous surveillance studies (15, 46). The proportion of isolates resistant to ampicillin, TMP-SMX, or both has increased substantially in the last decade, and these agents are no longer appropriate for empirical treatment in most parts of the country. The comparison of NARMS data with national laboratory-based surveillance data collected through PHLIS for the same time period suggests that NARMS captures a demographically representative sample of the total population of individuals infected with *Shigella* in the United States.

Few *Shigella* isolates were resistant to nalidixic acid ($n = 19$) or ciprofloxacin ($n = 1$), and none was resistant to ceftriaxone. While these agents may offer reliable results when used for empirical treatment of shigellosis, clinicians and public health officials should anticipate increasing resistance to them as their use increases. Clinical experience also supports the use of macrolides, such as azithromycin, for therapy of shigellosis; however, NARMS does not currently include data on susceptibility to any representative of this class of antimicrobial agents (24).

A majority of isolates that were resistant to nalidixic acid also had decreased susceptibility to ciprofloxacin. Although clinical failures to treat nalidixic acid-resistant *Shigella* infection with fluoroquinolones have not been reported, clinicians should be aware that failures have been reported for *Salmonella* infections caused by strains that were resistant to nalidixic acid and had decreased susceptibility to ciprofloxacin (13, 40).

There were regional differences in the prevalence of TMP-SMX resistance among *Shigella sonnei* isolates. TMP-SMX resistance was found in 30% of *S. sonnei* isolates in the Midwest NARMS sites compared to 66% and 80% of *S. sonnei* isolates from the Northeast and West, respectively. A similar trend was seen for isolates of *S. sonnei* resistant to both ampicillin and TMP-SMX. The geographic differences in resistance patterns may be a result of clonal spread of isolates. One study conducted during an outbreak of shigellosis in a day-care center revealed that pulsed-field gel electrophoresis and antimicrobial susceptibility patterns were identical in all outbreak strains, while nonoutbreak strains isolated during the outbreak period had indistinguishable pulsed-field gel electrophoresis patterns but different antimicrobial susceptibility patterns. Therefore, the occurrence of outbreaks of shigellosis in areas captured by NARMS sites in the Northeast or West regions could account for the geographic differences in resistance patterns observed. Because molecular typing of organisms is not routinely performed by NARMS, we cannot exclude this possibility.

Our study confirms that infection with *S. sonnei* is particularly common among young children and among women aged

20 to 40 years, while *S. flexneri* infections occur more frequently in adult men aged 30 to 49 years. The higher median age among persons infected with *S. flexneri* and the relatively greater proportion of *S. flexneri* isolates in adult men may reflect the occurrence of sexually transmitted shigellosis in the population of men who have sex with men (MSM), although outbreaks of *S. sonnei* infections have also recently been reported in this community (12, 45). Our data indicate that persons with TMP-SMX-resistant *S. sonnei* infection were more likely to be men and were older than persons with TMP-SMX-susceptible infections. The recently reported outbreak of *S. sonnei* among MSM in California was caused by a TMP-SMX-resistant strain. Furthermore, a high rate of human immunodeficiency virus infection has been reported among MSM with *S. sonnei* infection in the Bay Area (6). The widespread use of TMP-SMX for prophylaxis of pneumocystis pneumonia among MSM may predispose this population of adult men to TMP-SMX-resistant *S. sonnei* infections. An extensive outbreak of TMP-SMX-sensitive *S. sonnei* infection among day-care center attendees in the mid-Atlantic regions from 2001 to 2003 may also have contributed to the relatively lower percentage of TMP-SMX resistance among *S. sonnei* isolates from the South and Midwest and to the lower median age of persons infected with TMP-SMX-susceptible strains (9).

Transmission of shigellosis is primarily fecal-oral, and no *Shigella* vaccines are available in the United States. Because it takes only few organisms to transmit the infection, hand washing has been promoted as the single most important control measure to reduce the spread of shigellosis (23) and is especially critical in limiting the spread of shigellosis among young children in day-care centers (28). Many state health departments require exclusion of food handlers, health-care workers, child-care providers, and children who attend day care centers while they are symptomatic and until one or more negative stool cultures have been obtained (41). In these situations, appropriate antimicrobial use can greatly reduce the inconvenience and cost incurred during outbreaks, since most persons will cease to excrete *Shigella* within 72 h of starting appropriate antimicrobial therapy compared with carriage of up to several weeks that can occur without therapy (26).

The high prevalence of antimicrobial resistance among *Shigella* isolates noted in this study limits safe and efficacious treatment options for shigellosis, particularly for children. Where resistance to ampicillin and TMP-SMX is common, appropriate antimicrobial agents for the treatment of shigellosis are limited to nalidixic acid or fluoroquinolones, ceftriaxone, or azithromycin (18, 24, 42). Nalidixic acid (55 mg/kg/24 h in four divided doses for 5 days), a narrow-spectrum quinolone, is effective and approved in the United States for treatment of children older than 3 months (18). The broader-spectrum fluoroquinolones, while effective in treating shigellosis in children, are not approved by the U.S. Food and Drug Administration for use in children aged 18 years or younger because some fluoroquinolones have been shown to cause cartilage damage in juvenile animals (8). However, ciprofloxacin, the fluoroquinolone most extensively studied in children, has been successfully used to treat acute invasive diarrhea in children without the development of joint abnormalities (25, 37). Fluoroquinolones may be justified in children, after risks and benefits of treatment are discussed with parents, when no other

oral agent is available and in cases of severe shigellosis caused by a multidrug-resistant strain (4).

Mobile genetic units, including plasmids, gene cassettes in integrons, and transposons, are important in the spread of resistant determinants among *Shigella* isolates (32, 47). Trimethoprim and sulfamethoxazole resistance is most commonly acquired through a plasmid-encoded variant of the dihydrofolate reductase enzyme (20). Ampicillin resistance arises as a result of beta-lactamases similar to TEM-1 or OXA-1, whose genes may be located on chromosomes, plasmids, or transposons (30). Although resistance to quinolones is commonly mediated through chromosomal mutations rather than mobile genetic units, certain plasmids have been shown to contribute to quinolone resistance by increasing the rate of spontaneous mutation (3, 5).

Reports of antimicrobial resistance trends in *Shigella* isolates from other countries raise the specter of wider resistance to nalidixic acid, fluoroquinolones, and ceftriaxone in the future.

A recent report from the United Kingdom revealed a 13% nalidixic acid resistance among *Shigella sonnei* isolates, with all isolates also exhibiting decreased susceptibility to ciprofloxacin (MIC, 0.25 to 1.0 mg/liter) (13). A study from Japan reports a 26% prevalence of nalidixic acid resistance among *Shigella sonnei* isolates (21). Fluoroquinolone resistance in *S. dysenteriae* isolates has been reported recently from Bangladesh, India, and Nepal (29, 38, 43, 44). Finally, with the gradual increase in extended-spectrum beta-lactamases detected in *Klebsiella* and other *Enterobacteriaceae*, an increase in extended-spectrum beta-lactamase-producing *Shigella* strains has been reported since 1999 (2). Ceftriaxone-resistant strains of *S. sonnei* and *S. flexneri* have been reported from Korea, Argentina, France, Turkey, and Taiwan (1, 17, 22, 33, 34). The beta-lactamases described in these reports included CTX-M-14, CTX-M-2, SHV-2, CTX-M-3, and CMY-2 type AmpC.

Our study has several limitations. Laboratory-based surveillance systems, such as NARMS, may overestimate the prevalence of resistance among *Shigella* isolates because persons infected with resistant strains who fail empirical treatment are more likely to get a stool culture for continued diarrhea than are persons infected with susceptible strains who respond to empirical therapy. In addition, NARMS sites include a non-weighted 37% sample of the U.S. population, limiting generalizability of the data. Nonetheless, there was good agreement between demographic data for patients captured by NARMS and the national *Shigella* laboratory-based surveillance system reported to PHLIS. This high degree of correlation suggests that NARMS surveillance captures a representative sample of all *Shigella* infections in the United States and that generalizations from NARMS data on antimicrobial resistance among *Shigella* isolates may be valid despite the fact that NARMS captures only every 10th isolate of *Shigella* received by participating public health laboratories. In 2003, NARMS surveillance was extended to all 50 state health department laboratories. Finally, interpretation of antimicrobial susceptibility breakpoints without also considering pharmacokinetics and pharmacodynamics of the drug is not straightforward. For example, in vitro resistance testing using axenic medium may reveal that extracellular *Shigella* isolates are susceptible to aminoglycosides. However, in vivo, the same *Shigella* isolate, which is a facultative intracellular pathogen, may be resistant to ami-

noglycosides due to their inability to permeate mammalian cells (39, 49). The CLSI recommends that susceptibility of *Shigella* to aminoglycosides not be reported to clinicians as "sensitive" because of this discrepancy (14).

Treatment for shigellosis is critical in persons who have severe disease, especially in children and the immunosuppressed. The small but evident increase in the proportion of *Shigella* isolates that are resistant to nalidixic acid may portend the loss of an important class of antimicrobial agents against *Shigella*. The continued monitoring of emerging resistance in *Shigella* isolates through NARMS will be essential to timely and appropriate recommendations for antimicrobial therapy.

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