

## Original Article

# Prevalence of Ciprofloxacin and Nalidixic Acid Resistant *Salmonella enterica* serovar Typhi in Bangladesh

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A total of 1,059 *Salmonella enterica* serovar Typhi were isolated from blood samples during January 2006 to October 2007 from urban rural facilities in Dhaka, Bangladesh, of which 980 (92.5%) isolates were nalidixic acid resistant. The minimum inhibitory concentrations (MIC) of ciprofloxacin (CIP) were determined for 127 nalidixic acid resistant *Salmonella enterica* serovar Typhi (NARST) strains (every fifth) isolated during 2006. Nine isolates were found to be resistant against CIP (3%) with high MIC (12 - >32 µg/mL). Only four isolates were found to be sensitive (MIC <0.125 µg/mL), whereas most of the isolates (N=113) showed reduced susceptibility (MIC 0.125 – 2 µg/mL) to CIP. All these isolates were subjected to molecular typing by multiplex PCR on VNTR (variable number tandem repeats) loci, which revealed eight different VNTR patterns. Almost all CIP resistant strains had similar genetic organization, identical to the most common VNTR type. Restriction fragment length polymorphism (RFLP) analysis of the *gyrA* gene revealed point mutations at Ser-83 and Asp-87 in all CIP resistant strains.

**Key words:** *Salmonella enterica* serovar Typhi, fluoroquinolone resistant, VNTR

## Introduction

Diseases caused by *Salmonella enterica* serovars are especially prevalent in developing countries. Typhoid fever is sometimes a fatal infection to adults and children that causes bacteremia and inflammatory destruction of the intestine and other organs. This requires an urgent treatment by the administration of appropriate antibiotics. Emergence of multidrug-resistant (MDR) *S. enterica* serovar Typhi strains in Asia in the beginning of 1990s, led to the widespread use of fluoroquinolones for treating enteric fever<sup>1-2</sup>. MDR typhoid fever endemic started in Bangladesh in 1990s, which reached to peak in 1994, and then declined and re-emerged in 2001 and 2002<sup>3</sup>. During the last decade, several treatment failures of *S. typhi* strains with decreased susceptibility to ciprofloxacin<sup>3-5</sup> have been reported and some studies also confirmed the presence of fluoroquinolone resistant *S. enterica* serovar Typhi strains<sup>6-8</sup>. Antibiotic-resistant *Salmonella* strains pose a significant threat to the development of reliable therapies. *S. enterica* serovar Typhi with resistance to nalidixic acid and decreased susceptibilities or resistance to fluoroquinolones have been increasingly reported in several countries including Bangladesh<sup>7, 9-13</sup>. Such strains have also been reported from other parts of the world, most of them have travel history to Asian countries particularly South East Asia<sup>6,14</sup>.

The mechanisms of fluoroquinolone resistance have been well studied. Single point mutation in the quinolone resistance-

determining region (QRDR) of the *gyrA* gene in *Salmonella* usually leads to resistance against nalidixic acid and to decreased susceptibility to ciprofloxacin<sup>7, 15-16</sup>. Turner *et al.*<sup>17</sup> reported that two substitutions in *gyrA* (Ser-83 → Phe or Tyr and Asp-87 → Asn) and one in *parC* (Glu-84 → Lys) confer complete fluoroquinolone resistance in *S. typhi*. Emergence of these resistant strains are associated with treatment failure or delayed response to ciprofloxacin<sup>9,18</sup>.

There are few reports on the occurrence of typhoid by absolute fluoroquinolone resistant strain in Bangladesh. Here, we report nine cases of typhoid fever in Dhaka, Bangladesh caused by *S. enterica* serovar Typhi with complete resistance to ciprofloxacin during 2006.

## Materials and Methods

### *Specimen collection and identification of Salmonella enterica serovar Typhi*

The patients were selected for specimen collection based on the clinical manifestation of typhoid fever diagnosed by physician from different sites at Dhaka City, Bangladesh. Blood samples were collected along with patient's demographic data from all age group of typhoid patients during 2006 to October 2007, as a part of routine diagnosis.

Blood cultures were performed by the lysis centrifugation method as described in our previous report<sup>19</sup>. Positive cultures were

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subsequently plated on blood, chocolate and MacConkey agar plates. Colonies of *Salmonella enterica* serovar Typhi were identified by standard biochemical and serological procedures<sup>20</sup>.

#### Determination of antibiotic sensitivity and MIC of ciprofloxacin

Antibiotic susceptibility testing was performed by disk diffusion method using discs (Oxoid, UK) containing ampicillin (10 µg), ceftriaxone (30 µg), cotrimoxazole (25 µg), chloramphenicol (30 µg), ceftazidime (30 µg), ciprofloxacin (5 µg) and nalidixic acid (30 µg). Based on the date of isolation, every fifth nalidixic acid resistant *S. enterica* serovar Typhi (NARST) isolated during the year 2006 were selected to determine the MIC of ciprofloxacin (CIP) by E-strip. All results were interpreted according to Clinical and Laboratory Standards Institute (CLSI) guidelines<sup>21</sup> and were analysed by SPSS version 11.5. The trend of multidrug resistance, which is defined as resistance to ampicillin, chloramphenicol and co-trimoxazole was also analysed during 2006 to October 2007.

#### Multiplex PCR on VNTR loci

All the isolates whose MIC was determined by E-strip test were subjected to molecular typing by multiplex PCR on variable number tandem repeats (VNTR) using primers flanking three VNTR loci (TR1, TR2 and TR3) (Table 1), as described by Liu et al.<sup>22</sup>. In brief, each 25 µL of reaction mixture contained 1.5 µL of the bacterial lysate suspension and 10 pmol each of the forward and reverse primers for TR1 and TR3, as well as 12.5 pmol each of the corresponding primers for TR2, in addition to 22 µL of the *Taq* PCR mastermix (QIAGEN GmbH, Hilden, Germany). After initial denaturation at 94°C for 5 min, the PCR reaction was run for 35 cycles at 94°C for 30 s, 55°C for 30 s, and 72°C for 1 min, followed by a final extension at 72°C for 7 min. The PCR products, along with a 100-bp DNA marker (Invitrogen, USA), were subjected to electrophoresis on a 2% agarose gel. In every case, positive and negative controls were run simultaneously.

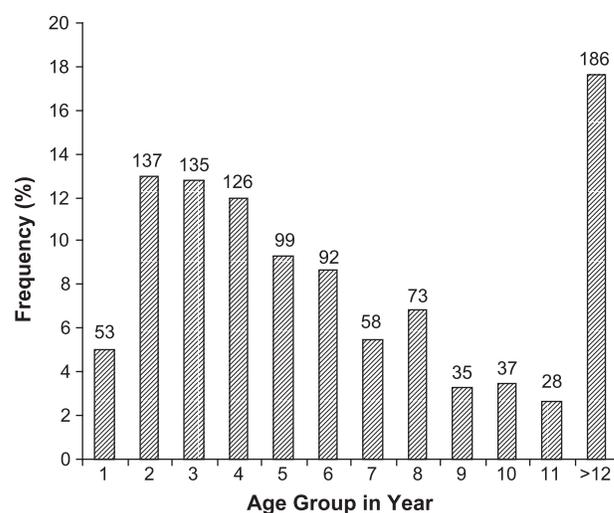
#### Restriction fragment length polymorphism (RFLP) analysis

PCR-RFLP was performed to detect common mutations related to fluoroquinolone resistance at the codon Ser-83 and Asp-87 of *gyrA* following the procedure described by Hirose et al.<sup>23</sup>. PCR was performed with the primers, *gyrA*-F, 5' TGT CCG AGA TGG CCT GAA GC 3' and *gyrA*-*HinfI*-as, 5' ATG TAA CGC AGC GAG AAT GGC TGC GCC ATA CGA ACG CTG GAG 3'. The 195-bp

amplified product contains two *HinfI* restriction sites at the codon corresponding to Ser-83 and Asp-87 of *gyrA*. Restriction digestions were performed in a total of 15 µL reaction mixture containing 8 µL of the above-mentioned PCR product, 1.5 µL of 10x digestion buffer and 10 U of *HinfI* for overnight at 37°C. The restriction digests were electrophoresed onto a 3% agarose gel. *HinfI*-digested PCR products of *gyrA* gene from a known nalidixic acid sensitive *S. enterica* serovar Typhi strain, and the undigested product from known ciprofloxacin resistant strain, were run as controls.

#### Results

During the period of the study (2006 to October 2007), a total of 1,059 *Salmonella enterica* serovar Typhi were isolated. The majority (57%; 605 out of 1059) of the *S. enterica* isolates were isolated in the year 2006. A significant higher proportion of cases (43%) occurred in patients younger than 5 years of age than in those 5 years of age (9%) or older. The highest rate (13%) of *S. enterica* serovar Typhi isolation was in the second year of life (Fig. 1).



**Figure 1.** Age distribution of typhoid fever cases caused by *S. enterica* serovar Typhi identified at a diagnostic referral center and Dhaka Shishu Hospital, Bangladesh during 2006-2007 ( $n = 1059$ ). Numbers above the bars show the total numbers of isolates

**Table 1.** PCR primers flanking the three VNTR loci of *S. enterica* serovar Typhi<sup>22</sup>.

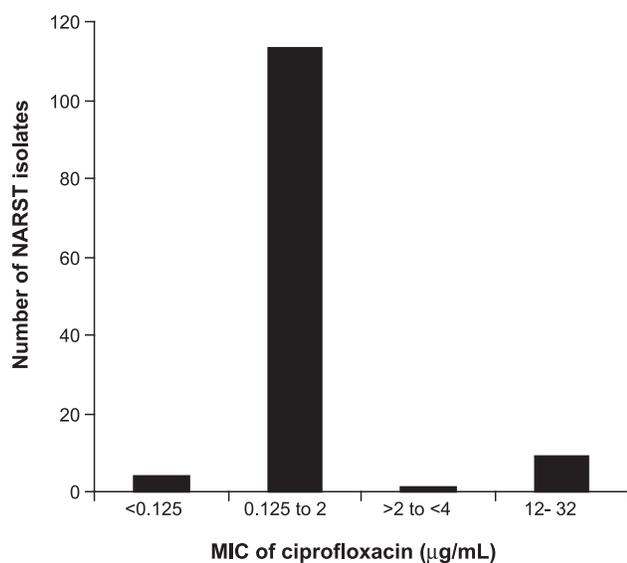
Primer	Nucleotide positions	Sequence (5'-3')	VNTR sequence
TR1F1	2017115-2017136	AGA ACC AGC AAT GCG CCA ACG A	(AGAAGAA) <sub>12</sub>
TR1R1	2017354-2017375	CAA GAA GTG CGC ATA CTA CAC C	
TR2F1	2556810-2556831	CCC TGT TTT TCG TGC TGA TAC G	(CCAGTTCC) <sub>27</sub>
TR2R2	2557299-2557320	CAG AGG ATA TCG CAA CAA TCG G	
TR3F1	2926145-2926166	CGA AGG CGG AAA AAA CGT CCT G	(CGCGGGGATCGGTTT ATCCCCGCTGG) <sub>3,3</sub>
TR3R1	2926668-2926689	TGC GATTGG TGT CGT TTC TAC C	

Antibiotic susceptibility patterns of 1,059 *S. enterica* serovar Typhi were analysed for seven antibiotics. Among them only 52 (4.9%) isolates were sensitive to all antimicrobial agents tested. Although no resistance to ceftriaxone (CRO) or ceftazidime (CAZ) was observed, 92.5% isolates were nalidixic acid (NA) resistant. Multidrug resistance (MDR) was noted in around 56% (593 out of 1059) of the isolates. Moreover, MDR strains were more likely to be resistant to NA, in comparison to sensitive strains (96.7% vs 3.3%).

The results of molecular typing and MIC for ciprofloxacin are presented in Table 2 and Figure 2 respectively. Although not considered resistance by current CLSI standard breakpoints, 89% (113 out of 127) isolates demonstrated a “reduced susceptibility” to ciprofloxacin (MIC value 0.125 to 2 µg/mL) and 9 isolates were completely resistant to CIP (MIC value 12 - >32 µg/mL) (Fig. 2).

**Table 2.** Different VNTR types and their corresponding amplicon size.

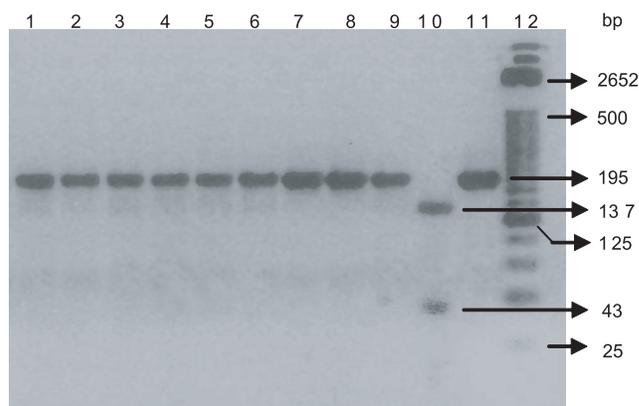
VNTR Type	Number of isolates	Amplicon size (bp)
A	93	~200, >600
B	14	> 200, > 600
C	12	>300, ~500
D	2	~150, >200, ~600
E	2	>250, >300, ~600
F	2	>250, ~600
G	1	200, < 400, > 600
H	1	>200



**Figure 2.** MIC of ciprofloxacin of some nalidixic acid resistant *Salmonella enterica* serovar Typhi ( $n = 127$ ) isolated in 2006. MIC value <0.125 µg/mL, Sensitive; MIC value 0.125 to 2 µg/ml, decreased susceptibility; MIC value >2 to <4 µg/mL, intermediate; MIC value 12 - >32 µg/mL, resistant

All the NARST isolates were subjected to molecular typing on VNTR loci through multiplex PCR. A total of 8 VNTR types (type A to type H) based on the number and size of bands were identified among 127 isolates (Table 2). Most of the isolates 73% (93/127) belonged to the VNTR type A and eight of the nine ciprofloxacin resistant isolates were also included in this group.

The fragments containing the *gyrA* QRDR of nine ciprofloxacin resistant isolates were amplified by PCR. The 195 bp PCR products of CIP resistant strains were not digested with *HinfI* (Fig. 3), suggesting point mutations at both Ser-83 and Asp-87 of *gyrA*. In contrast, the PCR product from nalidixic acid susceptible control strain cleaved at both *HinfI* sites at the codon corresponding to Ser-83 and Asp-87 of *gyrA* (Fig. 3).



**Figure 3.** PCR-RFLP Pattern of *S. enterica* serovar Typhi based on digestion of *gyrA* gene (195 bp) with *HinfI* restriction enzyme. Lanes 1-9: Treated gyrase PCR product of  $Cip^R$  strains; Lanes 10: Treated gyrase product of  $NA^S$  strain; Lanes 11: Non-treated gyrase PCR product of  $Cip^R$  strain; Lanes 12: 25 bp DNA Ladder (Invitrogen)

## Discussion

Typhoid fever is still endemic in developing countries with fatal infection in children and occasionally in adults. In the present study the disease was found highest in young children ranged within 2-3 years. This result corroborates with our previous study in which we reported the highest incidence of the disease in children younger than 5 years of age and more than one fourth of cases occurred in the first two years of life<sup>24</sup>. The prevalence of *S. typhi* infection in younger children (<5 years) has also been reported by several other groups<sup>25-27</sup>.

Using the breakpoints recommended by the CLSI<sup>21</sup>, nine typical *S. enterica* serovar Typhi isolates resistant to ciprofloxacin were isolated in this study. However, most of the isolates (89%) were found with decreased susceptibility to ciprofloxacin (MICs ranged from 0.125 - 2.0 µg/mL). The isolates with decreased ciprofloxacin susceptibility were also uniformly resistant to nalidixic acid. Recently, *S. enterica* serovar Typhi isolates with decreased ciprofloxacin susceptibility (MIC, >0.125 µg/mL) have become

the subject of worldwide attention<sup>28</sup>. Although none of these isolates were phenotypically “resistant” according to the current CLSI guidelines, treatment failure in a substantial proportion of our patients treated with ciprofloxacin has been observed. The documentation of clinician as well as microbiological failure suggests a need to re-evaluate the interpretive MIC breakpoints of fluoroquinolones, since the current reference standards are unable to distinguish between fluoroquinolone susceptible isolates and isolate with reduced susceptibility<sup>7,29</sup>.

The multiplex PCR based VNTR profiling revealed eight different types of VNTR patterns, among them 93 (73%) isolates showed two bands corresponding to ~200, >600 bp (type A). Eight out of nine ciprofloxacin resistant isolates also gave the same banding pattern as type A. As most of the isolates gave the same banding pattern like resistant strains and MIC of these strains ranged from 0.125 to 2 µg/mL (reduced susceptible group), therefore indicating all of these isolates are prone to resistance to fluoroquinolones.

Our previous report identified three ciprofloxacin resistant isolates in 2005, afterward we identified nine resistant strains with high MIC value (12 - >32 µg/mL) in 2006. These results suggest that NARST with reduced susceptibility to ciprofloxacin is now endemic in Bangladesh. Only a few cases of fluoroquinolone resistant isolates have been reported up to now, but the increasing number of isolates resistant to nalidixic acid with decreased susceptibility to ciprofloxacin is a matter of concern. There is also an urgent need to reevaluate fluoroquinolones breakpoints for *S. enterica* serovar Typhi.

More recently, *Salmonella* spp. isolates, exhibiting reduced susceptibility to ciprofloxacin but sensitive to nalidixic acid has been observed<sup>30</sup>. Therefore, screening method for detection of reduced susceptible group to ciprofloxacin by nalidixic acid disc diffusion need reconsideration. Until then, laboratory should include MIC method along with standard disc diffusion, particularly in endemic region.

This study has the limitation that patient information was collected retrospectively, only after confirmation that the isolates were *S. enterica* serovar Typhi. Furthermore, we could not confirm the mutation in *gyrA*, which contain the quinolone resistance determining region, by DNA sequencing. The emergence of highly ciprofloxacin-resistant isolates may be due to the overuse of this drug in a population with high prevalence of NARST, as predicted by Hirose et al.<sup>23</sup>. Therefore, now it is a prime concern to consider new, effective and alternative choice of drug in forthcoming days to combat against typhoid caused by ciprofloxacin resistant *S. enterica* serovar Typhi. All isolates investigated here including ciprofloxacin-resistant isolates were susceptible to ceftriaxone and azithromycin (data not shown), which might indicate that these antibiotics could still provide an appropriate therapy for typhoid fever. But the increased therapeutic cost, together with difficulties in intravenous administration can be a serious handicap in developing countries like Bangladesh.

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