

Full Length Research Paper

Emerging threats of antibiotic resistance in *Salmonella typhi* and *Salmonella paratyphi A* among enteric fever cases of Dhaka, Bangladesh

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Enteric fever is a severe public health threat because of the rising antibiotic resistance of *Salmonella* species in developing countries, especially in its endemic areas like Bangladesh. This retrospective study was aimed to assess the effectiveness of a range of 17 commonly used antimicrobials against *Salmonella Typhi* and *Salmonella Paratyphi A* isolated from 601 enteric fever cases in Dhaka, Bangladesh. Conventional biochemical tests were used to identify *Salmonella* strains and the Kirby-Bauer disc diffusion method to perform the antibiotic sensitivity in SAIC Digital Diagnostic Lab, Dhaka. The 2017 Clinical Laboratory Standard Institute (CLSI) guideline was employed to interpret the antibiogram results, and statistical software SPSS (version 22.0) to analyze the obtained data. The number of male patients (54.74%) dominated over their female counterparts (45.26%). The patients aged from 1 month to 75 years, with a mean of 19.74±12.79 years. Among 601 *Salmonella* spp. isolates, *S. Typhi* (56.57%) prevailed over *S. Paratyphi A* (43.42%). Both strains showed >85% antimicrobial insusceptibility to three major antibiotics: ciprofloxacin, gentamicin, and amikacin. *S. Typhi* (65.29%) showed significantly greater resistance to azithromycin compared to *S. Paratyphi A* (14.9%) ($p < 0.001$). Both pathogens reported over 95% sensitivity to ceftriaxone, cefixime, ceftazidime, amoxiclav, cephalixin, aztreonam, imipenem, and cefuroxime. To conclude, this study found an increased antibiotic resistance of *Salmonella* spp. to commonly prescribed antibiotics. These findings would help physicians and policymakers make informed decisions and provide better treatment to the affected patients.

Key words: *Salmonella*, antimicrobials, antimicrobial insusceptibility, antibiotic sensitivity, Dhaka.

INTRODUCTION

Enteric fever is a life-threatening systemic illness caused by Gram-negative *Salmonella Typhi* and *Salmonella*

Paratyphi A (Crump and Mintz, 2010). It attacks almost 16 million people each year and causes over 153,000

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deaths worldwide; notably, most of them belong to South Asia and sub-Saharan Africa. In 2017, nearly 17 million people worldwide got infected, and 117,000 patients lost their valuable lives with a heightened mortality of 4 to 5% (Global Burden of Disease Study, 2017). Its widespread prevalence in the developing and tropical regions like Asia and Africa is primarily due to the existing inadequate food and water safety. Likewise, this contagious fatal disease has also become endemic in Bangladesh (Crump and Mintz, 2010; Kirk et al., 2015). Between 2003 and 2004, Bangladesh reported enteric fever incidence as 200 episodes per 100,000 individuals each year compared to 394.2 episodes per 100,000 individuals in South Asia (Saha et al., 2018). One recent study by Ahmed et al. (2017) explored the bacterial etiology of bloodstream infections and found *S. Typhi* and *S. Paratyphi A* as the most frequently isolated organism with a high percentage of multidrug-resistant (MDR) strains (Ahmed et al., 2017). Worryingly, younger children in Bangladesh have experienced the highest incidence of enteric fever compared to Vietnam and other comparable regions (Brooks et al., 2005).

This deadly infection is regarded as “typhoid” when caused by *S. Typhi* and “paratyphoid” fever when by *S. Paratyphi*. These pathogens can transmit through the oral or fecal routes of patients and manifest morbidity through multiple signs: fever, abdominal pain, and non-specific symptoms, including nausea, vomiting, headache, and anorexia (Connor and Schwartz, 2005; Sur et al., 2007). When ingested, these *Salmonella* species bacteria colonize the small and large intestines, invade the gastrointestinal barrier, and then spread to the vital organs such as the liver, spleen and bone marrow (Raffatellu et al., 2008). However, due to increasing resistance of *S. Typhi*, the available antibiotics that can be considered for effective treatment are decreasing day by day (Das et al., 2017; Saha et al., 1997). This situation has been deteriorating abruptly in low and middle-income countries because of the higher antimicrobial resistance of *S. Typhi* and *S. Paratyphi A* strains. Multiple factors like incomplete treatment, overuse, and over-the-counter sales of antibiotics may contribute to this public health concern of antimicrobial resistance. Several studies confirmed that *S. Typhi* was first reported MDR against ampicillin, chloramphenicol, and cotrimoxazole in the early 1970s and ciprofloxacin in the early 1990s (Olaro and Galindo, 1973). Nowadays, roughly 90% clinical isolates from the urban settings of endemic regions showed decreased sensitivity to ciprofloxacin (Das et al., 2017; Iyer et al., 2017). Later, this trend also shifted to other classes of antibiotics such as azithromycin and ceftriaxone (Das et al., 2017). A recent study from Pakistan also revealed that *S. Typhi* induced extensive drug-resistance to ciprofloxacin and ceftriaxone (Klemm et al., 2018). Therefore, this study was carried out to investigate the current antibiotic susceptibility patterns of *S. Typhi* and *S. Paratyphi A*. Its findings would benefit healthcare professionals in making informed decisions

and providing better treatment for enteric fever patients in the coming days.

METHODOLOGY

Study design and setting

A retrospective study spanning approximately one year (January 2019 to November 2019) was conducted based on the laboratory records of the SAIC Digital Diagnostic Lab database, Dhaka. In total, 601 blood culture-positive samples collected from the enteric fever patients were assigned for the study. A semi-structured checklist was used to extract all cultures and antimicrobial sensitivity test results of patients from the laboratory records notebook.

Isolation and identification of *Salmonella* spp.

Gram-staining and conventional biochemical methods were used to identify the *Salmonella* isolates (Figure 1). A culture media enriched with Selenite broth was used to support the likely growth of pathogens (Figure 2). Following the inoculation, the media was incubated overnight at 37°C and sub-cultured into *Salmonella-Shigella* agar, blood agar, and Mac-Conkey agar. Triple sugar iron (TSI) agar was initially used to differentiate the isolated *Salmonella* strains, resulting in alkaline slant, acidic butt, and H₂S production. *S. Typhi* produced H₂S but not gas, whereas *S. Paratyphi A* produced gas and some *S. Paratyphi A* produced H₂S after 72 h. Both strains were motile but showed negative reactions in indole, citrate, and urea tests.

Antimicrobial susceptibility test (AST)

To determine the antibiotic susceptibility of *Salmonella* isolates, the Kirby-Bauer disc-diffusion method was performed on Muller-Hinton agar plates shown in Figure 2, (Bauer et al., 1966). Antibiotics used were selected based on their 2017 Clinical Laboratory Standard Institute (CLSI) guideline (CLSI, 2017), local prescription by physicians, and availability in the market. All isolates were tested against 17 different types of antibiotics from 8 classes: β -lactamases (Ampicillin-10 μ g, Aztreonam-30 μ g, Amoxicillin-Clavulanic acid- 30 μ g), Carbapenem (Imepenem-10 μ g), Aminoglycosides (Gentamycin-10 μ g, Amikacin-30 μ g), Cotrimoxazole (Co- trimoxazole-25 μ g), Cephalosporin (Cefepime 30 μ g, Ceftriaxon 30 μ g, Cefixime 5 μ g, Ceftazidime 30 μ g, Piperacillin 75 μ g, Cephalexin 30 μ g, Cefuroxime 30 μ g), Fluoroquinolone (Ciprofloxacin 5 μ g), Tetracycline (Tetracycline-30 μ g), and Macrolide (Azithromycin-10 μ g). Subsequently, the results of AST were interpreted according to the CLSI 2017 guideline.

Statistical analysis

The data were tabulated and illustrated graphically using Microsoft Excel-2019 and subsequently analyzed by the statistical software, SPSS-22. The descriptive results were represented as a percentage, relative frequency, mean \pm standard deviation (SD). At last, to find the association between the types of *Salmonella* spp. infection with patients' attributes, and antibiotic sensitivity against the tested antibiotics, Chi-square tests and Independent Sample t-test were applied.

Ethical considerations

The Institutional Review Board and chairperson of the SAIC Digital

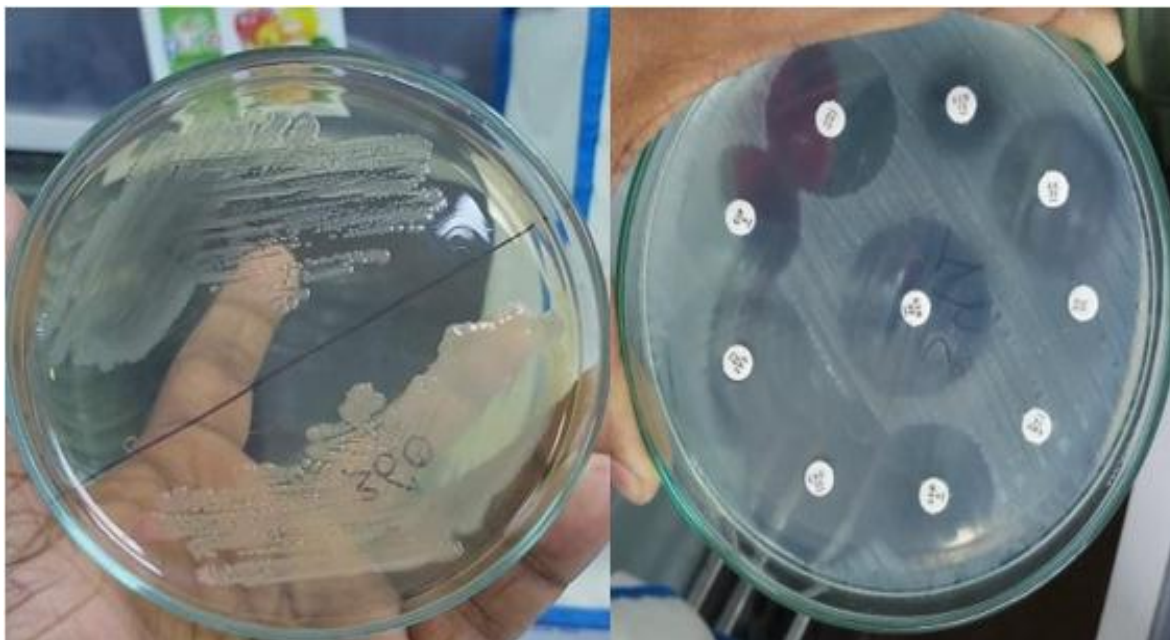


Figure 1. Biochemical tests.



Figure 2. Culture and sensitivity tests.

Diagnostic Lab, Dhaka, acknowledged the required ethical approval for the study. It was ensured that the patients selected for the study

had not received any antibiotics before 8 h of their sample collection.

Table 1. Distributions of positive cases based on sex and age of the patients.

Patients' sex and age	<i>Salmonella</i> spp.		Total [n (%)]	Statistical Tests
	<i>Salmonella</i> Paratyphi A [n (%)]	<i>Salmonella</i> Typhi [n (%)]		
Sex of the patients				
Male	145 (44.1)	184 (55.9)	329 (54.74)	$\chi^2=0.123$; $p=0.726$
Female	116 (42.6)	156 (57.4)	272 (45.26)	
Total n (%)	261 (43.42)	340 (56.57)	601 (100)	
Age of the patients (years)				
<5	31 (47.7)	34 (52.3)	65 (10.82)	$\chi^2=6.184$; $p=0.186$
5-20	115 (40.4)	170 (59.6)	285 (47.42)	
21-40	104 (48.4)	111(51.6)	215 (35.77)	
41-60	8 (29.6)	19 (70.4)	27 (4.49)	
>60	3 (33.3)	6 (66.7)	9 (1.50)	
Mean±SD of age (years)	19.87±11.97	19.64±13.39	19.74±12.79	$t=0.218$, $p=0.827$
Median age (years)	19.00	17.00	18.00	
Range of age	-	-	1 month to 75 years	

χ^2 =Chi-square Value, p =significance value at LS = 0.05.

RESULTS

Among 601 *Salmonella* isolates, 340 (56.57%) and 261 (43.42%) were confirmed as *S. Typhi* and *S. Paratyphi A*, respectively. The number of male patients (54.74%) predominated their female counterparts (45.26%). But, the distribution of male and female patients based on their infections either by *S. Typhi* or *S. Paratyphi A* was similar ($p>0.05$). Males and females suffered more from *S. Typhi* than *S. Paratyphi A*; about 60% males and 57% females tested positive for *S. Typhi*. The patients aged from 1 month to 75 years, with a mean of 19.74 ± 12.79 years. The average age of the patients infected by *S. Typhi* and *S. Paratyphi A* was nearly the same: 19.64 ± 13.39 , and 19.87 ± 11.97 years, respectively. The majority of the patients, almost 83%, were 5-40 years old. Patients of the 5-20 years group accounted for the highest, 47.42%, among all enteric fever cases, followed by the adult group, 21-40 years, contributing to 35.77% enteric fever cases. The least number of patients (1.5%) belonged to the oldest age group, >60 years. When *S. Typhi* and *S. Paratyphi* cases were distributed within different age groups, the number of typhoid patients outnumbered the paratyphoid patients in each age group. Within the groups of 41-60 and >60 years, the typhoid patients nearly doubled that of paratyphoid. The infection by both pathogens was most common among the age groups of 5 to 20 years, followed by 21-40 years (Table 1).

As shown in Figure 3, *S. typhi* and *S. paratyphi A* were highly insensitive (>85%) to ciprofloxacin, gentamycin, and amikacin. On the other hand, nearly 10-20% cases by both pathogens had developed resistance to cotrimoxazole, piperacillin, and ampicillin. Interestingly,

5 out of 17 antimicrobials tested showed invariable efficacy against nearly all typhoid and paratyphoid cases: cefixime, ceftazidime, cephalixin, aztreonam, and amoxicillin. Ten antibiotics were highly sensitive against *S. Typhi*; they all showed over 95% susceptibility (ceftriaxone 99.71%, ceftazidime 99.71%, cefepime 99.65%, cefixime 99.41%, cephalixin 98.51%, cefuroxime 98.23%, imipenem 97.35%, amoxiclav 397.31%, aztreonam 97.30% and tetracycline 96.51%). In striking resemblance with *S. Typhi*, 8 out of those 10 antimicrobials had over 95% efficacy against *S. Paratyphi A* as follows: ceftriaxone 99.2%, cefixime 98.9%, ceftazidime 98.9%, amoxiclav 98.1%, cephalixin 97.7%, aztreonam 96.5%, imipenem 96.2%, and cefuroxime 96.2%. On the other hand, *S. Typhi* demonstrated as high as over 85% resistance to the following antibiotics (gentamycin 99.12%, amikacin 99.41%, and ciprofloxacin 85.50%); however, *S. Typhi* showed lower resistance against other remaining antimicrobials (azithromycin 65.29%, cotrimoxazole 22.65%, piperacillin 21.32%, and ampicillin 19.53%) (Table 2).

Similar to the resistance shown by *S. typhi*, *S. paratyphi A* was found to be sensitive to cefepime 83.1%, tetracycline 93.5%, cotrimoxazole 87.3%, piperacillin 86.4%, and amikacin 91.9%. Likewise, *S. Paratyphi A* too showed over 85% insensitivity to the antibiotics (gentamycin 99.1%, amikacin 98.5%, and ciprofloxacin 88.5%, followed by cotrimoxazole 12.7%, piperacillin 13.6%, and azithromycin 14.9%) (Table 2). When the sensitivity of each antibiotic was distributed against the type of *Salmonella* spp., several significant variations ($p<0.05$) were observed in their sensitivity. Cefepime showed significantly uneven resistance to *S. Typhi* (.35%) and *S. Paratyphi A* (16.9%) ($p<0.001$).

Table 2. Patterns of antibiotic sensitivity of both *Salmonella*.

Antibiotics	Sensitivity	<i>Salmonella</i> spp.		Chi-square	p
		<i>Salmonella</i> Typhi [n (%)]	<i>Salmonella</i> Paratyphi A [n (%)]		
Cefepime	S	285 (99.65)	217 (83.1)	49.20	<0.001**
	R	1 (0.35)	44 (16.9)		
Ceftriaxone	S	339 (99.71)	259 (99.2)	0.663	0.416
	R	1 (0.29)	2 (0.8)		
Imipenem	S	331 (97.35)	251 (96.2)	0.677	0.411
	R	9 (2.65)	10 (3.8)		
Tetracycline	S	332 (96.51)	244 (93.5)	2.380	0.123
	R	13 (3.78)	17 (6.5)		
Cefixime	S	338 (99.41)	258 (98.9)	0.564	0.453
	R	2 (0.59)	3 (1.1)		
Ceftazidime	S	338 (99.71)	258 (98.9)	0.664	0.413
	R	1 (0.29)	2 (1.1)		
Cephalexin	S	331 (98.51)	250 (97.7)	0.583	0.445
	R	5 (1.29)	6 (2.3)		
Cotrimoxazole	S	263 (77.35)	227 (87.3)	9.752	0.002*
	R	77 (22.65)	33 (12.7)		
Piperacillin	S	262 (78.68)	216 (86.4)	5.76	0.016*
	R	71 (21.32)	34 (13.6)		
Aztreonam	S	324 (97.30)	251 (96.5)	0.286	0.593
	R	9 (2.70)	9 (3.5)		
Ampicillin	S	272 (80.47)	239 (91.9)	15.49	<0.001**
	R	66 (19.53)	21 (8.1)		
Cefuroxime	S	333 (98.23)	251 (96.2)	2.415	0.120
	R	6 (1.77)	10 (3.8)		
Ciprofloxacin	S	49(14.50)	30 (11.5)	1.160	0.280
	R	289 (85.50)	231 (88.5)		
Gentamycin	S	3 (0.88)	1 (0.9)	0.557	0.456
	R	337 (99.12)	260 (99.1)		
Amikacin	S	2 (0.59)	4 (1.5)	1.315	0.252
	R	336 (99.41)	257 (98.5)		
Amoxyclav	S	325 (97.31)	255 (98.1)	0.378	0.539
	R	9 (2.69)	5 (1.9)		
Azithromycin	S	118 (34.71)	222 (85.5)	152.370	<0.001**
	R	222 (65.29)	39 (14.9)		

Note: S - Sensitive, R - Resistant; *Statistically significant at LS=.05, **Highly statistically significant at LS=.001.

increased sensitivity of *S. Typhi* to imipenem. Two studies in Indonesia and China also noticed decreased resistance of *S. Typhi* to imipenem (Lugito and Cucunawangsih, 2017; Yaxian et al., 2015). However, we found alarmingly heightened resistance of *S. Typhi* against ciprofloxacin and azithromycin. Two relevant studies found a similar trend revealing excessive resistance of azithromycin and ciprofloxacin as 95.29 and 90.0%, respectively (Rahman, 2015; Vlieghe et al., 2012). Similarly, decreased ciprofloxacin susceptibility for *S. Typhi* has been witnessed by some studies in India recently (Chandel and Chaudhry, 2001). In addition, a study in Pakistan reported the enhanced resistance of *S. Typhi* for ciprofloxacin, that is, consistent with our finding, but that same study found reduced sensitivity to ampicillin which is not consistent with our finding (Qamar et al., 2014). *S. typhi* was also highly resistant to antibiotics like gentamycin and amikacin. In sharp contrast to us, a community-based study in Indonesia showed almost no resistance against ceftriaxone or ciprofloxacin (Punjabi et al., 2013). The antibiotic resistance pattern may vary among the countries.

Furthermore, the current study revealed that *S. paratyphi A* was greatly sensitive to cefepime, ceftriaxone, imipenem, tetracycline, cefixime, ceftazidime, cephalexin, cotrimoxazole, piperacillin, aztreonam, amikacin, amoxiclav and cefuroxime. In agreement with this, *S. Paratyphi A* showed complete sensitivity to ceftriaxone (Bhatia et al., 2007). Interestingly, like *S. Typhi* strain, *S. Paratyphi A* also became resistant to ciprofloxacin. But, unlike *S. Typhi* which showed considerable insensitivity to azithromycin, *S. Paratyphi A* was sensitive against the same antibiotic. Earlier studies, contrarily to our outcomes, found azithromycin as highly sensitive to both *Salmonella* spp. (Chandey and Multani, 2012). We also observed a strikingly resemblance between *S. Typhi* and *S. Paratyphi A* as they both demonstrated similar enhanced insensitivity to two other antibiotics: gentamycin and amikacin. In contrast, Naheed et al. (2010) found that all *S. Paratyphi A* isolates were susceptible to all antimicrobial agents they tested. In Bangladesh, alarmingly, both *S. Typhi* and *S. Paratyphi A* lost the susceptibility to azithromycin.

Azithromycin's insusceptibility to both *S. Typhi* and *S. Paratyphi A* poses an emerging public health concern as treatment failures have been reported (Molloy et al., 2010). Over-use of ciprofloxacin and azithromycin resulting from over-the-counter availability and easy oral administration, coupled with incomplete dose treatment by them might contribute to their high antibiotic resistance in Bangladesh. In the present study, not any single antibiotic had complete susceptibility to the total *S. typhi* isolates tested. Unless this increasing antibiotic resistance rate for *Salmonella* is checked, options for treating enteric fever cases would be lost shortly. Bangladesh Government should cryingly implement a national guideline on the proper usage of antibiotics.

Conclusion

The study unraveled the current antibiotic resistance patterns of *S. Typhi* and *S. Paratyphi A* to help medical practitioners so that they can make informed decisions and provide better treatment for enteric fever patients. This study revealed male and children were more susceptible to enteric fevers. Both *S. Typhi* and *S. Paratyphi A* were equally highly resistant to ciprofloxacin, gentamicin, and amikacin. Several antimicrobials presented significant variation in resistance against *S. Typhi* and *S. Paratyphi A*. Researchers and policymakers could find this study helpful in prioritizing their research scopes to tackle the upcoming challenges of antibiotic resistance among enteric fever patients.

CONFLICT OF INTERESTS

The authors have not declared any conflict of interests.

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