Phenotypic and genotypic characterization of multidrug-resistant salmonella in a tertiary care hospital

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Abstract
This is a prospective laboratory based observational study on genotypic and phenotypic characterization of multidrug resistant salmonella conducted over a period of 1 year in a tertiary care hospital. About 2280 blood samples from patients suffering from PUO and suspected cases of enteric fever were received from various departments like medicine, paediatrics, septic labour ward, clean labour ward and general surgical ward. The main goal of the study was to identify resistance patterns as well as the mechanisms contributing to resistance in salmonella. Enteric fever due to S.Typhi was seen to be 4 times more than S.paratyphi A. 144 samples were positive for Salmonella Typhi among 2280 samples. Out of which, 16 isolates showed multidrug resistant & from these, 10 drug resistant plasmids were detected. By transformation process, plasmids were transformed into E.coli. Antimicrobial testing confirmed multi drug resistance in these 10 transformed E.coli (100%). These results confirm that plasmids are responsible for drug resistance in all the strains of MDR Salmonella. It is by careful selection of appropriate antibiotics according to antimicrobial testing will avoid the emergence of MDR Salmonella in the future.

Keywords: Enteric fever, Multi-drug resistance, Plasmids, Transformation, Ampicillin, Co-trimoxazole, Chloramphenicol.

Introduction
Human infections with Salmonella are most commonly caused by ingestion of food, water or milk contaminated by human or animal excreta and humans are the only known reservoir for Salmonella Typhi. Food materials also get contaminated via polluted water or by the food handlers. Typhoid fever is a major cause of morbidity and mortality with an estimated global incidence of 21.6 million cases and 216,000 deaths per year. In developing countries, its annual incidence ranges from 12 to 622/100,000 persons. Salmonellalaenterica serovar Typhi (S. Typhi) is responsible for the majority of cases followed by S. enterica serovar para typhi A (S. paratyphi A) that causes 20% of the cases. In the last two decades, the worldwide emergence of multidrug-resistant strains of Salmonella has led to virtual withdrawal of chloramphenicol and its replacement with fluoroquinolones and third-generation cephalosporins. (1)

Chloramphenicol was the first antibiotic introduced in 1948 against Salmonella Typhi and was found very effective. However, resistance was developed two years after its introduction and it caused a major problem until 1972. (2) Ampicillin and co-trimoxazole were the effective alternatives till the development of multidrug-resistant strains. (3) The presence of nalidixic acid resistance is a marker for decreased susceptibility to fluoroquinolones and should be tested for when dealing with multidrug-resistant strains. Since the fluoroquinolones – ofloxacin and ciprofloxacin were used as alternatives in multidrug-resistant strains, which concomitantly showed resistance to nalidixic acid strains. Hence, the dose of fluoroquinolones has to be increased. (4)

Third generation cephalosporins – ceftriaxone, cefixime and azithromycin, a macrolide antibiotic are the drug of choice for multidrug-resistant typhoid fever.

Contributing factors for multidrug-resistant Salmonella Typhi are overuse and misuse of antibiotics, inappropriate prescribing practices along with the intrinsic microbiologic plasmid mediated factors.

In 2001 the entire genome of a multidrug-resistant isolate of Salmonella Typhi was sequenced. Salmonella share more than 70-80% of genes with other bacteria like E.coli. Another feature of Salmonella Typhi genome is the presence of over 200 inactivated genes which are felt to be related to the adaptation of the bacteria to the human host and possibly its ability to invade human tissue. Drug resistance is encoded in a transmissible plasmid. The development of additional horizontal genes in the Salmonella Pathogenicity Islands (SPI) represented the separation of the E.coli and Salmonella lineages and allows for the targeting of intestinal epithelial cells by Salmonella. Genetic studies have shown that resistance is encoded on H11 incompatibility plasmid and is transferable. (5)

Aims and Objectives
To study the prevalence of Salmonella Typhi at a tertiary care hospital and to identify the multidrug-resistant Salmonella Typhi among the positive isolates.
To isolate the plasmids from these MDR Salmonella Typhi. To prove genotypically, the plasmid mediated drug resistance in these isolates by transformation of isolated plasmids into E.coli, later confirmed by disc diffusion and agar dilution method.
**Materials and Method**

The present study was conducted at GRH attached to Madurai Medical College. Ethical clearance was obtained. The period of study was one year and the study population consisted of 2280 patients of different age groups attending outpatient department and the patients admitted in different wards viz. Paediatrics, Medicine, Septic labour theatre, clean labour theatre and General surgical ward at Government Rajaji Hospital, Madurai. Blood samples were collected from the study subjects who had the following signs and symptoms.

**Clinical features:** Fever which is remitting and diurnal type, nausea, vomiting, abdominal pain, diarrhea, loss of appetite, headache, lethargy and confusion.

**Clinical Signs:** Rose spots, abdominal tenderness, splenomegaly, hepatomegaly.

**Sample collection:** Blood was collected using strict aseptic precaution. In adults, 5 ml of blood was drawn and transferred to the blood culture bottle with 50ml of brain heart infusion broth swabbing the cap with 70% alcohol. Similarly in children, 1-2 ml of blood was collected in 20 ml of sterile BHI broth in blood culture bottles. Collected samples were transferred to the microbiology laboratory within 1 hour.\(^{(5)}\)

**Processing of Specimens:** In the microbiology laboratory, the specimen bottles were checked for any leakage or damage and incubated at 37°C for 48 hrs.

The specimens were inoculated on Nutrient Agar, Macconkey Agar, Blood Agar plates. The plates were incubated at 37°C for 24hrs.\(^{(5,6)}\)

**Identification of Bacteria**

**Colony morphology:** On Nutrient agar plate, salmonellae colonies were large 2-3 mm grey white, moist, translucent, low convex, discrete, smooth surface with entire edges.

On Macconkey agar, colonies were 1-3 mm in size and were non lactose fermenting colonies. On Blood agar plates, non hemolytic colonies were observed. Colonies that were presumptively identified as *Salmonella* isolates were subjected to the following tests.

GNB on Gram staining, positive Catalase test, Oxidase test-negative, Indole-negative, positive - MR test, negative – VP test, negative citrate test and negative - urease test. TSI agar media was alk/acid with minimalH₂S in case of *Salmonella Typhi* whereasH₂S is not produced by *Salmonella para typhi A* and profuse H₂S production in *para typhi B*. *Salmonella Typhi* were found to be fermentative in Oxidation – fermentation test.

*Salmonella* fermented glucose, mannitol and mannose forming acid and no gas in case of *Salmonella Typhi* and gas produced by *Salmonella Para Typhi A*.\(^{(5,6)}\)

**Slide agglutination tests:** Slide agglutination tests were conducted with the specific anti *Salmonella* agglutinating sera containing *Salmonella Typhi* and *para typhi A & B* which are used for the rapid serotyping of *Salmonella* species. The agglutination was watched by naked eye.\(^{(5,6)}\)

**Antibiotic susceptibility test**

**Disc diffusion method:** The species of *Salmonella* viz. *Salmonella Typhi* and *paratyphi A* were further subjected to antimicrobial susceptibility testing by the Kirbybaur disc diffusion method. The antibiotic discs used were ampicillin (10µg), co-trimoxazole (25µg), chloramphenicol (30µg), from Hi media. The inoculum was prepared and standardized to 0.5 McFarland turbidity. The inoculum was then spread across the surface of the Mueller Hinton agar plate of 9 cm size to give a confluent growth. Antibiotic discs were placed on the agar surface 24 mm apart. Three discs were kept on the surface of the plate at a time. After overnight incubation, the diameter of the growth inhibition was measured by an antibiotic scale (Hi media) and the sensitivity and resistance patterns were identified. The isolates which showed resistance to ampicillin, co-trimoxazole and chloramphenicol were taken as multidrug-resistant *S.Typhi and S.paratyphi A*.\(^{(7,8)}\)

**Agar dilution tests:** Minimum inhibitory concentration tests are performed using an agar dilution method. Specific volumes of antimicrobial solutions are dispensed into premeasured volumes of molten and cooled Mueller Hinton agar. A series of plates, containing varying concentrations of each antimicrobial agent ampicillin 50µg/ml, chloramphenicol 25 µg/ml and co-trimoxazole 25 µg/ml were prepared in sterile distilled water to give a final concentration ranging from 100 – 1600 µg / l. Growth control plates without antimicrobial agents were prepared. The agar is allowed to solidify and then a standard number of test bacteria are spot inoculated onto each plate. After overnight incubation, MIC is read as the lowest concentration of the antimicrobial agents that inhibits the visible growth of the test bacterium. The shelf life of agar dilution plates is only one week and the plates were stored at 2 to 8°C. These isolates were kept separately and preserved in refrigerator at 4 to 5°C for further studies. Genotypically plasmid profile analysis was done for all these isolated strains using gel electrophoresis method.\(^{(7,8)}\)

**Plasmid DNA isolation by alkaline lysis method (Sambrooke et al 1989):**\(^{(9)}\) Plasmids are non obligate, circular extra chromosomal bacterial replicons. Plasmid DNA isolates require separation of DNA from the chromosomal DNA from the bacterial cell as well as from the polysaccharides, lipids and proteins that constitute the cell. Subsequent manipulation especially enzymatic modification of the plasmid DNA requires that it be free of impurities.

**Procedure:** Sterile Eppendorff were taken and to which normal saline 1.5 ml / peptone water was added. A loopful of isolated colonies were added and incubated for 18-24 hours. It was mixed in vortex and micro centrifuged for 5 minutes at 10,000 rpm. After centrifugation, the supernatant was discarded and tubes
were drained on paper towel. 0.1 ml of GTE mixture was taken (Glucose + Tris chloride + EDTA) and added to the deposit, mixed in vortex and incubated for 5 minutes at room temperature. 0.2 ml of NaOH with sodium dodecyl sulphate was added, mixed in a vortex and then kept on ice for 5 minutes. 150μl of potassium acetate was added, mixed in a vortex and kept on ice for 5 minutes. After which it was centrifuged for 3 minutes at 10,000 rpm. Carefully the supernatant was taken to a new eppendorff tube, 800 μl of 80% alcohol was added and kept for 2 minutes at room temperature. Centrifuged the tube for 1 minute at 10,000 rpm. A milky white thin layer or pellet was seen at the bottom. The supernatant was discarded with the help of a micro pipette. 1 ml of ice cold 100% ethanol was added to the pellet and kept for 3 minutes and centrifuged at 10,000 rpm for 3 minutes. The supernatant was removed and pellet was air dried. 30 μl of TE buffer was added to the pellet and plasmid was stored at -20°C.

Electrophoresis: The apparatus was washed with 1 x TAE buffer. 1% of agarose was prepared with 50 ml of TAE buffer. 3μl of ethidium bromide was added to the agarose solution (Use gloves while adding this) and was poured to the electrophoresis plate containing comb. The agarose slab was then fixed into electrophoresis apparatus. The plasmid was mixed with 6x loading dye (fermentas) along with a control and run at 100 V for 60 minutes and the bands were read in a gel-doc system.

Transformation of the resistant plasmids into competent cells (E.coli): 9,10 500 μl of the plasmid DNA was added to 500 μl of one shot chemically competent E.coli (Invitrogen, CA) and mixed gently. The mixture was incubated on ice for 5 to 30 minutes. (E.coli shrinks by this method) Then the mixture was kept in water bath at 42 to 42.5°C for 30 to 50 seconds without shaking, so that the cells will expand. Then the tubes were transferred immediately to the ice. 250 μl of LB broth (Lysogenic broth) were added to the mixture(to enhance cell growth at room temperature). The tubes were tightly capped and shaken horizontally (200 rpm) at 37°C for 1 hour. Transformed mixture of the competent cells was kept separately. Antibiotic susceptibility testing of transformed competent cells (E.coli) by disc diffusion method using lysogenic broth agar and agar dilution method was done as described earlier.

About 25 to 200 μl from each transformation was spread on a prepared selective plate (Lysogenic agar) containing the following antibiotics, ampicillin 50 μg / ml, chloramphenicol 25μg / ml and co-trimoxazole 25μg / ml were added. The transformed mixture was spread evenly on to LB agar using ‘L’ rod and incubated overnight at 37°C. The plates were observed for resistant patterns. The antibiotic susceptibility test results of transformed competent cells were compared with the antibiotic susceptibility test of the previous MDR Salmonella isolates.

Results
The study was conducted over a period of 1 year and 2280 samples were collected. All the samples were sub cultured on Nutrient agar, Blood agar, Macconkey agar, confirmed by biochemical tests and slide agglutination test. Out of 2280 blood samples, 144 (6.31%) were positive for Salmonella Typhi by culture, biochemical tests and slide agglutination test and 43 cases (1.88%) were positive for Salmonella paratyphi A. Results are given in Table 1.

Chi-Square test (X²) showed that there was a highly significant difference between typhoid fever cases and paratyphoid fever cases (X² =56.88; df = 1; p<0.001). Enteric fever due to Salmonella Typhi was seen to be four times more than that caused by Salmonella paratyphi A in this study. 187 Salmonellae isolates were subjected to antibiotic susceptibility test by disc diffusion and dilution methods using ampicillin, chloramphenicol and co-trimoxazole. By disc diffusion method, it was found that out of 144 Salmonella Typhi isolates, 16 (11.11%) showed resistance to all three drugs (ampicillin, chloramphenicol and co-trimoxazole) and out of 43 Salmonella paratyphi A, none exhibited resistance to all three drugs. The results are depicted in Table 2.

The 144 isolates were also tested for antibiotic susceptibility test by agar dilution method and it was found that 16 samples were resistant for ampicillin (ranging from 100 – 400 mg/l), chloramphenicol (ranging from 100-400mg/l) and co-trimoxazole dilution (ranging from 200 – 1600 μg/l) as shown in Table 3.

It was inferred that 11.11% samples were multidrug-resistant by agar dilution method. Plasmid isolation studies were done on these 16 isolates. Plasmids were isolated from 10 of these MDR isolates. This is depicted in Table 4.

It was inferred that plasmids were detected in 62.5% of multidrug-resistant isolates.

The isolated plasmids were transformed to competent cells (E.coli) by transformation process successfully. Then the antimicrobial susceptibility of 10 transformed E. coli was performed against three antimicrobial compounds namely ampicillin, co-trimoxazole and chloramphenicol. The transformed E.coli were found to be resistant to all three antimicrobialagents→chloramphenicol, co-trimoxazole and ampicillin by disc diffusion and agar dilution method. This is depicted in Tables 5 & 6.

Out of 10 transformed competent cells (E coli), all the 10 isolates (100%) showed resistance to ampicillin, chloramphenicol and co-trimoxazole by disc diffusion method. Out of 10 transformed competent cells, all the 10 (100%) isolates showed resistance to ampicillin, chloramphenicol and co-trimoxazole by agar dilution method in dilution ranging from 100 to 1600 mg/ml. These results confirm the successful transformation of
E. coli with plasmid DNA of multidrug-resistant Salmonella Typhi.

**Table 1: Comparison of positive isolates**

<table>
<thead>
<tr>
<th>S. No.</th>
<th>Organism isolated</th>
<th>Positive by Culture &amp; Biochemical tests</th>
<th>Positive by slide agglutination test</th>
<th>Percentage (n=2280)</th>
</tr>
</thead>
<tbody>
<tr>
<td>1.</td>
<td>Salmonella Typhi</td>
<td>144</td>
<td>144</td>
<td>6.31</td>
</tr>
<tr>
<td>2.</td>
<td>Salmonella paratyphi A</td>
<td>43</td>
<td>43</td>
<td>1.88</td>
</tr>
</tbody>
</table>

**Table 2: Resistant patterns by Disc Diffusion method**

<table>
<thead>
<tr>
<th>S. No.</th>
<th>Isolates</th>
<th>Nos.</th>
<th>Resistant (Zone of inhibition in mm)</th>
<th>%</th>
</tr>
</thead>
<tbody>
<tr>
<td></td>
<td></td>
<td></td>
<td>Ampi. (&lt;13)</td>
<td>Chlor (&lt;12)</td>
</tr>
<tr>
<td>1.</td>
<td>Salmonella Typhi</td>
<td>144</td>
<td>16</td>
<td>11.11</td>
</tr>
<tr>
<td>2.</td>
<td>Salmonella paratyphi A</td>
<td>43</td>
<td>-</td>
<td>-</td>
</tr>
</tbody>
</table>

**Table 3: Resistant pattern of Salmonella Typhi by Agar dilution method (MIC values mg/l)**

<table>
<thead>
<tr>
<th>S. No.</th>
<th>Antibiotics</th>
<th>No. of strains tested</th>
<th>50</th>
<th>100</th>
<th>200</th>
<th>400</th>
<th>800</th>
<th>1600</th>
<th>3200</th>
<th>6400</th>
<th>Percentage (n=2280)</th>
</tr>
</thead>
<tbody>
<tr>
<td>1.</td>
<td>Ampicillin</td>
<td>144</td>
<td>-</td>
<td>5</td>
<td>4</td>
<td>7</td>
<td>-</td>
<td>-</td>
<td>-</td>
<td>-</td>
<td>100</td>
</tr>
<tr>
<td>2.</td>
<td>Chloramphenicol</td>
<td>144</td>
<td>-</td>
<td>5</td>
<td>5</td>
<td>6</td>
<td>-</td>
<td>-</td>
<td>-</td>
<td>-</td>
<td>100</td>
</tr>
<tr>
<td>3.</td>
<td>Co-trimoxazole</td>
<td>144</td>
<td>-</td>
<td>-</td>
<td>4</td>
<td>6</td>
<td>3</td>
<td>3</td>
<td>-</td>
<td>-</td>
<td>100</td>
</tr>
</tbody>
</table>

**Table 4: Plasmid isolation from multidrug-resistant Salmonella Typhi**

<table>
<thead>
<tr>
<th>S. No.</th>
<th>Total numbers tested for Plasmids</th>
<th>No. of plasmids isolated</th>
<th>Percentage (n=16)</th>
</tr>
</thead>
<tbody>
<tr>
<td>1.</td>
<td>16</td>
<td>10</td>
<td>62.5</td>
</tr>
</tbody>
</table>

**Table 5: Resistance of transformed E. coli to antimicrobial compounds by disc diffusion method**

<table>
<thead>
<tr>
<th>Antimicrobial agents</th>
<th>Resistant</th>
<th>% (n=10)</th>
</tr>
</thead>
<tbody>
<tr>
<td>Chloramphenicol</td>
<td>10</td>
<td>100</td>
</tr>
<tr>
<td>Ampicillin</td>
<td>10</td>
<td>100</td>
</tr>
<tr>
<td>Co-trimoxazole</td>
<td>10</td>
<td>100</td>
</tr>
</tbody>
</table>

**Table 6: Resistant pattern of transformed competent cells (E. coli) by agar dilution method (MIC in mg/l)**

<table>
<thead>
<tr>
<th>S. No.</th>
<th>Antibiotics</th>
<th>No. of strains tested</th>
<th>50</th>
<th>100</th>
<th>200</th>
<th>400</th>
<th>800</th>
<th>1600</th>
<th>3200</th>
<th>6400</th>
</tr>
</thead>
<tbody>
<tr>
<td>1.</td>
<td>Ampicillin</td>
<td>10</td>
<td>2</td>
<td>3</td>
<td>3</td>
<td>2</td>
<td>-</td>
<td>-</td>
<td>-</td>
<td>-</td>
</tr>
<tr>
<td>2.</td>
<td>Chloramphenicol</td>
<td>10</td>
<td>2</td>
<td>5</td>
<td>2</td>
<td>1</td>
<td>-</td>
<td>-</td>
<td>-</td>
<td>-</td>
</tr>
<tr>
<td>3.</td>
<td>Co-trimoxazole</td>
<td>10</td>
<td>2</td>
<td>1</td>
<td>2</td>
<td>3</td>
<td>2</td>
<td>-</td>
<td>-</td>
<td>-</td>
</tr>
</tbody>
</table>

**Discussion**

The present study was carried out to know the phenotypic and genotypic characterization of multidrug-resistant Salmonella Typhi. Blood samples collected from patients, suspected of enteric fever and PUO at a tertiary health care setting - Government Rajaji Hospital, Madurai. The drug resistant genes present on plasmids were demonstrated by isolation of plasmids and further characterization was done after transformation.

The study showed that enteric fever due to Salmonella Typhi was seen to be more common than that caused by Salmonella paratyphi A (6.31% and 1.88% respectively). In accordance with the study, Raveendran et al (2008),(11) Prabhakar et al (1996),(12) Sweta et al (2000),(13) and Madhulika et al (2004)(14) also found 1.66%, 2.7%, 3.32% and 12.11% Salmonella Typhi isolates respectively in their studies. Raveendran et al (2008)(11) explained that there were convalescent carriers in the population during their study period and Prabhakar et al (1996)(12) pointed out that the habits of hand washing techniques were lacking in their study group and they had the habit of sharing food from the same plate. Perhaps, these factors might have contributed to the increased incidence of S. Typhi in their studies. More isolation of S. Typhi in this present study may be due to close contact of children with asymptomatic carriers in the population because most of the cases in the study population were from endemic areas where carriers were more common.
common. However, as carrier study was not conducted, it could not be proved.

In this present study, multi drug resistance was shown to exist in and around Madurai District. It was found that out of 144 Salmonella Typhi isolates, 11.11% of them were multidrug-resistant Salmonella Typhi. In accordance with this study, Jesudasan et al (1992), Raveendran et al (2008), OH et al (1994), Uma Saberwal (1992) also found 11%, 14.9%, 17.4% and 18.3% MDR isolates respectively in their studies. But, studies by Prabhakar et al (1996) (Ciraj et al, 1991), Ishaq et al (1990) and Bhutta et al (1991) found 61.4%, 57.9%, 30% and 20% MDR strains respectively in their studies and explained that the emergence of MDR might be due to the early rate of relapse. They also explained that the other reasons for emergence of MDR could be due to indiscriminate use of antibiotics, inappropriate prescribing practices and also due to intrinsic microbiological plasmid mediated factors. In this present study, plasmid mediated factor was proved to be the cause of emergence of MDR by Sambroke et al (1989) method.

In this present study, a single plasmid of 120 kb was isolated from 62.5% of the MDR strains. Similarly, Karmakar et al (61.84%) (Jesudasan et al 59.5%) and Mandal et al (60.2%) also isolated plasmids of 120 kb from MDR strains. In all these studies, the samples were blood, the cases were suspected cases of enteric fever and the method of study followed was Amyes and Gould. Also studies by Mandal et al (2003) indicated the presence of R-plasmid in S.Typhi isolates by transconjugation method (Protocol of Jevanan et al). It was demonstrated by him that R-plasmids isolated from E.coli could convert antibiotic sensitive S. Typhi to antibiotic resistant S. Typhi. Transconjugation studies by Senthilkumar and Prabhakaran (2005) showed that in conjugation studies, (Cappuccino & Sherman procedure) the isolates having R+ factor showed the transfer of drug resistance through conjugation and it was determined by the development of antibiotic resistance in recipients. Bader Jehan Farooqui (1993) did both conjugation (Miller) and transformation studies. In transformation studies he used host E.coli HB 101 and the plasmids were transformed successfully. He also did conjugation experiments and found that the plasmids are conjugative in nature. In this present study, transformation process was carried out to transfer isolated plasmids into competent (E.coli) cells. (invitrogen, CA) and proved to be 100% successful. Moreover the transformation process of plasmid transmission is an easy technique, results can be obtained within a shorter period and cost wise it is also a cheaper method.

All the transformed competent (E.coli) cells were subjected to disc diffusion and agar dilution method. It was found that the transformed E.coli also showed resistance to all three antimicrobials as similar range as donor cells as in the studies of Bader Jehan Farooqui (1993). Thus, the plasmid mediated drug resistance proved in this study may be due to various factors including the acquisition of plasmid encoded inactivating enzymes like beta lactamas and chloramphenical acetyl tranferase.

**Conclusion**

The study revealed 11.11% of S.Typhi to be drug resistant for 3 drugs. 62.5% of these showed plasmids, same transformed to competent cells (E.coli). The transformed cells showed 100% resistance to the three antimicrobials. Thus, confirming that a single plasmid is responsible for carryover of the drug resistance in all strains of MDR S.Typhi. This study proved the emergence of MDR Salmonella Typhi at GRH, Madurai and it was plasmid-mediated. It is a signal for the clinicians to prevent future epidemic of MDR Salmonella in the community by choosing proper antibiotics by drug sensitivity, to avoid improper usage and drug abuse during clinical management.

**References**


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