Typhoid and Growth Characteristics of *Salmonella paratyphi* A obtained from Blood sample of a Paratyphoid Case

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**ABSTRACT**

Paratyphoid fever has been a disease of public health problem in Asian countries. In India clinical cases of paratyphoid had been emerging on escalating rates. This problem requires a consistent monitoring of the isolation throughout the nation to analyze the spectrum of the *Salmonella* enteric serovar Paratyphi A. This study examined isolation of *Salmonella paratyphi* A from a clinical sample of paratyphi A obtained from Yashoda hospital, Ghaziabad. This sample was characterized by biotyping as well as serotyping. The characterization of the bacteria was reconfirmed at National Salmonella and Escherichia Centre, Kasauli. The isolate serotyping was based on somatic O and phase 1/2 flagellar antigens by agglutination tests with antisera according to the Kauffmann White scheme. The clinical isolate was confirmed to be a case of *Salmonella paratyphi* A by biotyping and serotyping. The bacteria achieved a complete growth when propagated in the laboratory in controlled temperature, humidity and pH. This study enlights the *Salmonella paratyphi* A characterization by biotyping and serotyping as well as cultivation techniqne in laboratory. The escalating rate of paratyphoid infections appears as a budding problem which requires the advancement in diagnostic techniques as well as effective vaccines for *Salmonella enterica* serovar Paratyphi A to deal with emerging cases of infections.

**Keywords**

Biotyping; Serotyping; Cultivation; *Salmonella paratyphi* A; India.

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**Introduction**

Enteric fever, includes typhoid fever caused by *Salmonella typhi* and paratyphoid fever caused by *S. enterica* serovar Paratyphi A and is one of the most important febrile illnesses in tropical and subtropical countries, with high rates of morbidity and mortality. *S. paratyphi* A can be isolated from the blood and feces from paratyphoid fever patients (1, 2). In India, however *Salmonella enterica* serotype Typhi remains predominating in causing enteric fever, isolation of *Salmonella enterica* serotype Paratyphi A causing the same disease, has also been reported increasingly in various parts of North and South India including regions like New Delhi, Chandigarh, Shimla, Kolkatta, Rourkela, Orissa, Calicut and Manipal (3,4,5,6,7,8,9,14).

Typhoid fever is estimated to have caused
21.6 million illnesses and 216,500 deaths globally in 2000, and based on one case of paratyphoid fever for every four cases of typhoid fever, *S. paratyphi* A is estimated to have caused an additional 5.4 million illnesses in 2000. These estimates, however, are known to vary widely between regions while the cases of enteric fever caused by *S. paratyphi* A are still increasing (16). The incidence of typhoid and paratyphoid fever is increasing especially in South Asia and South Africa, where the annual incidence is 622/100,000 and 233/100,000, respectively (17).

*S. paratyphi* A has the flagellar antigens and is a member of somatic O group A with more prevalent phage type I. For serotyping of *Salmonella*, lipopolysaccharide epitopes in bacterial membrane (O antigens) and flagella proteins (H antigens) should be identified with the respective antibodies (15). Biochemically, the organism resembled *S. enterica* serotype Typhi; particularly having negative citrate, arginine, ornithine, arabinose, and rhamnose reactions. However the identity of both organisms can be confirmed by full serotyping via Kauffman-White scheme, in which the *S. enterica* serotype Paratyphi A isolate is assigned the antigenic formula I 2,12:a:1, whereas the *S. enterica* serotype Typhi isolate is assigned the antigenic formula I 9,12[Vi]:d: (18). The given study is exploring the biochemical and serotyping characteristics of *Salmonella paratyphi* A isolate obtained from a clinical paratyphoid case with emphasis on analysis of growth characteristics of the isolated bacteria.

**Materials and Methods**

For the biotyping, all media and biochemicals were obtained from Hi Media Lab. Pvt. Ltd., India. The biotyping of isolate *S. paratyphi* A was performed by conventional biochemical tests (10,11) and confirmed by serotyping on the basis of somatic O and phase 1 and phase 2 flagellar antigens by agglutination test using standard *Salmonella* agglutinating factor sera in accordance to the scheme of Kauffmann-White (12,13). The isolates were cultured on nutrient agar, Mac-conkey agar and blood agar plates.

The sera raising was performed in rabbits by following an immunization schedule with 0.25 % formalin killed *S. paratyphi* A suspension of 0.5 ml, 1.0 ml, 1.5 ml, 2.0 ml dose by subcutaneous route on day 1,2,7 and 11. The animals were test bled after one week of the last injection and tested for homologous titres by agglutination test. The titre of ≥ 1:1000 for O sera and ≥ 1:1200 for H sera was the satisfactory level for testing purpose. The animals were bled by cardiac bleeding method. Blood was collected in dry sterile test tubes and then stored in a refrigerator (+4ºC) overnight. The blood clot was removed by centrifugation and all the sera from different bleedings were pooled together and filtered by 0.45µm micro-filter.

For serotyping purpose an 18 hour broth culture of the clinical isolate was taken. It was then 0.25 % formalinized and kept at room temperature for 30 minutes. The culture was centrifuged at 2,000 rpm for 15 minutes. The supernatant was discarded and the deposit was used for serotyping work. 4-5 drops of the centrifuged deposit were placed on a clean glass slide. To the first drop was applied a drop of H: a factor antisera. The agglutination was positive, the *Salmonella* serotype was identified. The tube agglutination test was also performed respectively. These confirmatory tests were performed by using standard diagnostic antigen and factor antisera for *Salmonella paratyphi* A.
For the serological specificity the bacterial culture of *S. paratyphi* A was grown on solid soyabean casein digest (SCD) media. The antiserum against *Salmonella paratyphi* A was added to the media plates with bacterial growth and these plates were then incubated at 37°C for overnight.

The studies for analysis of growth pattern analysis were also carried out (20,21). A fresh colony of bacteria was inoculated into SCD media. 2.5 ml of the turbid growth was inoculated to 50 ml of SCD media and the flask was incubated at 37°C for overnight. 7.5 ml of the turbid growth obtained was inoculated to 150 ml of SCD media and the flask was incubated at 37°C for overnight at pH 7. Next day 50 ml of the turbid growth was inoculated to 4 flasks having 1.5 lts SCD media in each flask and these flasks were incubated at 37°C /125 rpm on rotary shaker. After 3 hours the turbidity appeared in the flasks. 40% sterile glucose was added to make final concentration of 5% glucose in each flask.

Added 2N-NH₄OH to make the pH 7.2 and again incubated them at 37°C /125 rpm on rotary shaker. The culture was grown in flasks at 37°C on rotary shaker for 12 hours. Kept the pH at 7.00 using 2N-NH₄OH. The culture was kept at 37°C for overnight incubation. Repeated Gram staining was performed to check the purity of culture at every stage. The bacterial growth was found pure and uniform. The bacterial samples were withdrawn to analyze the absorbance at 600nm (19). The pH was adjusted to 7.2 using 2N-NH₄OH and formalin at the final concentration of 1% added for final inactivation. The culture was kept for inactivation at room temperature for 24 hours. This *Salmonella paratyphi* A was further used for the harvesting of lipopolysaccharide for the O-specific polysaccharides.

**Results and Discussion**

*S. paratyphi* A was sub-cultured on blood agar and Mac-Conkey agar, after overnight incubation at 37°C, blood agar showed non pigmented (grey-white) colonies of size 1-2 mm, opaque, non-hemolytic, moist, circular with a smooth convex surface and the entire edge. The growth on MacConkey agar consisted of non lactose fermenting colonies of similar morphology. Colonies were catalase positive and oxidase negative. Gram stained smear from the growth revealed Gram-negative bacilli, 2-4 μm × 0.6 μm in size, non-capsulate and non-sporing. The bacilli were motile.

Biotyping confirms *Salmonella paratyphi* A by cultures which were Lactose negative, Catalase positive, Oxidase negative, Indole negative, Methyl Red positive, Vogus Proskaur negative, Citrate negative, Urease negative, Triple Sugar Iron showed K/A glucose acid with gas, Nitrate reductase positive and Lysine negative.

The positive agglutination of *Salmonella paratyphi* A antisera with *Salmonella paratyphi* A bacteria was observed. The colonies showed serological specificity by creating haloes of immune-precipitation when grown on SCD agar media containing antiserum against *Salmonella paratyphi* A.

Absorbance studies were conducted to analyze the growth pattern of the *Salmonella paratyphi* A. The samples of bacterial growth were collected at various steps during propagation of bacteria and the optical density at 600nm was checked. The results are given in Table no.-01 and in Diagram No.-01.

There was an up-scaling in the absorbance at 600nm of the bacteria. As the volume of media increases the O.D of the culture was found in increasing order. The samples were
obtained at 12 hours of growth and the culture was in the log phase. This indicates that the *Salmonella paratyphi* A seed was propagated with better culturist behavior.

**Table 1** Growth Analysis of *Salmonella paratyphi* A

<table>
<thead>
<tr>
<th>S.No.</th>
<th>Volume of Media</th>
<th>Incubation time</th>
<th>Purity (Gram Staining)</th>
<th>OD at 600nm</th>
</tr>
</thead>
<tbody>
<tr>
<td>1.</td>
<td>2 ml</td>
<td>Nil</td>
<td>Pure</td>
<td>0.201 (Autozero)</td>
</tr>
<tr>
<td>2.</td>
<td>20 ml</td>
<td>12 hour</td>
<td>Pure, Uniform</td>
<td>1.375</td>
</tr>
<tr>
<td>3.</td>
<td>50 ml</td>
<td>12 hour</td>
<td>Pure, Uniform</td>
<td>1.215</td>
</tr>
<tr>
<td>4.</td>
<td>100 ml</td>
<td>12 hour</td>
<td>Pure, Uniform</td>
<td>1.555</td>
</tr>
<tr>
<td>5.</td>
<td>3000 ml</td>
<td>12 hour</td>
<td>Pure, Uniform</td>
<td>2.398</td>
</tr>
</tbody>
</table>

The ‘gold standard’ for identifying the cause of an infection is the isolation and identification of the causative agent of disease. In the absence of a viable bacterium, antibody tests can give evidence of infection provided that suitable immunoassays, based on well-characterized antigens, are used. Biotyping characters are considered as markers of major importance in the typing of salmonellae. On the otherhand, serotyping is based on the long standing observation that the microorganisms from the same species can differ in the antigenic determinants expressed on the cell surface. Serotyping method is used to differentiate isolates of *Salmonella* beyond the subspecies level. The reference sera used for serotyping contain antibodies to the panel of the lipopolysaccharide and flagellar antigens of *S. paratyphi* A. The *S. paratyphi* A bacteria when cultivated at temperature of 37°C at pH of 7.2-7.4 in presence of additional glucose ratio then the log phase can be achieved within 12 hours. Higher volumes of media lead to better bacterial yields in comparison to lower volumes of media.

The present study endorses the observation of *Salmonella paratyphi* A as rapidly
emerging pathogen of enteric fever. So, continued surveillance on the mutual analysis by biotyping and serotyping of Salmonella serovar paratyphi A, the multidrug resistance, diagnosis and antimicrobial susceptibility from different regions of the country will help in up-dating the knowledge for proper therapeutic cure and vaccine development.

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References


