

Salmonella serovars associated with bacteraemia infection in persons infected with human immunodeficiency virus with low CD4⁺ cell counts in Akwa Ibom State, Nigeria

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ABSTRACT

Background: Certain serovars of *Salmonella* show a much higher predilection for causing bacteraemia as opportunistic infections in human immunodeficiency virus (HIV) infected persons.

Objectives: The study was undertaken to characterize serovars of *Salmonella* isolated from bacteraemia infections in HIV/AIDS patients in Akwa Ibom State and to determine the relationship of the infection with the low CD4⁺ cell counts of the subjects.

Materials and Methods: A cross-sectional descriptive epidemiological study was conducted among 300 HIV/AIDS patients and 105 non-HIV subjects. Blood culture samples were collected and cultured on *Salmonella-Shigella* agar (SSA) and Deoxycholate citrate agar (DCA) for recovery of *Salmonellae*. Isolates were identified using conventional tests and Microgen™ GN ID System (Microgen Bioproducts). Serotyping was done using *Salmonella* antisera (Statens Serum Institut, Denmark). Estimation of CD4⁺ cell counts was by Partec flow cytometry.

Results: *Salmonella* species and *Salmonella enterica arizonae* were identified. Seven serovars were obtained; *S. typhi*, *S. paratyphi* A, *S. paratyphi* B, *S. paratyphi* C, *S. typhimurium*, *S. choleraesuis* and *S. enteritidis*. The prevalence of *Salmonella* associated bacteraemia was significantly higher ($p < 0.05$) in HIV persons with 13.7% in contrast to 3.8% non-HIV subjects. Regardless of the HIV/AIDS sero-status, subjects with CD4⁺ counts below 200 μ L had the highest rates of *Salmonella* associated bacteraemia.

Conclusion: *Salmonella typhi*, *S. paratyphi* A, *S. paratyphi* B, *S. paratyphi* C, *S. typhimurium*, *S. choleraesuis* and *S. enteritidis* are identified as serovars of *Salmonella* which are endemic in the study area and associated with bacteraemia infection with significantly higher occurrence in subjects with low CD4⁺ counts compared to HIV seronegative subjects.

Keywords: *Salmonella*, Bacteraemia, HIV infection, CD4⁺ counts.

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INTRODUCTION

Bacteraemia is defined as the presence of bacteria in the bloodstream (1). Bacteria can enter the bloodstream through severe complications of infections, dental procedures, catheterization, other foreign bodies entering the arteries and veins and during surgery involving the mucous membranes (2). Individuals with human immunodeficiency virus (HIV) infections are at greater risk of bacteraemia because of their defective immune status (3-5).

The rates of *Salmonella* invasive infections and mortality are higher in infants, elderly and people living with HIV and hemoglobinopathies (6). *Salmonella* species have emerged as a major public health problem in developing countries, particularly sub-Saharan Africa, including Nigeria (7,8). The increasing prevalence of people living with HIV/AIDS has increased the occurrence of both the typhoidal and non-typhoidal salmonellae in bacteraemia in sub-Saharan Africa (9, 10).

There are two species of *Salmonella* complex namely *Salmonella bongori* and *Salmonella enterica* (11). The species of *Salmonella enterica* has six subspecies with more than 2579 serovars currently identified (12). Certain serovars of *Salmonella* show a much higher predilection for causing bacteraemia and these serovars differ with geographic location (13, 14).

Infection with HIV is associated with a progressive decrease of CD4⁺ T-Cells in the immune system, resulting in an increase in viral load. The normal CD4⁺ T-Cell count for an adult is over 1,000 cells/ μ L (15). In HIV pathogenesis, the primary target is the cells bearing CD4⁺ markers on their surfaces (16-18). Thus, CD4⁺ cell count determination is the best validated predictor of the likelihood of developing series of opportunistic infections as well as an important parameter to initiate prophylaxis that helps to reduce the risk and severity which would have resulted in death of the patients (19, 20). Moreover, screening for opportunistic infections is an aspect of care and support to these immunosuppressed persons (18).

Therefore, this study aimed at screening and identifying *Salmonella* serovars causing bacteraemia in HIV/AIDS patients in the study area and to determine the relationship between CD4⁺ cell counts and the distribution of *Salmonella* infections in HIV-infected persons.

MATERIALS AND METHODS

Study Design

This research work is a cross-sectional descriptive study undertaken to isolate and characterize *Salmonella* serovars associated with bacteraemia infection in persons infected with HIV/AIDS patients with CD4⁺ cell counts of less than 1,000 cells/ μ L (15), in Akwa Ibom State, Nigeria.

Ethical approval

Approval for the work was obtained from the Ethical Committee of the University of Uyo Teaching Hospital (UUTH), Akwa Ibom State, Nigeria. Consents were also sought from the subjects who were willing to volunteer. Consents for minors (below 18 years) were obtained from the parents and guardians.

Collection of samples

Fresh blood samples for culture were collected from a total of 300 confirmed HIV/AIDS infected persons and 105 HIV negative people (control subjects) control who consented to be enrolled into the study.

Inoculation of Samples

Five millilitres (5mL) of blood sample was collected intravenously from each patient and 1mL of the blood was dispensed into blood cultured bottles containing 5mL tryptone soy broth medium and was incubated at 37°C for 48 hours in a standard incubator (DNP-9052 Laboratory incubator, TCA, England). Subculture was made on *Salmonella-Shigella* agar (SSA) plates, while the remaining 4mL was dispensed into blood sample bottles containing ethylene diamine tetra-acetic acid (EDTA) as anticoagulant for HIV test and for CD4⁺ cell estimation.

Identification of isolates

All purified colonies were identified using the different biochemical tests (15,21,22) and Microgen™ GN ID System (Microgen Bioproducts, USA), an identification system for all currently recognized Enterobacteriaceae were used to confirm the species of *Salmonella* isolates obtained in the study.

Serotyping of Salmonella

Serotyping of *Salmonella* isolates was carried out using *Salmonella* antisera and monoclonal antibodies for serological confirmation. Polyvalent "H" (poly A –E + Vi) antisera and "O" (O:2, O:4, O:7, O:9) antisera (Statens Serum Institut, Denmark) were used.

Estimation of CD4⁺ cell counts

Estimation of CD4⁺ cell counts was conducted using Partec flow cytometry method. The analysis involves the use of cytoflow beads with a Partec flow cytometry counter machine. Flow cytometry uses laser to excite fluorescent antibody, probes specifically for CD4⁺ and other cell surface markers, to distinguish one type of lymphocyte from another.

The method involves measuring 20µL of fresh whole EDTA blood (0.02mL) and this quantity of blood was added to 20µL of CD4⁺/MAb (0.02mL of counting beads) in a test tube and incubated in the dark at room temperature of 37°C for 15 minutes after which, eight hundred microlitres (800µL/0.8mL) of phosphate buffered saline was added to the blood sample and run on the cytometry machine. Thereafter, the machine automatically generated the report of the quantity of the red blood cells present and the number of CD4⁺ in the cells.

Statistical analysis

Percentages were employed to express the prevalence and the frequency of occurrences of the infection. Analysis of Variance (ANOVA) was used to determine the significant level of *Salmonella* associated infections in HIV sero-positive subjects and to determine the role between the infection and CD4⁺ cell counts of the subjects studied.

RESULTS

Prevalence of *Salmonella* bacteraemia among HIV/AIDS and the controls

Prevalence of *Salmonella* bacteraemia among HIV/AIDS and the control subjects is presented on Figure 1. The results revealed that of the 300 HIV/AIDS patients studied, a total of 41 (13.7%) had *Salmonella* associated bacteraemia while 4 (3.8%)

of the blood samples of the 105 control subjects yielded *Salmonella* species. The prevalence in the test group differed significantly from that in the controls ($p < 0.05$). The results of biochemical characterizations and Microgen™ GN ID System identification of the isolates are presented on Table 1. The isolates were identified as *Salmonella enterica* species and *Salmonella enterica Arizonae* respectively. The biochemical reaction results obtained from the biochemical tests carried out on the test isolates are shown in Table 2.

Table 3 shows serotyping results of *Salmonella* isolates obtained in the study. *Salmonella* isolates agglutinated the polyvalent H antisera used as genus specific antisera. Further, Results from agglutination of O2, O4, O7, O9 and Vi antisera further identified 7 different serovars including *S. typhi*, *S. paratyphi* A, *S. paratyphi* B, *S. paratyphi* C, *S. typhimurium*, *S. choleraesuis*, and *S. enteritidis*.

The recovery rates of different serovars of *Salmonella* isolated from blood cultures of HIV/AIDS subjects and the control subjects are presented on Table 4. A total of 58 *Salmonella* isolates agglutinated the polyvalent H antisera used as genus specific antisera. These isolates were obtained from 41 HIV/AIDS patients and 4 control subjects. Some subjects had more than one serovar of *Salmonella*. *Salmonella typhi* was the most commonly observed *Salmonella* serovars identified from the samples in our study, making up 32% (17/53) of the *Salmonella* isolates in HIV/AIDS patients studied, and 40% (2/5) from the control subjects.

It was observed that irrespective of the HIV/AIDS status, subjects with low CD4⁺ cell counts of $\leq 200\mu\text{L}$ had higher percentages *Salmonella* associated bacteremia (Table 5). Data analysis showed a significant relationship ($p < 0.05$) between CD4⁺ cell counts of HIV/AIDS subjects and the prevalence of *Salmonella* bacteraemia.

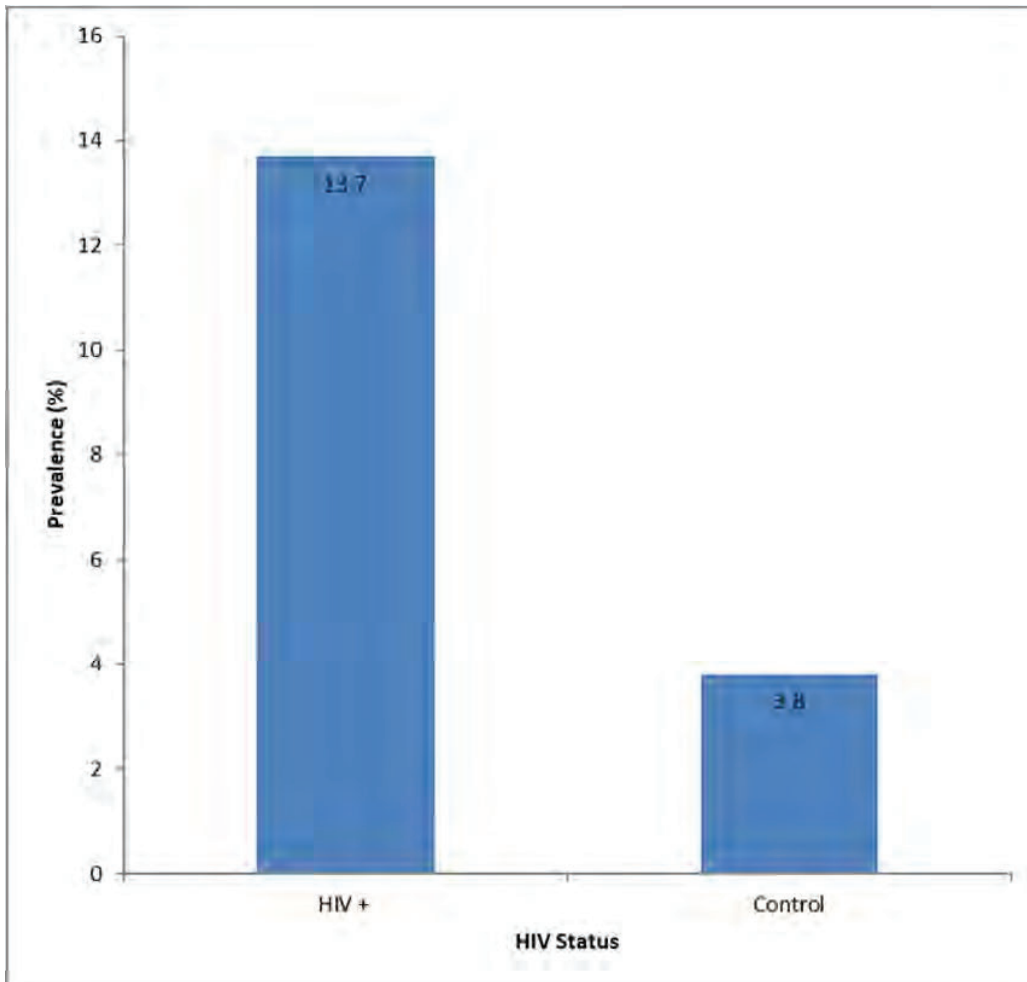


Figure 1: Prevalence of *Salmonella* associated bacteremia among HIV/AIDS patients and the control subjects in the study

Table 1. Biochemical characterizations and Microgen identification of the isolates obtained in the study

Isolates	Biochemical Tests														
	TDA	Cit	Mot	Oxi	Glu	Lys	Orni	Man	H2S	Xyl	ONPG	IT	UT	VP	Nit
S. spp	+	+	+	-	+	+	-	+	±	-	-	±	-	-	+
S. ariz	±	±	+	-	+	±	±	+	±	+	±	±	-	-	+

***S spp** : *Salmonella enterica* species, **S ariz**: *Salmonella enterica Arizonae*, **TDA**: Tryptophan deaminase test, **Cit**: Citrate Utilization test, **Mot**: Motility test, **Oxi**: Oxidase test, **Glu**: Glucose fermentation test, **Lys**: Lysine decarboxylase test, **Orth**: Orthinine decarboxylase test, **Man**: mannitol fermentation test, **H2S**: Hydrogen sulphide production test, **Xyl**: Xylose oxidation test, **ONPG**: beta-galactosidase test, **IT**: Indole production test, **UT**: Urease test **VP**: Voges Prokauer Test, **NT**: Nitrate reduction test. + = positive reaction, - = negative reaction, ± = variable.

Table 2. Biochemical reactions obtained from the serovars of *Salmonella* isolates in the study.

Salmonella serovars	Biochemical Tests and Their Reactions									
	KIA	Cit	Lys	Lac	Orth	ONPG	IT	H2S	Mot	
<i>S. typhi</i>	A	-	+	-	-	-	-	+	+	
<i>S. paratyphi A</i>	AG	-	-	-	-	+	-	-	+	
<i>S. paratyphi B</i>	AG	+	-	-	+	+	-	+	+	
<i>S. paratyphi C</i>	AG	+	+	-	-	-	+	+	+	
<i>S. typhimurium</i>	AG	+	+	-	+	+	+	+	+	
<i>S. choleraesius</i>	AG	+	+	-	-	+	+	-	+	
<i>S. enteritidis</i>	A	+	-	-	-	+	-	+	+	

***KIA**= Kligler iron agar test, **Cit**: Citrate Utilization test, **Lys**: Lysine decarboxylase test, **Lac**: Lactose fermentation, **Orth**: Orthinine decarboxylase test, **ONPG**: betagalactosidase, **IT**: Indole production test, **H2S**: Hydrogen sulphide production test, **Mot**: Motility test, + : Positive reaction, - : Negative reaction.

Table 3. Serotyping results of *Salmonella* isolates obtained from blood cultures

Antisera used	Serotypes of <i>Salmonella</i> species	'O' & Vi antigens possessed by the isolates	'H' antigens possessed by the isolates
O2	<i>S. paratyphi A</i>	1, 2, 12	a
O4	<i>S. paratyphi B</i>	1, 4, 5, 12	b
	<i>S. typhimurium</i>	1, 4, 5, 12	i
O7	<i>S. paratyphi C</i>	6, 7 (Vi)	c
	<i>S. choleraesius</i>	6, 7	y
O9	<i>S. typhi</i>	9, 12(Vi)	d
	<i>S. enteritidis</i>	4, 9, 12	g

Table 4: Recovery rate of *Salmonella* serovars from blood of HIV/AIDS subjects and the control subjects

<i>Salmonella</i> species	No. (%) of isolates from HIV/AIDS (n =53)	No. (%) of isolates from control subjects (n = 5)
<i>S. typhi</i>	17 (32.1)	2 (40.0)
<i>S. paratyphi A</i>	7 (13.2)	0 (0.00)
<i>S. paratyphi B</i>	9 (17.0)	1 (20.0)
<i>S. paratyphi C</i>	5 (9.4)	1 (20.0)
<i>S. typhimurium</i>	4 (7.6)	0 (0.00)
<i>S. choleraesius</i>	6 (11.3)	0 (0.00)
<i>S. enteritidis</i>	5 (9.4)	1 (20.0)

Table 5: Relationship between CD4⁺ cell counts and the distribution of *Salmonella* bacteraemia in HIV/AIDS patients.

CD4 ⁺ Counts/ μ L	(%) of HIV/AIDS subjects with <i>Salmonella</i> bacteraemia (n=41)	(%) of controls subjects with <i>Salmonella</i> bacteraemia (n=4)
\leq 200	21 (51.2)	2 (50.0)
201-300	11 (26.8)	1 (25.0)
301-400	3 (7.32)	1 (25.0)
401-500	2 (4.88)	0 (0.00)
501-600	3 (7.32)	0 (0.00)
601- 700	1 (2.44)	0 (0.00)
\geq 701	0 (0.00)	0 (0.00)

DISCUSSION

In this study, the 13.7% prevalence of *Salmonella* isolation from blood of HIV/AIDS subjects is of public health concern. Immunocompromised persons are more susceptible to *Salmonella* infection especially from its life-threatening bacteraemia (23). According to Keddy *et al.*, (2009) and Sharma *et al.*, (2010), *Salmonella* organisms especially *S. typhi*, infection in people living with HIV/AIDS may result in complications like septic shock, meningitis, and local abscess formation. The isolation of *S. typhi* and other serovars including the non-typhoidal ones in this study, corroborates the work of (25, 26) Taramasso *et al.*, (2016), Adisa *et al.*, (2017). *Salmonella* is known to cause serious infections ranging from febrile illness, typhoid fever, gastroenteritis and could lead to death (27). Previous studies have reported more than 2.16 million episodes of typhoid occurred worldwide, resulting in 216,000 deaths, and is described as a common systemic infection in tropical regions (28).

People living with HIV/AIDS are predisposed to bacteraemia and their systemic spread is uninhibited due to their defective adaptive immune system (29).

The isolation of serovars *S. typhimurium* and *S. enteritidis* agrees with the works of Brent, *et al.*, (2006) and Gordon *et al.*, (2010) who reported these organisms as the leading cause of community-acquired bloodstream and focal infection in Sub-Saharan Africa. Their study further highlighted that non-typhoidal *Salmonella* (NTS) serotypes were predominant among the HIV/AIDS infected persons and children with HIV, malaria, and malnutrition (30).

In this study, the majority of HIV/AIDS infected persons had low CD4⁺ cell counts with a higher recovery rate of *Salmonella* serovars. Other studies have reported that in HIV infection, the primary targets by the virus are the cells bearing CD4⁺ markers on their surfaces, hence the gradual decrease in CD4⁺ marker receptors (16-18). Fisk *et al.*, (2005) noted that the dysfunction observed with HIV/AIDS infected patients is usually as a result of opportunistic infections and the vulnerability to *Salmonella* infections which correlates inversely with the CD4⁺ lymphocyte count.

CONCLUSION

This study isolated two species of *Salmonella* and seven serovars from 13.7% of the HIV/AIDS infected persons in the study area, and their distribution largely depends on the level of CD4⁺ cell counts of the subjects. Therefore, emphasis should not be laid only on retroviral drugs for the HIV/AIDS infected persons, but also regular screening of the subjects for bacteraemia as well as advising the subjects to consume a balanced diet including fruits to boost their immune systems, maintain personal hygiene and sanitation coupled with provision of available *Salmonella* vaccines by Government to HIV/AIDS infected individuals so to eliminate *Salmonella* associated infections among this group.

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