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Occurrence of Salmonella serovars implicated in Salmonellosis in North-western part of Nigeria

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Abstract

Accurate diagnosis of clinical Salmonella enterica and differentiation into its serovars is important from a public health and epidemiological point of view. Standard cultural, biochemical and polymerase chain reaction techniques were employed to isolate and identify Salmonella enteric serovars implicated in salmonellosis. This study was carried out between June 2015 and May 2016 in some selected hospitals in North-Western Nigeria. Four hundred and twenty stool samples were collected from patients clinically diagnosed of pyrexia and/or gastroenteritis. Of the 420 samples collected, 27 were positive for Salmonella, given rise to an overall prevalence rate of 6.4%. One hundred and fifty two samples were collected from \leq 1-5yrs age-group and 19 (12.5%) were positive for Salmonella species. Out of the 199 male patients recruited for this study, 16 (8.0%) had Salmonella infection while 11 (5.0%) of 221 female patients recruited had Salmonella infection. Sex of the patients has no significant association with the infection (p>0.05). Out of the 27 Salmonella isolated, six different serovars were encountered. S. typhi and S. enteritidis were the most frequently encountered; given rise to 25.9% prevalence rate each. The second most frequently encountered was S. typhimurium with 18.5% prevalence rate. The least encountered serovar was S. paratyphi B which had 3.7% prevalence rate. Out of the 27 Salmonella isolates, 17 (63%) were isolated from patients presenting with mix symptoms and S.Typhi was the most frequently encountered, given rise to 35.3%. Seven isolates (25.9%) of the 27 Salmonella were isolated from patients presenting with diarrheoa only and S. enteritidis (57.1%) was the most frequently encountered. However, frequency of occurrences of serovars has no significant association to the salmonellosis symptoms (p>0.05). The prevalence rate in elderly patients (\geq 60yrs) was also high, for out of the 18 samples collected from them, 5 (27.8%) were positive. The age-group with least prevalence rate was 16-40yrs; all samples collected from them were negative for Salmonella infection. Salmonella infection was significantly associated with the age groups with p<0.05.Out of the 16 cases of salmonellosis in male patients, 7 (43.8%) were typhoidal, while 9 (56.3%) were nontyphoidal. Among female patients, 11 salmonellosis cases were recorded, 5 (45.5%) were typhoidal and 6 (54.5%) were non-typhoidal. This result showed that there was reduced in Salmonella infection in part of North Western Nigeria. However, there is need to sustain good sanitary practices in order to curtail increase and reduced children's exposure to the infection. Key Words: PCR; Salmonella Serovars; Salmonellosis; Prevalence.

INTRODUCTION

Salmonella enteric comprised of six subspecies; they are S. enteric subsp. enterica, S. enteritica subsp. salamae, S. enteric subsp. arizonae, S. enteric subsp. diarizonae, S. enteric subsp. indica, and S. enteric subsp. houtenae (Porwolliket al., 2003). Of these six subspecies, only S. enteric subsp. enterica is associated with disease in warm-blooded animals. As of 2002, there are over 2,541 serovars identified within S. enteric subsp. enterica (CDC, 2010). Salmonella species cause wide range of human diseases such as enteric fever, gastroenteritis and bacteriamia (Bennasar *et al.*, 2000).

Typhoid salmonellosis is a global problem with an estimated 12-33 million people infected in the world (Kohinur *et al.*, 2010). Non-typhoidal salmonellosis is self-limiting; however, it may lead to systemic symptoms in children, the elderly and immunocompromised (Rayamajhi *et al.*,2008). Bacteremia is reported to occur in 3 to 10 percent of cases of non-typhoidal salmonellosis and in such situations, antimicrobial therapy is lifesaving (White *et al.*, 2001).

Salmonella isolates are serotyped according to the Kaufmann-White scheme using somatic (O), capsular (VI) and flagellar (H) antigens that are present in the cell surface of Salmonella. The O factors determine the grouping, while the H factors completely define the serotype identity of a Salmonella strain (Popoff, 2001; Fitzgerald et al., 2003).

According to Herrera-Leon et al (2007), molecular serotyping methods are based on the detection of the same antigens as the Kauffmann-White Salmonella. Molecular methods including multiplex PCR offer a consistent and high-throughput approach to typing etiologic agents (Levy et al., 2008).

PCR identification of Salmonella is important or critical for surveillance, improving prevention and control of food-borne diseases. It has allowed rapid detection, identification of sources, control of outbreaks, and also identification of emerging serotypes and new mechanisms of transmission (Fritzgerald et al, 2003).

Therefore, as the salmonellosis becomes so prevalent and life-threatening within the study area couple with no or limited similar studies conducted more especially exploring the use of molecular approach as such the need for the research. However, the objective of this work was to detect and identify Salmonella enteric serovars implicated in salmonellosis in part of North-Western Nigeria using monoplex and multiplex PCR techniques.

MATERIAL AND METHODS

Sampling Sites

Six Hospitals were selected for this study in three states of North-Western Nigeria. These Hospitals were Hajia Gambo Sawaba General Hospital, Major Ibrahim Abdullahi B Memorial Hospital, Hasiya Bayero Public Hospital, Muhammadu Abdullahi Wase Specialist Hospital, General Hospital Dutsin-ma and General Hospital Daura. The Hospitals are located in Kaduna, Kano and Katsina States of North-Western Nigeria respectively.

Ethical Consideration, Patient's Consents, Inclusion and Exclusion Criteria:

Approvals for the study were obtained from Kaduna (with Ref NO: MON/ADM/744/VOL.1/286), Kano (with Ref NO: REF/OFF/797/T.I/24) and Katsina (with Ref NO: KHSMB/S.7/VOL.VII) States Ministry of Health respectively (See attached on appendix I-III). Patients' consents were obtained prior to collection of samples. The inclusion criteria were consented outpatients clinically diagnosed by clinicians of having either pyrexia, gastroenteritis or both. The exclusion criteria include non-consented patients and anyone that does not satisfy the inclusion criteria.

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Sample Size, Structured Questionnaire and Sample Collection

Seventy stool samples were collected from each of the hospital. This amounted to 420 from the six Hospitals. The samples were collected between January 2015 and December 2015. Structured questionnaires (attached on appendix IV) were administered to the consented patients to obtain information on the socio-demographic data and possible risk factors associated with Salmonella infection. Stool specimens were collected from outpatient clinically diagnosed by clinicians having either pyrexia, gastroenteritis or both.Patients were grouped into 5 Age-groups of $\leq 1-5$, 6-15, 16-40, 41-59and ≥60.

Pre- Enrichment and Enrichment of Specimen for Isolation

A loopful of stool sample collected from each patient was directly inoculated into 45ml of buffered peptone water and incubated at 37°C for 16hrs. Aloopful of culture from buffered peptone was inoculated into 9ml of selenite-F broth (Oxiod, Cambridge, UK) contained in test tubes (Akinyemi et al., 2007). This was incubated for 18hrs at 37°C for bacteriological culture.

Culture and Biochemical Characterization

A loopful of the culture from selinite-F broth was streaked on xylose lysine deoxycholate (XLD) agar (Oxoid, Cambridge, UK) and incubated overnight at 37°C. The plates of red with black centers colonies or red colonies only on XLD (OIE, 2010) were picked and subculture on XLD inorder to have pure culture for biochemical confirmation. The suspected colonies were then streaked onto Nutrient Agar Slant and incubated overnight at 37°C and later stored in refrigerator. The suspected Salmonella isolates were purified again, by streaking on XLD Agar plate from Nutrient Agar Slant and incubated overnight at 37°C. The pure isolated colonies were then used to characterized the organisms biochemically. Series of conventional biochemical testes wereconducted to srceen for Salmonella isolatesas recomended by WHO, (2010). Such biochemical tests were Urease production, Lysine Iron Agar (LIA), Triple Sugar Iron (TSI) Agar, Indole, Methyl Red and Voges-Proskauer tests.

DNA Extraction

DNA of the biochemically characterized Salmonella species were extracted using QuickgDNA[™] Miniprep (D3024). Two milliliter of an overnight grown Salmonella culture in Luria-Bertani Broth was transferred into 2ml micro centrifuge tube and spinned at 5000rpm for 10minutes.

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The supernatant was removed and 500µl of genomic lysis buffer was added directly to the cell pellet. The pellet was re-suspended by vortexing for 4-6 seconds and allowed to stand for 5-10minutes at room temperature. The mixture was then transferred to a Spincolumn in a collection tube. This was centrifuged at 10000rpm for one minute. The collection tube was discarded with flow through. The Spin column was transferred to a new collection tube and 200µl of DNA pre-wash buffer was added to it. This was then centrifuged at 10000rpm for 1minute. Five hundred microlitre of g-DNA buffer was added to the spin column and centrifuged at 10000rpm for 1minute. The spin column was then transferred to a microcentrifuge tube and 50µl DNA elution buffer was added to the spin column. This was incubated for 2-5minutes at room temperature and thereafter, centrifuged at top speed for 30 seconds to elude the DNA. The eluted DNA was stored at -20°C for monoplex and multiplex PCR confirmation and typing of Salmonella species respectively. All procedure according to manufacturers guidelines.

Monoplex PCRDNA Amplification

(Fset of primer pair GTGAAATTATCGCCACGTTCGGGCAA Rand TCATCGCACCGTCAAAGGAACC) with 284bp was used to confirm Salmonella species (Galan et al., 1992). It is specific for the invAgene located on the Salmonella pathogenicity island 1 (Galan et al., 1992; Rahnet al., 1992). The primer was commercially available from QIAGEN Operon. The primer was dissolved according to the instruction provided by the manufacturer. The amplification was performed in a final volume of 50µl in microamplification tube (PCR tube). The reaction mixture consisted of 5ml of the DNA template, 5ml of PCR buffer (75mM Tris-HCl, pH 9.0, 2mM MgCl2, 50mM KCl, 20mM (NH4)2SO4), 1ml deoxynucleoside triphosphate(dNTPs) (40 mM), 1ml (1U AmpliTaq DNA polymerase), 25 pmol of each primer (invAF, invA R), and the volume of the reaction mixture was completed to 50ml using DDW. The thermal cycler was adjusted as follows: Initial denaturation at 94°C for 5 min, followed by 35 cycles of (denaturation at 94°C for 1 min, annealing at 55°C for 1 min and extension at 72°C for 1 min). Final extension was carried out at 72°C for 10 min and the PCR products were stored in the thermal cycler at 4°C until they were collected.PCR products were resolved by electrophoresis in 1.5-2% (w/v) agarose gels and visualized under UV light after ethidium bromide staining (Sambrook et al., 1989) using 100bp molecular weight marker.

17 ISSN: 2616 - 0668 Multiplex PCR Typing of Salmonella enteric

Set of primer pairs were used. The primers were commercially available from QIAGEN Operon and Bioneer. The primers were dissolved according to the instruction provided by manufacturer. Primers for O-serogrouping were selected based on the *rfb* gene clusters specific for Salmonella serogroups A, B, C1, D and E, as well as primers based on viaB gene to detect Vi-positive strains as reported by Levy et al, (2008) and Herrera-Leon et al, (2007). Primers used in H typing were also designed by Levy et al. (2008). An internal control (primers P1-P2 that amplify oriC) was incorporated into the system as described by Levy et al, 2008. The multiplex PCR amplification was performed in a final volume of 25µl in micro-amplification tube (PCR tube). Each reaction contained 1X of PCR buffer, 25mM MgCl₂, 0.25mM deoxynucleoside triphosphates, 0.4µM (each) of primers prt-F, prt-R, rfbJ-R, vi-F, vi-R, wzxC1-F, wzxC1-R, tyvD-F, tyvD-R, wzxE-F, wzxE-R, 0.2µM of control primers (P1-P2) and 1.75U PromegaTaqDNA polymerase. Five microliters of a briefly centrifuged, boiled isolate suspension was used as template. PCR was performed in a Mastercycler with a heated lid cover. This prevented evaporation and eliminated the use of mineral oil. The cycling parameters of the PCRs were as follows: denaturation at 95°C for 5minutes, followed by 35 cycles at 95°C for 40 seconds, 50°C for 30 seconds and a final extension at 68°C for 7 minutes. A negative control consisting of double distilled water was included. Multiplex PCR products were separated in 1.5% agarose gel by unidirectional electrophoresis using 0.5X TBE buffer and visualized by staining with ethidium bromide. Fragments (amplicons) sizes were determined by comparison with 100bp DNA ladders. The serovars were then identified.

RESULTS

Serovars

The conventional biochemically suspected *Salmonella* isolates (48) were reconfirmed using monoplex PCR. Out of the 48biochemically suspected *Salmonella* isolates, 27 were molecularly confirmed to be *Salmonella* species given rise to overall prevalence rate of 6.4% (Table 1).

Out of the 199 samples collected from male patients, 16 (8.0%) had *Salmonella* infection. Two hundred and twenty one samples were collected from female patients and 11 (5.0%) had *Salmonella* infection (Fig. 2). Out of the 27 *Salmonella* isolated, S. *typhi* and S. *enteritidis* were the most frequently encountered, given rise to 25.9% prevalence rate each. The second most frequently encountered was S. *typhimurium* with 18.5% prevalence rate. The

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least encountered serovar was S. *paratyphi* B which had 3.7% prevalence rate (Table 2 and 3). Seventeen (63%) isolates were isolated from patients presenting with mix symptoms. Out of this 17 isolates, S. *typhi* was the most frequently encountered, given rise to 35.3%. Seven (25.9%) from the 27 isolates were found to be isolated from patients presenting with diarrhea only and S. *enteritidis* (57.1%) was the most frequently encountered (Table 4).

One hundred and fifty two samples were collected from ≤ 1 -5yrs age-group and 19 (12.5%) were positive for *Salmonella* species. Out of this 19 positive samples, 8 (42.1%) were typhoidal *Salmonellae* and 11 (57.9%) were non-

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typhoidal Salmonellae. The prevalence rate in elderly patients (\geq 60yrs) was also high, for out of the 18 samples collected from them, 5 (27.8%) were positive. Out of this 5 positive samples, 3 (60%) were typhoidal Salmonellae and 2 (40%) were non-typhoidal Salmonellae (Table 5).

The overall cases of salmonellosis in male patients were 16. Out of this 16 cases, 7 (43.8%) were typhoidal, while 9 (56.3%) were non-typhoidal. Among female patients, 11 salmonellosis cases were recorded, 5 (45.5%) were typhoidal and 6 (54.5%) were nontyphoidal (Fig. 5).

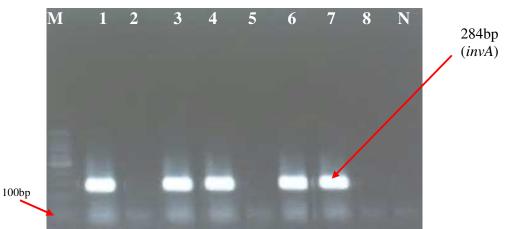
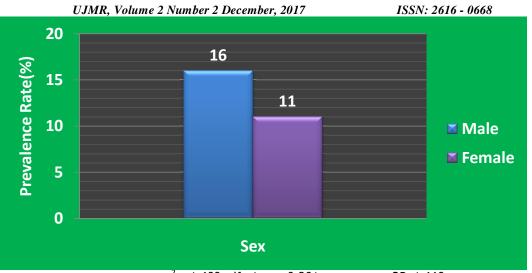


Fig. 1: One of the Electrophoresis Gel Pictures of Monoplex PCR amplified *invA* Genes of *Salmonella* Isolated from Hasiya Bayero Public Hospital (HB), Kano. Key: Lane M: PCR Ladder Maker (100bp); Lane 1, 3, 4, 6 and 7: Bands (Amplicons) of *invA* Genes; Lane 2, 5 and 8: No *invA* Genes; Lane N: Negative Control.

Table 1: Prevalence Rate of *Salmonella* Infection in the Selected Hospitals in Part of North-Western Nigeria.

Hospitals	No. of Samples	No. Positive (%)	
HG	70	6 (8.6)	
MA	70	3 (4.3)	
HB	70	5 (7.1)	
MW	70	5 (7.1)	
DT	70	6 (8.6)	
DA	70	2 (2.9)	
Total	42027 (6.4)		

Key: No.: Number, **HG**: Hajiya Gambo Sawaba Hospital Kaduna, **MA**: Mahor Ibrahim Abdullahi Hospital Kaduna, **HB**: Hasiya Bayero Public Hospital Kano, **MW**: Muhammadu Abdullahi Wase Specialist Hospital Kano, **DT**: General Hospital Dutsin-Ma Katsina, and **DA**: General Hospital Daura Katsina.



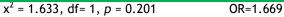


Fig. 2: Prevalence of Salmonella Infection in Respect of Sex in the Selected Hospitals in Part of North-Western Nigeria

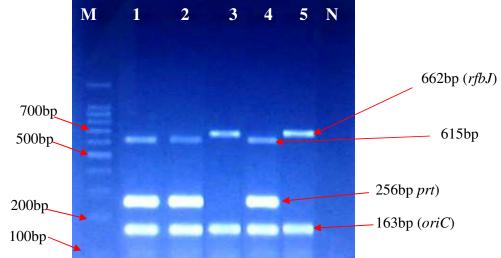


Fig. 3: One of the Electrophoresis Gel Pictures of Multiplex PCR Amplified O-groups, P1-P2 and Vi of *Salmonella* Isolated from Hasiya Bayero Public Hospital (HB), Kano. Key: Lane M: PCR Ladder Maker (100bp); Lane 1-5: Amplicons of O-Group and Vi genes; Lane N: Negative control.

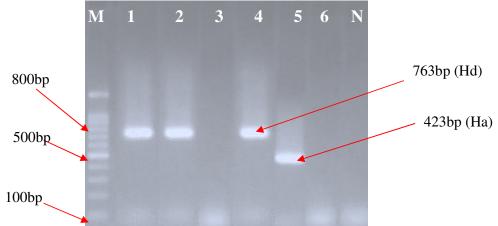


Fig. 4: One of the Electrophoresis Gel Pictures of Multiplex PCR Amplified Phase 1 Flagellar Antigens of
Salmonella Isolated from Hajia Gambo Sawaba General Hospital (HG), Zaria City, Zaria
Key: Lane M: PCR Ladder Maker (100bp); Lane 1, 2, 4 and 5: Amplicons of Phase 1 Flagellar Genes; Lane 3 and
6:No Phase 1 Flagellar genes; Lane N: Negative control.
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UJMR, Volume 2 Number 2 December, 2017 Table 2: Results of Multiplex PCR Salmonella Typing ISSN: 2616 - 0668

ID	P1-						0	Inference				
	P2	Ha	Hb	Hd	Prt	rfbJ	wzxC1	Туv	Vi	wzxE	Group	
HG29	+	-	-	+	+	-	-	+	+	-	D	S. typhi
HG35	+	-	-	+	+	-	-	+	+	-	D	S. typhi
HG37	+	-	-	-	+	-	-	+	-	-	D	S. enteritidis
HG41	+	-	-	+	+	-	-	+	+	-	D	S. typhi
HG63	+	+	-	-	+	-	-	-	-	-	Α	S. paratyphiA
HG70	+	-	-	-	-	+	-	-	-	-	В	S. typhim
MA3	+	-	-	+	+	-	-	+	+	-	D	S. typhi
MA18	+	-	-	-	+	-	-	+	-	-	D	S. enteritidis
MA23	+	-	-	-	+	-	-	+	-	-	D	S. enteritidis
HB10	+	-	-	-	+	-	-	+	-	-	D	S. enteritidis
HB15	+	-	-	-	+	-	-	+	-	-	D	S. enteritidis
HB21	+	-	-	-	-	+	-	-	-	-	В	S. typhim
HB41	+	-	-	-	+	-	-	+	-	-	D	S. enteritidis
HB46	+	-	+	-	-	+	-	-	-	-	В	S. paratyphiB
MW8	+	-	-	-	-	+	-	-	-	-	В	S. typhim
MW28	+	-	-	-	-	+	-	-	-	-	В	S. typhim
MW30	+	-	-	-	+	-	-	+	-	-	D	S. enteritidis
MW42	+	-	-	+	+	-	-	+	+	-	D	S. typhi
MW63	+	-	-	-	-	-	+	-	-	-	C1	S. infantis
DT7	+	+	-	-	+	-	-	-	-	-	Α	S.paratyphiA
DT29	+	+	-	-	+	-	-	-	-	-	Α	S.paratyphiA
DT53	+	-	-	-	-	+	-	-	-	-	В	S. typhim
DT58	+	-	-	+	+	-	-	+	+	-	D	S. typhi
DT59	+	-	-	-	-	-	+	-	-	-	C1	S. infantis
DT67	+	-	-	-	-	-	+	-	-	-	C1	S. infantis
DA46	+	-	-	+	+	-	-	+	+	-	D	S. typhi
DA68	+	+	-	-	+	-	-	-	-	-	Α	S.paratyphiA

Key: ID = Isolate Number, **HG**: Hajiya GamboSawaba Hospital Kaduna, **MA**: Mahor Ibrahim Abdullahi Hospital Kaduna, **HB**: HasiyaBayero Public Hospital Kano, **MW**: Muhammadu AbdullahiWase Specialist Hospital Kano, **DT**: General Hospital Dutsin-Ma Katsina, and **DA**: General Hospital Daura Katsina, S. *typhim* = S. *typhimurium*

Table 3: Distribution of Clinical Salmonella enteric Serovars in the Selected Hospitals in part of North-Western Nigeria

Hospital s	Ν	S. typhi	S.paratyphi A	S. paratyphi B	S. enteritidis	S. typhimurium	S. infantis
HG	6	3(50)	1(16.7)	0(0)	1(16.7)	1(16.7)	0(0)
MA	3	1(33.3)	0(0)	0(0)	2(66.7)	0(0)	0(0)
HB	5	0(0)	0(0)	1(20)	3(60)	1(20)	0(0)
MW	5	1(20)	0(0)	0(0)	1(20)	2(40)	1(20)
DT	6	1(16.7)	2(33.3)	0(0)	0(0)	1(16.7)	2(33.3)
DA	2	Ì(50)	ı́(50)	0(0)	0(0)	0(0)	0 (0)
Total	27	7(25.9)	4(14.8)	1(3.7)	7(25.9)	5(18.5)	3(11.1)

x² = 26.314, df = 25, *p* = 0.391

Key: N: Number of Isolates; Figures in parenthesis indicates percentages.**HG**: Hajiya Gambo Sawaba Hospital Kaduna, **MA**: Mahor Ibrahim Abdullahi Hospital Kaduna, **HB**: Hasiya Bayero Public Hospital Kano, **MW**: Muhammadu Abdullahi Wase Specialist Hospital Kano, **DT**: General Hospital Dutsin-Ma Katsina, and **DA**: General Hospital Daura Katsina

UJMR, Volume 2 Number 2 December, 2017 ISSN: 2616 - 0668 Table 4: Distribution of *Salmonella* Serovars in Respect to Salmonellosis Symptoms.

(Isolates=27)							
Symptoms	No. Isolates	S. Typhi	S. PA	S. PB	S. En	S. Tm	S. In
Fever	2(7.4)	0(0)	2(100)	0(0)	0(0)	0(0)	0(0)
Constipation	0(0)	0(0)	0(0)	0(0)	0(0)	0(0)	0(0)
Diarrhea	7(25.9)	0(0)	0(0)	0(0)	4(57.1)	2(28.6)	1(14.3)
Vomiting	1(3.7)	1(100)	0(0)	0(0)	0(0)	0(0)	0(0)
Mix-Symptoms	17(62.9)	6(35.3)	2(11.8)	1(5.9)	3(17.6)	3(17.6)	2(11.8)

x² = 21.908, df= 15, *p* = 0.110

Key: No.: Number; PA: Paratyphi A; PB: Paratyphi B; En: Enteritidis; Tm: Typhimurium; In: Infantis; Number in parenthesis indicates percentages

Table 5: Prevalence of Typhoidal and Non-Typhoidal Salmonellae Infection in Respect to Age-Group Patients in Selected Hospitals in Part of North-Western Nigeria.

Age-Groups	No. of Sample	No. Negative (%)	No. Positive (%)	Salmonellae Infection		
(yrs)				TS(%)	NTS(%)	
<1-5	152	133 (87.5)	19 (12.5)	8 (42.1)	11 (57.9)	
6-15	109	107 (98.2)	2 (1.8)	0 (0)	2 (100)	
16-40	106	106 (100)	0 (0)	0 (0)	0 (0)	
41-59	35	34 (97.1)	1 (2.9)	1 (100)	0 (0)	
≥60	18	13 (72.2)	5 (27.8)	3 (60)	2 (40)	
Total	420	393 (93.6)	27 (6.4)	12 (44.4)	15 (55.6)	

 $x^2 = 34.802$, df= 4, p = 0.000



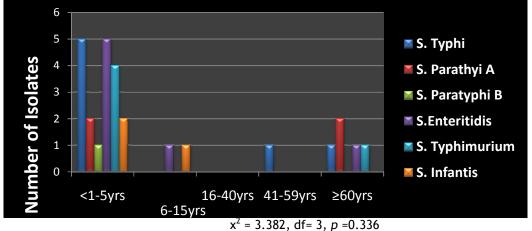


Fig. 5: Distribution of *Salmonella* Serovars among Age-Group Patients attending Selected Hospitals in Part of North-Western Nigeria.

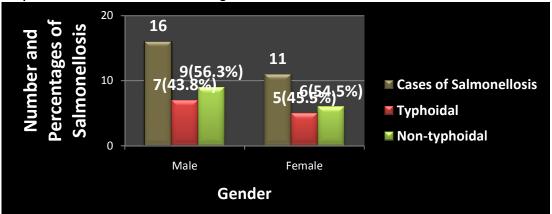


Fig. 6: Prevalence rate of Typhoidal and Non-TyphoidalSalmonellosis in Male and Female Patients attending Selected Hospitals in Part of North-Western Nigeria

DISCUSSION

The occurrence rate was lower in our study compared to other reports elsewhere in Nigeria; in Zaria, Gbonjubola *et al.* (2009) reported 14.1% cases, in Katsina State, Abdullahi *et al.* (2014) reported 15% cases and in Lagos, Muibat *et al.* (2015) reported 35% cases. The decrease in the occurrence (in this study) may be due to better awareness, more effective diagnosis and treatment. More so, out of the 70 samples collected from each of the Hospital, Hajia Gambo Sawaba

General Hospital Kaduna and General Hospital Dutsin-ma Katsina satate had 8.6% prevalence rate each, Hasiya Bayero Public Hospital and Muhammad Abdullahi Wase Specialist Hospital all from Kano state had 7.1% each and General Hospital, Daura from Katsina state had the least prevalence rate of 2.9%. However, there was no any significant association between the sampled Hospitals and the infection (p>0.05).

The frequency of the infection is higher in male (8.0%) than in female (5.0%) (Fig.2 and5). This finding was in agreement with the finding of Zailani et al. (2004), who reported higher frequency in male than in female. Also, Isa et al. (2013), reported higher prevalence rate in male (43.3%) than female (39.5%) in Biu, Borno, State, Nigeria. Although, there was no significant association between the infection and sex (p>0.05), the difference could be as a result of the fact that outdoor activities are more pronounced in males than females in North-Western Nigeria, there by exposing the males to high risk of infection. It may also be due to the fact that male are more exposed to infection because they tend to eat more road side and fast foods. Most reported cases of salmonellosis are due to contaminated food and water (Threlfall and Ward, 2001 and Wright et al, 2005).

In this studysix different serovars were encountered. S. typhi and S. enteritidis were the most frequently encountered serovars, given rise to 25.9% prevalence rate each. The second most frequently encountered was S. typhimurium with 18.5% prevalence rate (Table 3). This finding conformed to the reports ofAkinyemiet al, (2007) and Abdullahi et al (2014). In most parts of the world, surveys have reported S. enteritidis and S. typhimurium as the major serovars found in non-typhoidal salmonellosis (Antoine et al., 2008; Sabaet al., 2013). In Ghana and most other African countries, they are also the most frequently isolated from diarrheal diseases (Bonkoungou et al., 2013), which is confirmed by this study.

On the other hand, there was no significant association between distribution and frequency of occurrence of the serovars in the sampled hospitals with p>0.05. However, S. typhi was

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more encountered in HG than any other hospital and it was not isolated in HB hospital. Out of the 5 Salmonella isolates isolated in HB, were non-typhoidal salmonellae. Non-4 typhoidal salmonellae are usually implicated in non-typhoidal salmonellosis which is selflimiting without the use of antibiotics except in children, elderly and immunocompromised patients (Rayamajhi, et al., 2008). HB is a pediatric hospital and most patients encountered were children, this accounted for more isolation of non-typhoidal salmonellae than typhoidal salmonellae in the hospital.

Comparison of distribution of serovars in respect of salmonellosis symptoms revealed no significant difference in distribution with P>0.05(Tables 4). However, 17(62.9%) of the serovars were isolated from patients presenting with mixed-symptoms (fever, constipation, vomiting and diarrhea), while 7 (25.9%) were isolated from patients presenting with diarrhea only. Most people with salmonellosis develop diarrhea, fever, vomiting, and abdominal cramps 12 to 72 hours after infectionas reported by previous research (Santos *et al.*, 2001).

This study has shown that non-typhoidal salmonellosis was more prevalence than typhoidal salmonellosis with prevalence rate of 55.6% and 44.4% respectively (Table 5). This is in conformity with a study in Lagos-Nigeria by Akinyemiet al. (2012), who recorded 64.5% and 35.5% prevalence rate for non-typhoidal and typhoidal salmonellosis respectively but it is in contrast with Abdullahi et al. (2014), who recorded more prevalence rate of typhoidal (56.5%)than non-typhoidal salmonellosis (45.5%) in Katsina-Nigeria. The possible reasons for increase in non-typhoidal salmonellosis in part of North-Western Nigeria may be due to changes in cultural habits of eating which increase the patronage of fast foods.

Furthermore, in this study, there was significant association between salmonellosis and age groups with p < 0.05. Out of the 27 salmonellosis cases recorded, 19 (70.4%) were from age group <1-5yrs and of this value, 11 (57.9%) were non-typhoidal. The age group ≥60yrs had 5 (18.5%) cases while 16-40yrs had no cases recorded against them (Table 5). This outcome may be due to the fact that children and elderly people have weak immunity and are more likely to go down with salmonellosis than the middle age groups. It may also be that the middle age groups usually get over the infection and this may not necessitate them to seek medical attention. Also in this study, it was discovered that most middle age groups usually indulged in self-medication before seeking medical attention and this might have get rid of the infection before samples were

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collected for laboratory diagnosis. More so, age group <1-5yrs patients were more encountered than any other age group. It was also reported previously that the highest rate of *Salmonella* infection is found in infants (130 isolates/100,000). One quarter to one third of pediatric typhoid patients are younger than 5 years, of which 6-21% is younger than 2 years (Zaki and Karande, 2011).

All the six different serovars encountered in this study were isolated from age group <1-5yrs with S. typhi and S. *enteritidis* been the most frequently encountered. Four of the 6different serovars including *typhi* and *enteritidis* were isolated from age group \geq 60yrs with S. *paratyphi* A most frequently encountered. These findings support the report of Saba *et al.* (2013) and Paragrigorakis *et al.* (2007), they reported that S. *enteritidis* and S. *typhi* are the most encountered serovars in non-typhoidal and typhoidal salmonellosis respectively.

CONCLUSION

This study showed that children and elderly patients were the most susceptible to

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Salmonella infection. There was significant association between salmonellosis and age groups with p<0.05. In this study, non-typhoidal salmonellosis was more occurrence than typhoidal salmonellosis. We therefore, conclude that, there isneed to sustaingood sanitary practices in order to curtail increasein the infection. Provision of safe drinking and potable water as well as healthcare facilities by authorities at various levels is needed, in order to check mate any possible outbreak of the infection.

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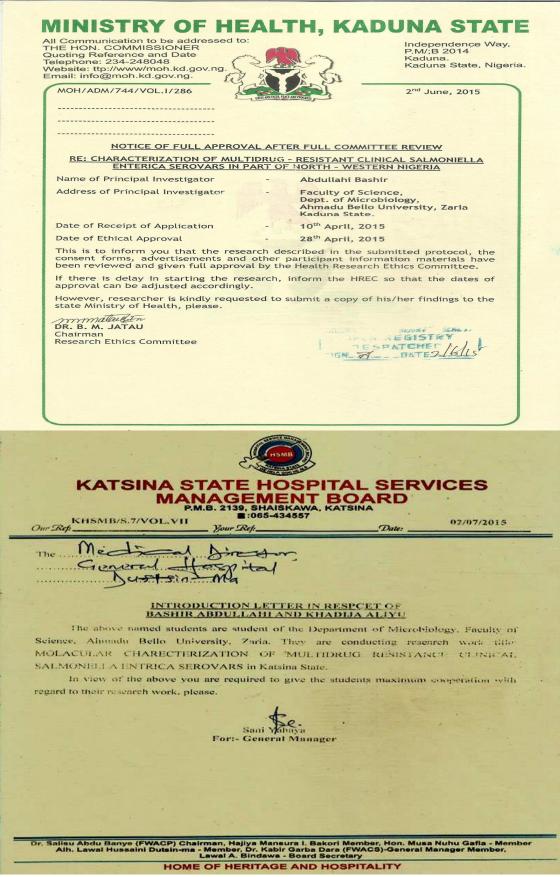
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Questionnaire!	Questionnaire!!	Questionnaire!!!
~	N	N

RESEARCH TITLE: "Molecular Characterization of Multidrug-Resistant Clinical Salmonella enterica Serovars in Part of North-Western Nigeria".

Patient number		Hospital			
Sex	Tick appropri	Age			
Yes 1. Fever	No If yes, for If yes, for If yes, for Borehole	<1wk or how long?	>2wk		
 8. Self Medication before Com 9. If 8 above is Yes, for how lo Ampicillin Cefotaxime Septri Clinical Signature/Date 	ng? days and wha	No It Antibiotic/s did you use Chloramphenicol Diagnosed	Other Dther B. Abdullahi		
			Investigator		