

Original Research Article

Primary Sources of Salmonella Species in Poultry Production Settings in Calabar, Cross River State, Nigeria

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Background: The poultry production settings have been frequently revealed as a major reservoir for Salmonella species. Salmonellosis is endemic in many countries, including Nigeria, where the poultry production industry is rapidly increasing. This study was designed to determine the gateways of entry of Salmonella into the poultry production setting so as to suggest measures to curb the introduction and maintenance of these organisms into the poultry production systems. **Methods:** Between the period of August 2013 and May 2014, 120 poultry feed samples were collected from commercial feed outlets, 120 water samples from poultry water storage tanks, 90 water samples from primary water sources (bore hole and pipe borne), 120 samples from both eggshell and the internal egg contents and 180 samples from day-old chicks within the poultry production settings in Calabar. Samples were collected aseptically and bacteriological analysis for the presence of Salmonella was performed according to the ISO 6579:2002 including standard bacteriological, biochemical and serologic techniques. The Chi-square parameter was used to determine the level of significance at 95% confidence level. **Results:** The rate of recovery of Salmonella species from the sources of poultry feed from commercial outlets, poultry water storage tanks and water from Primary water sources (bore hole and pipe borne) were respectively 7.5%, 11.7% and 3.3%. However, there was no statistical significant difference ($P\text{-value} = 0.084$) in the rate of recovery of Salmonella species from these sources, hence implicating these sources as primary routes for the entry of Salmonella species into the poultry production setting. There was also no statistical significant difference in the rate of recovery of Salmonella species from eggs and day-old chicks suggesting that the source of infection of day-old chick is from the incubation of contaminated eggs. **Conclusion:** The results of this study also suggest a relationship between eggshell contamination and internal egg content contamination as well as the contamination of the egg contents even before egg formation.

Keywords: Primary-sources, Salmonellosis, Eggshell-contamination, Internal-egg-content contamination

INTRODUCTION

The poultry production setting is a major reservoir for Salmonella species, the agent of Salmonellosis. Salmonellosis is still a major problematic zoonosis responsible for significant morbidity in humans and animals worldwide. In spite of the several attempts to minimize the causative agents to near zero tolerance, the disease remains a major threat to the commercial production of poultry and is endemic in Nigeria [1, 2]. Several outbreaks of salmonellosis in humans have been linked to the consumption of eggs, egg-containing foods,

poultry meats and other poultry products contaminated with the Salmonella organisms [3].

The increased outbreaks of human salmonellosis have been attributed to infection of breeder and layer flocks, the ability of some Salmonella serovars to cause infection of the ovaries and oviduct of birds and subsequently their eggs, the increased consumption of poultry and poultry products, and the abnormal storage temperature of eggs which allow the Salmonella organisms to increase in numbers [4,5,6]. The

vertical transmission of the infection from the ovaries to the eggs is termed trans-ovarian transmission. Apart from the host-adapted *Salmonella* serovars (*S. pullorum* and *S. gallinarum*), *S. enteritidis*, *S. typhimurium* and *S. heidelberg*, have been frequently implicated to infect eggs by trans-ovarian transmission [7,6].

However, this is not the only route by which eggs may become contaminated. In the poultry production setting, day-old chicks from hatcheries may become infected with *Salmonella* via contacts with formites and eggshells contaminated with faecal matter or dust and other debris present in poultry houses. *Salmonella* organisms belonging to a variety of serovars can contaminate eggs by penetration of the egg shell into the internal egg contents [8,9] or by trans-ovarian transmission [6]. The surpluses of hatching eggs may be diverted from incubation to be used as table eggs. When such eggs are contaminated with the *Salmonella* organisms they become a source of human infections when consumed [10].

The growth and survival of *Salmonella* species in water can be maintained given the right conditions. Water reservoirs for poultry use can readily favour the growth and maintenance of *Salmonella* species consequently leading to the infection of birds in the flock. Feeds and feed ingredients amongst others are important sources whereby the *Salmonella* organisms can be introduced in the poultry production setting with the consequent infection of the entire flock as well as humans and other animals [11, 12].

In order therefore to control *Salmonella* and other pathogenic microorganisms to near zero tolerance in the poultry production setting, strict hygiene and sanitation measures should be taken into account at every stage involved in the production of poultry. This study was therefore aimed at investigating the major gateways of introduction and maintenance of the *Salmonella* organisms in the poultry production setting by determining the rate of contamination of feeds from commercial outlets, poultry water reservoirs and water from primary sources as well as eggs and day-old chicks as primary sources of *Salmonella* species.

MATERIALS AND METHODS

This study was conducted in Calabar, Cross River State, Nigeria between August 2013 and May 2014. A total of 630 specimens consisting of 180 from day-old chicks, 120 egg pools, 120 from poultry feed in commercial outlets, 120 from poultry water reservoirs and 90 from primary water sources (Pipe-born and bore-hole) located within the Calabar Metropolis.

Sample collection

The day-old chicks were collected from 6 hatcheries and were immediately analyzed upon arrival in the Pentamed Scientifics Laboratory (No. 50 Atamunu, Calabar, Nigeria). Cloacal swabs and organs (liver and yolk sac) were collected from each day-old chick. 10 g of poultry feed was collected from commercial feed outlets into dry sterile polythene bags. 100 ml of water was collected aseptically directly from the sources and transported to the Laboratory within 4 h. Eggshell samples were collected by placing the intact egg in a sterile polythene bag containing 10 ml modified Buffered Peptone Water (10%) (HARDY Diagnostics, 1430 West McCoy Lane, Santa Maria, CA 93455, USA) and the egg was thoroughly rubbed through the bag for 1 minute without breaking it [13]. The diluents were then used as the eggshell sample and incubated at 37°C for 18

hours the internal egg contents were collected aseptically by first disinfecting the eggshell with 30% hydrogen peroxide (H₂O₂) solution, followed by 70% ethanol and burning off the alcohol briefly. The disinfected egg was broken using a sterile blade and sanitized plastic gloves. The egg content was enriched in modified Buffered Peptone Water and incubated at 37°C for 18h [13]

Isolation and identification of *Salmonella* species

The isolation of *Salmonella* species from poultry feeds, poultry water reserves and day-old chicks and eggs was carried out in accordance with the ISO 6579:2002. The reagents were obtained from HARDY Diagnostics, 1430 West McCoy Lane, Santa Maria, CA 93455, USA. By means of a sterile pipette, 1 ml and 0.1 ml of the non-selective enrichment (modified Buffered Peptone Water) was inoculated into 9 ml of Muller-Kauffmann Tetrathionate-novobiocin (MKTTn) and Modified Semi-solid Rapaport-Vassiliadis (MSRV) selective enrichments respectively. These were respectively incubated at 37°C and 42°C in separate incubators for 24 h.

After incubation, a loop full from each of the selective enrichments was streaked onto both Brilliant Green Agar (BGA) and Xylose Lysine Desoxycholate Agar (XLDA) selective plates in order to obtain distinct *Salmonella*-like colonies. On BGA, typical *Salmonella*-like colonies appear as 1-2 mm pink colonies which convert the agar from green to red, whereas on XLDA, they appear as pink colonies with or without the presence of black centres (indicating the production of H₂S).

The typical *Salmonella*-like colonies on BGA and XLDA were then picked by means of a sterile wire needle in order to make smears on microscope slides for gram staining as well as to stab and streak on pre-prepared Triple Sugar Iron Agar (TSIA) and Christensen agar (CA) slants. Those *Salmonella*-like colonies that showed Gram-negative small rods by microscopy, produced alkaline slope/acid butt with or without the production of H₂S and gas, on TSIA slant and Urease negative on Christensen agar slant were considered suggestive of *Salmonella*. They were then sub-cultured on nutrient agar slants overlaid with sterile paraffin oil and kept in a cool, dark corner prior to further biochemical reactions [14].

Confirmation of *Salmonella* species was conducted based on the standard biochemical techniques in order to identify the isolates which belong to the genus *Salmonella*. This involved the use of Lysine Decarboxylation (LCD) test, β-galactosidase test, Acetone production test and Indole production test. Serotyping of obtained *Salmonella* isolates was further supplemented by means of the commercially available polyvalent *Salmonella* antisera kit (Denka Seiken Co. Ltd. Tokyo, Japan) specific for all group and type-factor *Salmonella* antigens. A loop full from *Salmonella* isolates that satisfy all the confirmation procedures were then emulsified with a drop of normal saline (0.85% NaCl) on a microscopic glass slide [14]. The preparation was gently stirred and observed for auto-agglutination. If there was no self-agglutination, a drop of the polyvalent antisera was added and gently agitated by rocking back and forth for about three minutes and observed for agglutination. Those that showed agglutination were considered to belong to the genus *Salmonella*.

Statistical Analysis

The data obtained in this study were analyzed by means of the Predictive Analytical Software (PASW) 18.0. The simple descriptive statistics was used to analyse the prevalence rate

of *Salmonella* species in the samples obtained from the production of poultry while the level of significance in the rate of recovery of the *Salmonella* species was determined using the Chi-square parameter. *P-values* of less than 0.05 were considered statistically significant.

RESULTS

Out of 330 samples consisting of poultry feed from commercial outlets (120), poultry drinking water from poultry water storage tanks (120) and water from primary sources (pipe-borne and bore-hole) (90), *Salmonella* isolates were respectively recovered thus: 7.5%, 11.7% and 3.3% (Table 1). There was no statistical significant difference in the rate of recovery of *Salmonella* isolates from these sources (*P-Value* = 0.084).

According to table 2, the rate of recovery of *Salmonella* isolates from egg samples and day-old chicks from hatcheries were 10.0% and 7.2% respectively. However, there was no statistical significant difference in the rate of *Salmonella* species from eggs and day-old chick samples (*P-value* = 0.402).

The Venn diagram shows that *Salmonella* species were recovered from $27 + 12 = 39$ (32.5%) of eggshells and $2 + 12 = 14$ (11.7%) of egg contents. 2 (1.7%) of the egg contents alone were contaminated with *Salmonella* species, while 27 (22.5%) of the egg shells alone were contaminated with *Salmonella* species. Furthermore, *Salmonella* species were recovered from 12 (10.0%) of both egg shells and egg contents. The differences in the rate of recovery of *Salmonella* isolates from the egg shells and the egg contents was statistically significant (*P-Value* = 0.000).

DISCUSSION

Salmonellosis is a significant problematic zoonosis worldwide from the point of view of public health and it accounts significantly to the drawbacks encountered by food production systems especially in poultry and poultry products [15]. The bacteriological analysis of the samples from commercial feeds, poultry water reservoirs and primary water sources (pipe-borne and bore-hole) revealed low rates of recovery of *Salmonella* species and this was in conformity with the results of the studies conducted in Zaria, Nigeria and Jamaica [12,16].

According to the results of this present study, in spite of the least rate of recovery of *Salmonella* isolates from the primary water sources (borehole and pipe-borne), there was no statistical significant difference in the rate of recovery of *Salmonella* species (*P-Value* = 0.084). This suggests that these sample sources (poultry feed from commercial outlets, poultry drinking water from reservoirs and water from primary sources) serve as important portals of entry of the *Salmonella* organisms into the poultry production system.

Therefore, it would be strongly recommended that water for poultry use should be treated. This is in conformity with the work of other researchers who demonstrated a significant difference in the *Salmonella* contamination rate of treated (1.0%) and untreated (7.5%) water for poultry use [12]. Moreover, the feed production industries and poultry feed vendors should observe strict hygiene and sanitation measures during feed processing, transportation and storage in order to minimise bacteriological hazards. Trans-ovarian contamination occurs when the *Salmonella* organisms reside in the reproductive tissue of an infected bird and are transferred to the internal contents of the egg during egg formation [17]. The

microbiological analysis of *Salmonella* from eggs and day-old chicks as revealed by this study correspond with the studies carried out by other workers [17] and in Brazil who isolated *Salmonella* species from day-old chicks and table eggs [18]. However, other workers showed considerably higher rates (37.5% and 24.3%, respectively) of recovery of *Salmonella* species from day-old chicks [19, 20]. Under normal circumstances, eggs are sterile and hence day old chicks. However, this current study revealed 7.2% rate of recovery of *Salmonella* species from day-old chicks. The results of this study, therefore, suggest two possible sources of the *Salmonella* infection, which is in agreement with literatures that eggs may become contaminated with the *Salmonella* organisms via vertical transmission (trans-ovarian transmission) and/or the contamination of the egg, post egg formation with the subsequent penetration of the *Salmonella* organisms into the internal egg contents [17].

The results of this study showed that *Salmonella* species were recovered from 10.0% of eggs destined for incubation and there was no statistical significant difference in the rate of recovery of *Salmonella* species from eggs destined for incubation and from day-old chicks (*P-value* = 0.402). This invariably suggests that, infection of day-old chicks is dependent upon the contamination of eggs destined for incubation and as a consequence serves as an important portal for the entry of the *Salmonella* species into the poultry production systems. On the other hand, the difference in the rate of recovery of *Salmonella* isolates from egg shells (32.5%) and egg contents (11.7%) in this current study was statistically significant (*P-value* of 0.000), implying that the contamination of eggshell is independent from the of contamination of egg contents.

This is in conformity with the works of Albuquerque *et al* and Gantois *et al* who distinguished between the trans-ovarian contamination of eggs from the contamination of egg post egg formation [8,17]. Trans-ovarian contamination usually occurs before or during egg formation in the oviduct while the contamination of eggshell after egg formation usually occurs when the formed egg makes contact with contaminated faeces in the cloaca during oviposition or when in contact with contaminated debris or matter found in the poultry setting post oviposition [8, 17]. However, *Salmonella* species were recovered from both the eggshell and the egg content, in 12 (10%) of the egg samples used in this study, hence suggesting the source of the contamination of the internal egg contents could be traced from the contaminated eggshell.

Several studies have demonstrated the penetration of *Salmonella* serovars from the eggshell into the egg contents [6, 17]. This implies that a contaminated environment can lead to the contamination of eggshell with the subsequent contamination of the internal egg contents. The contamination of eggs destined for incubation consequently leads to the production of infection of day-old chicks and hence the entry of *Salmonella* species in the poultry production setting. In like manner, when such eggs are diverted for food, this may lead to the infection of the consumers.

Finally, the results of this study revealed that only 2 (1.7%) of the egg content samples were contaminated with *Salmonella* species without eggshell contamination. Such contamination of internal egg contents without eggshell contamination can most likely be implicated to have occurred at the level of the oviduct rather than post egg formation, hence trans-ovarian transmission.

Table 1: Distribution of Salmonella species by poultry feed and water sources

Source of sample	Number of sample tested	Number positive of sample	Percentage (%) of sample positive
Poultry feed from commercial outlets	120	9	7.5
Poultry water reservoirs	120	14	11.7
Primary water source	90	3	3.3
Total	330	26	7.9

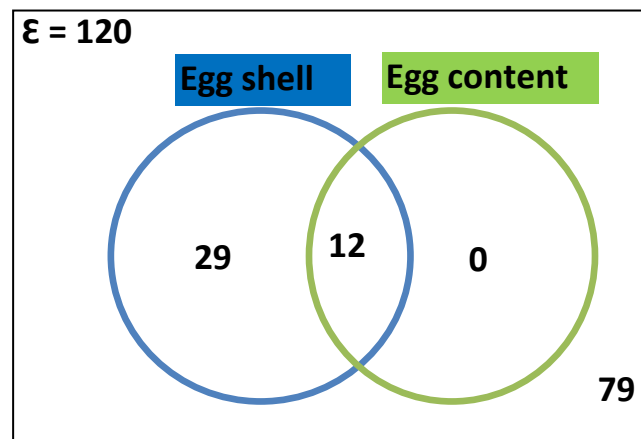
$\chi^2 = 4.958$, $df = 2$, $P\text{-value} = 0.084$ (two-tailed).

Table 2: Distribution of Salmonella species by eggs and day-old chicks from hatcheries

Type of sample	Number examined	Number infected (%)
Eggs	120	12 (10.0)
Day-old chicks	180	13 (7.2)
Total	300	25 (8.3)

$\chi^2 = 0.727$, $df = 1$, $P\text{-value} = 0.402$ (two-tailed) (Fisher's Exact Test for 2 X 2 tables)

Venn diagram of the distribution of Salmonella species in egg contents and egg shell



$\chi^2 = 20.458$, $df = 1$, $P\text{-value} = 0.000$ (two-tailed) (Fisher's Exact Test for 2 X 2 tables)

CONCLUSION

The results of this study have demonstrated that in spite of the major advances in technology in the appropriation of adequate sanitation, there is yet a dearth in the implementation of these strategies. The water and feed given to the poultry birds are frequently contaminated which poses both the birds and the consumers at risk of the infection. Eggs may become contaminated prior or post oviposition and consequently infection of chicks from hatcheries as well as infection to the consumers of egg and egg products.

It is therefore necessary to implement standardized bacteriological hazard control points at the level of feed production, transportation and storage. It would also be necessary to appropriately treat water for poultry use. Chicks and layer flocks should be screened for bacteriological hazards before introduction into the poultry production setting. There is

also a need for a surveillance control mechanism to monitor and check these control strategies in order to ultimately minimize or eliminate the hazard of *Salmonella* contaminations and hence curb human salmonellosis.

AUTHORS' CONTRIBUTIONS

NY Y and BEB conceived the study; carried out the statistical analyses; field interviews; and drafted the manuscript. Authors read and approved the final manuscript.

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Competing Interests

None declared.

Ethical Approval

Not required

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