



Isolation and Serological Identification of Current *Salmonella* Species Recovered from Broiler Chickens in Egypt

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ABSTRACT

This study was carried out to identify and serotype *Salmonella* (*S.*) species from suspected broiler chickens and healthy chickens in seven Egyptian governorates (KafrEl-sheikh, Fayoum, Qalubia, Menofiya, Garbia, Cairo, and Giza). Five hundred specimens (liver, spleen, caecum, and bone marrow) were collected from poultry farms and slaughterhouses. Samples were only collected from slaughterhouses in Cairo and Giza governorates. Forty-nine *Salmonella* isolates were recovered, and their colonies were characterized on specific media. Those *Salmonella* isolates were morphologically and biochemically identified. Furthermore, 33 out of 500 samples were serologically confirmed as *Salmonella* serovars by the slide agglutination test based on O and H antigen using 140 commercial polyvalent and monovalent diagnostic *Salmonella* antisera with a prevalence rate of 6.6%. They were serotyped as *S. blegdam* (39.4%), *S. typhimurium* (21.2%), *S. montevideo* (9.1%), *S. agama* (6.1%), *S. gueuletapee* (6.1%), *S. salamae* (6.1%), *S. enteritidis* (3%), *S. infantis* (3%), *S. kentucky* (3%), and *S. virchow* (3%). The prevalence of *Salmonella* serovars in KafrEl-sheikh, Fayoum, Qalubia, Menofiya, Garbia, Cairo, and Giza was 8.5%, 4%, 8%, 12%, 8%, 0%, and 0%, respectively. Finally, continuous monitoring of *Salmonella* serovars in chickens is essential for better prevention and reduction of its zoonotic risk.

Key words: *Salmonella*, Serovars, Isolation, Biochemical identification, Serological identification, Broilers.

INTRODUCTION

Salmonellosis has a great economic impact in poultry industry, due to high morbidity, mortality and reduced production (Khan et al. 1998; Rostagno et al. 2006; Abd El-Hack et al. 2021).

Also, Salmonellosis has a great zoonotic and public health importance in Egypt and worldwide. It causes an important zoonotic food borne disease characterized by mortalities, gastroenteritis, and/or septicemia in humans (Majowicz et al. 2010; Newell et al. 2010; Eng et al. 2015; Balasubramanian et al. 2019; Girh et al. 2019; Haley et al. 2019; Wang et al. 2020). Chickens were considered the main source of *Salmonella* outbreaks because they act as carriers of this pathogen in their guts. The main reservoir for *Salmonellae* is contaminated poultry and eggs (Antunes et al. 2016). Salmonellosis in poultry may occur by one or more strains of genus *Salmonella* either in acute or chronic form (Hofstad et al. 1992). The disease in chickens

characterized by significant economic losses due to high morbidity, mortality especially in young chicks, and reduced production (Khan et al. 1998; Rostagno et al. 2006). Additionally, *Salmonellae* can be transmitted vertically from broiler breeder chickens to their progeny (Lister 1988; Barbour et al. 1999; Wibisono et al. 2020).

Salmonellae are gram-negative, facultatively anaerobic, and non-spore-forming bacteria belong to family Enterobacteriaceae. They biochemically produce carbon dioxide and hydrogen gases from D-glucose, and typically hydrogen sulfide is produced by most *Salmonellae* species. All *Salmonellae* are aerogenic except for *Salmonella* serovar Typhi which never produces gas which does not produce urease, oxidase, and indole (Popoff and LeMinor 2001). About 2600 different *Salmonella* serovars have been described worldwide (Mezal et al. 2014). These serovars were serologically identified according to their somatic (O) and flagellar (H) antigens (Amagliani et al. 2012). European Food Safety Authority

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(EFSA) stated the prevalence of *S. infantis* (29.2%) and *S. enteritidis* (13.6%), followed by *S. kentucky* (6.2%) and *S. typhimurium* (4.4%) as the most isolated *Salmonella* serotypes in broiler chickens in European union (EFSA, 2010). Furthermore, different *Salmonella* serovars have been reported in Egypt (Abd El-Ghany et al. 2012; Halawa et al. 2016; El-Sharkawy et al. 2017; Mahmoud et al. 2018; Elkenany et al. 2019; Awad et al. 2020). Thus, continuous isolation and characterization of the most prevalent serovars are essential for improvement of salmonellosis prevention and reduce its public health hazard.

Thus, this study aimed to isolate and characterize the most prevalent *Salmonella* serovars from broiler chicken's farms and slaughter houses in 7 governorates in Egypt in the period between December 2019 till May 2021.

MATERIALS AND METHODS

Ethical Approval

This study was approved by Ethical Committee for Medical Research at the National Research Centre, Egypt (19157).

Chicken Flocks

This study was conducted on 500 samples of freshly dead chickens showed diarrhea from 5 governorates (KafrEl-sheikh, Fayuom, Qalubia, Menofiya, and Garbia), and freshly slaughtered broiler chickens from 2 governorates (Cairo and Giza). All birds were subjected to postmortem examination.

Bacterial isolation and identification

Sampling

For bacteriological examination, a total number (no.) of 500 samples were collected from (liver, spleen, caecum, and bone marrow) from freshly dead broiler chickens in 5 governorates: KafrEl-sheikh (no.200), Fayuom (no.50), Qalubia (no.50), Menofiya (no.50) and Garbia (no.50). Samples from freshly slaughtered broiler chickens were also collected from Cairo and Giza governorates (no.100). All samples were labeled and transported in cool boxes to the laboratory.

Culture Media

Bacterial enrichment was carried out using pre-enrichment buffered peptone water (Oxoid, UK) and incubated at 37°C for 18 hours for *Salmonella* spp. A total amount of 0.1 ml of the pre-enriched culture of suspected *Salmonella* spp. were inoculated into enrichment Rappaport Vassiliadis (RV) broth (Oxoid, UK) at 41°C for 24 h. A loopful of the enriched broth was streaked onto (S.S) agar (Oxoid) and (XLD) agar (Oxoid, UK) and incubated at 37°C for 24 h. Then, the growing colonies were examined for bacterial growth according to (Quinn et al. 1994) and (Collee et al. 1996). The suspected colonies (pink colonies with or without black center on XLD and colorless colonies with black centre on S.S agar) were picked up and purified for further investigation (Quinn et al. 1994). A loopful of the colony was stained with Gram stain and examined under a microscope according to Merchant and Packer (1967). Then kept in semi-solid agar for biochemical analysis (Quinn et al. 2002).

Biochemical Identification

The proper biochemical characterization was carried out using triple sugar iron agar (TSI), urea splitting ability in Christensen's urea agar, carbon utilization in Simmon's citrate agar, lysine iron agar and indole production in tryptone broth (Sneath et al. 1986).

Serological Typing of *Salmonella* Species

The obtained *Salmonella* isolates were subjected to serological identification according to Edward and Ewing (1972) through the application of a standard slide agglutination test based on O and H antigen using 140 commercial polyvalent and monovalent diagnostic *Salmonella* antisera (SINIF Co., Germany).

Statistical Analysis for Prevalence Calculation

The prevalence (P) in percentage was calculated using the formula $P = d/n$, where d is the number of positive samples analyzed at that point in time and n is the total number of chickens sampled at that point in time according to Thrusfield (2005).

RESULTS

Isolation of *Salmonella* spp. was attempted from 500 field samples (liver, spleen, caecum, and bone marrow) that were collected from poultry farms and poultry slaughterhouses of broiler chickens from seven Egyptian governorates (KafrEl-sheikh, Fayuom, Qalubia, Menofiya, Garbia, Cairo and Giza). Postmortem examination of the collected samples from freshly dead birds was shown nodular myocarditis, pericarditis and hepatitis. However, samples which collected from slaughterhouses had no lesions.

Salmonella suspected isolates showed smooth red colored colonies with black center on XLD and (S-S) agar. *Salmonella* produces colorless colonies with black centers due to H₂S production. Our result revealed that a suspected 49 *Salmonella* isolates out of 500 collected samples were detected by morphological and bacteriological examination. The isolated bacteria were Gram-negative, non-spore-forming and short rod-shaped single or paired in an arrangement under the microscope. These isolates were subjected to further biochemical identification where the suspected *Salmonella* isolates were positive to TSI, carbon utilization in Simmon's citrate agar and lysine iron agar but negative to urea and indole tests as in Table 1. Results of biochemical identification revealed that 49 suspected *Salmonella* isolates were biochemically positive.

Further serological identification using the slide agglutination test based on O and H antigen using 140 commercial polyvalent and monovalent diagnostic *Salmonella* antisera revealed that 33 *Salmonella* serovars were confirmed as *Salmonella* with a prevalence rate of 6.6% (33/500) as in Table 2. A total number of 33 *Salmonella* serovars were serotyped as *S. blegdam* (13 isolates), *S. typhimurium* (7 isolates), *S. montevideo* (3 isolates), *S. agama* (2 isolates), *S. gueuletapee* (2 isolates), *S. salamae* (2 isolates), *S. enteritidis* (1 isolate), *S. infantis* (1 isolate), *S. kentucky* (1 isolate), and *S. virchow* (1 isolate) as shown in Fig. 1. Results of serotyping demonstrated that the most predominant serotype was *S. blegdam* followed by *S. typhimurium* then *S. Montevideo*. On the

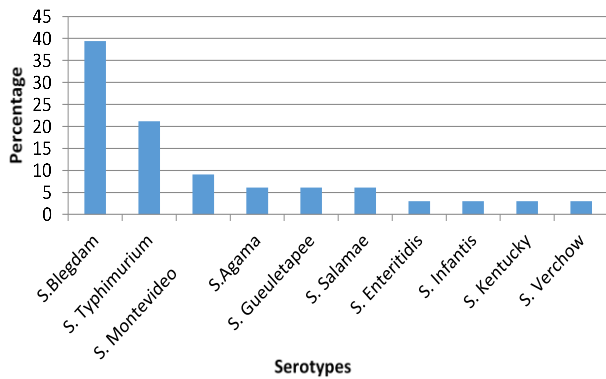


Fig. 1: Percent of serotyped *Salmonella* serovars.

Table 1: Biochemical profile of *Salmonella* isolates

Test	Result
TSI	+(Alkaline)\Slant\Acid bed\H ₂ S
Urease	-
Simmons citrate	+
Lysine	+
Indoie	-

TSI: triple sugar iron agar, H₂S: Hydrogen sulfide production, +: positive reaction, -: negative reaction

Table 2: Prevalence of *Salmonella* isolates in different governorates

Governorates	Number of Identified samples	Number of Identified Salmonellae	Percent
KafrElsheikh	200	17	8.5
Fayoum	50	2	4
Qalubia	50	4	8
Menofiya	50	6	12
Garbia	50	4	8
Cairo & Giza Poultry Slaughter houses	100	0	0
Total	500	33	6.6

other hand, *S. agama*, *S. gueuletapee*, and *S. salamae* had the same ratio. As well as *S. enteritidis*, *S. infantis*, *S. kentucky* and *S. Verchow* had the same and the least ratio of isolation.

The prevalence of *Salmonella* spp. in each governorate was shown in Table 2. Our result revealed that the high prevalence rate was detected in Kafr El-sheikh governorate (8.5%) and the lowest prevalence rate was detected in Cairo and Giza poultry slaughterhouse (0%).

DISCUSSION

Salmonella infection in poultry is considered one of the most important bacterial diseases causing heavy economic losses through mortality and reduced production (Haider et al. 2004; Rajan et al. 2017). Chickens can be infected with a wide variety of *Salmonella* serovars. Some serovars are host-specific for chicken such as *S. Pullorum* and *S. Gallinarum*, while other serovars are capable of infecting a wide range of hosts as *S. typhimurium* and *S. enteritidis* (Foley et al. 2008; Girh et al. 2018). Although there were more than 2600 identified serovars, identification and classification of *Salmonella* isolates into different serovars remain as a critical topic for studying the prevalence and surveillance. On the other hand, Salmonellosis is considered a major food borne pathogen in most countries of the world especially in developing countries (Soultose et al. 2003; Carraminana et al. 2004;

Chai et al. 2017; Wessels et al. 2021; Ehuwa et al. 2021; EL-Saadony et al. 2022).

In the current study, isolation of *Salmonella* spp. was carried out from 500 field samples (liver, spleen, caecum, and bone marrow). All samples were collected from poultry farms and slaughterhouses of broiler chickens from seven Egyptian governorates. *Salmonella* suspected isolates showed smooth red colored colonies with black center on XLD agar while on (S-S) agar. *Salmonella* produces colorless colonies with black centers due to H₂S production. It has been previously reported that pink colonies with or without black centers were typical for *Salmonella* on XLD. Many cultures of *Salmonella* spp. may also produce large colonies with glossy black centers or may appear as almost completely black (Sujatha et al. 2003; Ramya et al. 2012; Rabie et al. 2012; Islam et al. 2016). Our results revealed that a suspected 49 *Salmonella* isolates out of 500 collected samples were detected by morphological and biochemical examination (Table 1).

Previously, *Salmonella* were recovered from 4 broiler chicken flocks from Qalubia governorate in an incidence rate 3.84%, 4.15%, 5.06% and 5.18% (Abd El-Ghany et al. 2012). Furthermore, an incidence rate 7.5% of *Salmonella* have been detected in broiler chicks during 1st week of life in 5 Egyptian provinces (Sedeik et al. 2019). Our results were also in accordance to Samanta et al. (2014) who identified 22 isolates (6.1%) of *Salmonella* out of the 360 samples from drinking water, cloacal swabs, feed, and eggs. Additionally, *Salmonella* prevalence was recorded as 6%, 5.8 % and 3.5% in Iran, Belgium, and Paraguay respectively (Jafari et al. 2007; Namata et al. 2009; Leotta et al. 2010). In contrast, an incidence rate 14.7%, 10.9% and 12.4% was reported by others (Murugkar et al. 2005; Abd El-Tawab et al. 2015; El-Sharkawy et al. 2017) respectively. Moreover, prevalence rate of 29% have been reported from broiler chickens in Lithuania, 20% in Italy, and 11% in the Netherlands (Van Overbeke et al. 2006; Pieskus et al. 2008). Additionally, a considerably higher prevalence rate was described in other studies in China (52.2%) (Yang et al. 2011), South Africa (51%) (Zishiri et al. 2016) and India (46%) (Srinivasan et al. 2014). It has been reported that the prevalence of *Salmonella* affected by geographical location, age of chickens, seasons of the year, the hygienic status of the farms, or antibiotic using regime (Sikder et al. 2005).

Serological identification demonstrated that 33 *Salmonella* serovars were confirmed as *Salmonella* with a prevalence rate of 6.6% (33 out of 500) as shown in Table 2. In our study, the most predominant serotypes were *S. blegdam* (39.4%) followed by *S. typhimurium* (21.2%) which belongs to O antigen serogroups D1 and B respectively. While, *S. enteritidis* (D1 serogroup), *S. infantis* (C1 serogroup), *S. kentucky* (C3 serogroup), and *S. virchow* (C1 serogroup) had the lowest ratio (3%) (Fig. 1). Both *S. enteritidis* and *S. typhimurium* were the most frequent serotypes isolated in Egyptian poultry farms (Abd El-Ghany et al. 2012; Halawa et al. 2016; El-Sharkawy et al. 2017; Elkenany et al. 2019). Moreover, it was confirmed that *S. kentucky* (C3 serogroup) had been identified as one of the most prominent *Salmonella* serovars isolated from broilers causing diarrhea and high mortalities in Egypt (Mahmoud et al. 2018). Additionally, it was reported that both *Salmonella enteritidis* and *Salmonella kentucky* were

the dominant serovars with 22.6% each (Awad et al. 2020). However, the percent of *S. enteritidis*, *S. virchow* and *S. kentucky* was reported to be 2.4%, 1.4% and 0.8% respectively (Sedeik et al. 2019). Finally, continuous monitoring of *Salmonella* serovars in chickens is essential for better prevention and reduction of its zoonotic risk.

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Author's Contribution

Nagwa S. Rabie, AA Samy and Kh M Elbayoumi carried out the research design, supervision for all the steps and revised the manuscript. Nagwa S Rabie, Hanaa S Fedawy and Ali M Ghetas: wrote and revised the manuscript. Hanaa S Fedawy, Dalia M Sedeeq, Abdelbaki MM, MA Bosila, Ali M Ghetas, Hoda M Mekky: isolation and identification of the microbial agent. Eman R Hassan, Zeinab MS Amin Girh, AA Samy and Ashraf MA Barakat: Collection of samples from different poultry farms and Cairo and Giza Poultry Slaughterhouses.

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Uncorrected Proof