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Bacteriological Studies on Swollen Head Syndrome in Broiler Chickens

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Abstract

Swollen head syndrome (SHS) in poultry is a serious problem for the commercial broiler industry causing great economic losses. In the present study, the prevalence of SHS was studied in 500 broiler chickens and identification of the causative bacterial agents was conducted focusing. Samples were collected from the serous fluid from subcutaneous of the infra orbital sinuses. The prevalence rate of SHS in examined broiler chickens was 20%. The results of bacteriological examinationrevealed that 92% samples showed bacterial isolation, 72% of them had single bacterial isolates while 20% had mixed bacterial isolates. On the other hand,8% had negative bacterial isolation.A total of 112 bacterial isolates were recovered. E. coli was the most prevalent; recorded as 80 isolates (60 single and 20 mixed) followed by P. aeruginosarepresented as 24 isolates (11 single and 13 mixed with E. coli), Klebsiella spp. represented as 4 isolates (one single 1% and 3 mixed with E. coli) and finally, Salmonella spp. represented as 4 isolates, all were mixed with E. coli.Results of serogrouping of E. coli isolates revealed that 6 O-serogroups were obtained of which O₁₁₉ was the most prevalent (30%) followed by O₁₈ (23.8%), O₁ (16.3%), O₁₂₅ (15%), O₁₅₈ (8.6%), and finally O_{169} (6.3%). On the other hand, serotyping of Salmonella isolates revealed the existence of the only serotype S. Kentucky; identifies in 50% of isolates, meanwhile 50% of isolates were untyped.

Keywords: Broiler chickens, Swollen head syndrome, E. coli, P. aeruginosa, Klebsiella spp., Salmonella.

INTRODUCTION

Swollen-head syndrome (SHS) is acute respiratory disease affecting the upper respiratory tract of domestic poultry; includingbroilers and broiler breeders, characterized by swelling of headand facial edema;in periorbital and infraorbital sinuses, resulted from subcutaneous

accumulation of inflammatory exudates in the head in response to secondary bacterial infection; usually *E. coli*, following the initial upper respiratory viral infection(**Shawki**, *et al.*, **2017 and Abdelmoezet** *al.*, **2019**).SHS was first described in the South America (**Morley and Thomson**, **1984**)and it has becomean important avian problem all over the world in the last few years (**Abdelmoezet** *al.*, **2019**).SHS can affect chickens of all ages mainly 4-6 weeks old causing considerable economiclosses in the avian industry due tomorbidity rate reaching to 10% and mortality about 3-4% as well as 2-3% reduction in the egg production (**Osman**, *et al.*, **2015**).

Although the etiology of SHS is uncertain anditsclinical appearance is quite variable, it is a multi-factorial disease. SHS may be resulted from a mixed infection with *E. coli* and paramyxovirus, coronavirus, or pneumovirus in which the initial viral infection causes acute rhinitis that permits *E. coli* to invade the subcutaneous facial tissues (Osman, *et al.*, 2015). The severity of the disease depends on presence of some environmental factors such as accumulation of ammonia, dust, overcrowding and bad ventilation (Abdelmoezet al., 2019). In addition to *E. coli*, there are many pathogen shadfarrelation with SHS as *Pseudomonas* spp., *Klebsiella* spp. and *Enterococcus* spp. (Osman, *et al.*, 2015).

The present study aimed to investigate the prevalence of SHS in broiler chickensas well as identification of the causative bacterial pathogens.

MATERIAL AND METHODS

Chickens.

A total of 500 diseased broiler chickens of different ages (2-5 weeks) from different farms in Beni-Suef and El-Fayoum Governorates were subjected to the present study during the period from January 2018 up to May 2019. These chickens were subjected to clinical and postmortem examinations to detect SHS lesion.

Samples.

A total of 100 serous fluid samples collected were collected aseptically from subcutaneous of the infra orbital area in the affected head of slaughtered diseased and freshly dead chickens.

Bacteriological examination.

The collected samples wereaseptically inoculated into tryptone soya broth and incubated aerobically at 37°C for 24 hrs. Then a loopful of the broth culture was streaked onto tryptone soya agar and MacConkey's agar and incubated aerobically at 37°C for 24-48hr. The lactose fermenting (pink) colonies were further inoculated onto eosin methylene blue (EMB) agar medium and incubated at 37°C for 18-24 hrs while pale colonies were streaked onto xylose lysine desoxycholate(XLD) agar and incubated aerobically at 37°C for 24 hrs. Also, the green coloured colonies were further inoculated onto cetrimide agar medium and incubated at 37°C for 18-24 hrs.

Identification of bacterial isolates.

Morphological and biochemical identification

All the recovered isolates were identified morphologically and biochemically according to schemes described by Collee *et al.* (1996) and Quinn *et al.* (2011). The following tests were used; oxidase, catalase, indole, methyl red,VogesProskauer, citrate utilization, urease,H₂S production on TSI,nitrate reduction and sugar fermentationfor glucose, lactose, sucrose, mannose, arabinose, maltose and mannitol. Other non-biochemical tests including pigment production, motility test and haemolysis onto blood agar were applied.

Identification by using Microbact(GNB 12A) Gram-negative system.

The appropriateMicrobactkit (Microbact24E, Oxoid) was used for identification of the isolates of Family *Enterobacteriaceae*members and other Gram-negative bacteria. Microbactstrips should only be used to identify pure cultures. It was used according to the manufacturer's instruction.

Serological identification:

Serogrouping of *E. coli* **isolates.** *E. coli* isolates were serogrouped by slide agglutination test using standard polyvalent and monovalent *E. coli* antisera according to **Quinn** *et al.* (2011).

Serotyping of *Salmonella*. *Salmonella* isolates were serotyped by slide agglutination test using diagnostic polyvalent and monovalent O and H *Salmonella* antisera(**DENKA SEIKEN Co., Japan**) according to Kauffman-white scheme (**Quinn** *et al.*, **2011**).

RESULTS

Prevalence of SHS in the examined broiler chickens.

Out of 500 broiler chicken, 100ones (20%) showed SHS lesions in clinical and PM examination.

Bacteriological examination.

The results represented in **table** (1) revealed that out of 100 serous fluid samples collected from subcutaneous of the infra orbital area in the broiler chicken heads, 92 samples (92%) revealed bacterial isolation, of which 72 samples (72%) had single bacterial isolates while 20 samples (20%) had mixed bacterial isolates. On the other hand, 8 samples (8%) had negative bacterial isolation.

Table (1): Results of bacteriological examination of samples collected from broiler chickens with SHS.

No. of samples	Positive bacterial isolation					Negative bacterial		
	Single	isolates	Mixed isolates		Total		isolation	
	No.	%	No.	%	No.	%	No.	%
100	72	72	20	20	92	92	8	8

^{%:} was calculated according to the number (No.) of the samples.

Prevalences of different bacterial pathogens recovered from SHS lesions in broiler chickens.

Form all collected samples, a total of 112 bacterial isolates were recovered. *E. coli* wasthe most prevalentrepresented as 80 *E. coli* isolates (80%); of which 60 single (60%) and 20 (20%) mixed with other bacteria. Then, *Pseudomonas aeruginosa* represented as 24 isolates(24%); of which 11 single (11%) and 13 (13%) mixed with *E. coli*. Afterthat, *Klebsiella* spp. represented as 4 isolates; of which one single (1%) and 3 (3%) mixed with *E. coli*. And finally, *Salmonella* spp. represented as 4 isolates(4%); all were mixed with *E. coli* (**Table 2**).

Table (2): Prevalence of different bacterial isolates from SHS in broiler chickens.

	Bacterial isolate	No. of isolates	%
	E.coli	60	60
Single isolates	P. aeruginosa	11	11
	Klebsiella spp.	1	1
	Total single isolates	72	72

	E.coli + P. aeruginosa	13	13
Mixed isolates	E.coli + Salmonella spp.	4	4
	E.coli + Klebsiellaspp.	3	4
	Total mixed infection	20	20
Negative bacterial isolation		8	8
Over all total		100	100

^{%:} was calculated according to the overall total number (No.) of the samples (n=100).

Serogrouping of *E. coli* isolates.

Out of 80 *E. coli* isolates, 6 O-serogroups were obtained. The serogroup O_{119} was the most prevalent represented in 24 isolates (30%) followed by serogroups $O_{18}(n=19; 23.8\%)$, O_1 (n=13; 16.3%), $O_{125}(n=12; 15\%)$, $O_{158}(n=7; 8.6\%)$, and finally $O_{169}(n=5; 6.3\%)$ (**Table 3**).

Table (3): Serogroups of *E. coli* recovered from swollen head broiler chickens.

E. coliSerogroup	No.	0/0
- O ₁₁₉	24	30
- O ₁₈	19	23.8
- O ₁	13	16.3
- O ₁₂₅	12	15
- O ₁₅₈	7	8.6
- O ₁₆₉	5	6.3
Total No. of isolates	80	100

^{%:} was calculated according to the total number (No.) of isolates (n=80).

Serotyping of Salmonella isolates.

Out of 4 *Salmonella* enterica isolates, only oneserotypes was identified as *Salmonella* Kentucky which were represented in 2 isolates (50%) meanwhile the other 2 isolates (50%) were untyped with the available antisera.

DISCUSSION

Poultry industry has reached great levels of development and became one of the most dynamic areas of agriculture. Poultry is the main protein feed of animal origin for human consumption worldwide (**Milsavljevic**, **2014**) due to efficiency of production cost as well as its short life cycle. In Egypt, great efforts were directed toward the poultry industry to meet the high requirements of animal protein.

SHS is a disease of upper respiratory tract and considered as one of these problems in last few years. The disease affects broilers and broiler breeders which resulted in inflammatory exudate beneath the skin (Seifi et al., 2015). SHS has been described as a multifactorial disease where the initial lesion mainly caused by viruses while the clinical signs are a consequence of secondary bacterial infections and the severity of the disease depends upon environmental factors (Abdelmoezet al., 2019). In this study, the prevalence of SHS in diseased 500 broiler chickensand identification of the different causative bacterial agents wereinvestigated.

In the present study, the results revealed that 20% of diseases chickens showed SHS

lesions in clinical and PM examination. Such prevalence wasnearly similar to that recoded by **Georgiades***et al.*(2001) who found that from 50 commercial broiler flocks concerning respiratory disease, signs of SHS were shown in 8 birds (16%).

In the current study, the results of bacteriological examination revealed that 92% of serous fluid samples collected from subcutaneous of the infra orbital sinuses of the broiler chickens showed bacterial isolation, 72% of them had single bacterial isolates while 20% had mixed bacterial isolates. On the other hand, 8% had negative bacterial isolation. These results are coincided with those obtained in Egypt by Abdelmoezet al. (2019) who recorded bacterial isolation from all samples of chickens with SHS (100%) of them 65% had single bacterial isolates while 35% had mixed bacterial isolates. Also, Ameret al. (2019) found that all heads samples were positive for bacteriological examination.

Regarding the prevalences of different bacterial pathogens recovered from SHS lesions in broiler chickens, the results illustrated in **table(2)** showed that a total of 112 bacterial isolates were recovered. *E. coli* was the most prevalent; recorded as 80 isolates (60 single and 20 mixed) followed by *P. aeruginosa* represented as 24 isolates (11 single and 13 mixed with *E. coli*), *Klebsiella* spp. represented as 4 isolates (one single 1% and 3 mixed with *E. coli*) and finally, *Salmonella* spp. represented as 4 isolates, all were mixed with *E. coli*. Such results were nearly similar to those recorded by **Abdelmoezet** al. (2019) whofound that 28/40 samples were positive for *E. coli* isolation (16 single and 12 mixed) and 10/40 samples were infected had *P. aeruginosa* isolates (4 single, 6 mixed with *E. coli* or *P. mirabilis*). Also, **Osman**, et al. (2015) examined bacteriologically 250 samples of SHS from broiler chicken and isolated 70*E.coli* (28%), 7*K. pneumoniae* (2.8%) and 5 *P. aeruginosa* (2%). **Georgiades** et al. (2001) found that *E. coli* was the most prevalent isolate from infraorbital sinuses of the affected birds as 87.5% of samples. Moreover, **Hafez and Löehren** (1990) and **Al-Ankariet** al. (2004) isolated pure *E. coli* culture from cases of SHS.

Additionally, in the current study there was an unprecedented result where 4*Salmonella* spp. (4%) were recovered from SHS lesions and that was considered; as we believe, the first record for *Salmonella* spp. as a cause of SHS.

Results of serogrouping of E. coli isolates were illustrated in **table** (3).Six O-serogroups were obtained of which O_{119} was the most prevalent (30%) followed by O_{18} (23.8%), O_{1} (16.3%), O_{125} (15%), O_{158} (8.6%), and finally O_{169} (6.3%).The distribution of O antigens was nearly similar to that reported in previous studies (**Gomiset al., 2001;Schouleret al., 2012 and Hasan** *et al.*, **2020**) who recovered nearly the same serogroups; beside other serogroups. On the contrary they differed from those obtained by **Wang** *et al.* (**2010**) who recovered 8 different serogroups; O_{65} , O_{78} , O_{8} , O_{120} , O_{2} , O_{92} , O_{108} , and O_{26} , and **Tanaet** *al.* (**2013**) who recovered 8 serogroupsE. coli including O_{2} , O_{8} , O_{15} , O_{73} , O_{86} , O_{102} , O_{115} and O_{139} .

On the other hand, serotyping of *Salmonella* isolates revealed the existence of the only serotype S. Kentucky; identifiesin 50% of isolates, meanwhile 50% of isolates were untyped. These results run parallel to those recorded by (**Radwanet al., 2016 andandHasan et al., 2018**) as S. Kentucky was one of the most frequent broiler chicken isolated serovars. Also, **Helalet al., (2019)** isolated different *salmonella* serotypes and found that 12.5% of isolates were S. Kentucky.

CONCLUSION

Swollen-head syndrome is a serious problem for the commercial broiler industry causing great economic losses. The prevalence rate of SHS in examined broiler chickens was 20%. *E. coli* is the most prevalent bacterial agent causing SHS followed by *P. aeruginosa*, *Klebsiella* spp. and *Salmonella* spp.

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