

## PREVALENCE OF VIRULENCE GENES OF SOME FOODBORNE BACTERIA IN CHICKEN MEAT PRODUCTS

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### ABSTRACT

This Study was carried out on 200 random samples of chicken meat products represented by chicken luncheon, chicken burger, chicken sausage and chicken shawerma (50 of each). Samples were randomly collected from different supermarkets and retailers of different sanitation levels at Mansoura city, El Dakahlia Province, Egypt and bacteriologically analyzed to assess the prevalence of *Staph. aureus*, *E. coli* and *S. spp.* and their enterotoxigenic virulence genes using PCR in some chicken meat products intended for direct consumption. The obtained results revealed that the prevalence of *Staph. aureus* in examined chicken luncheon, chicken burger, chicken sausage and chicken shawerma were 6%, 2%, 2% and 2%., respectively. While *E. coli* were 2%, 4%, 0% and 2% in examined samples respectively and *S. spp.* was isolated by 2% from shawerma only. The isolated *S. typhimurium* harbor *invA* and *stn* genes. The isolated *E. coli* showed presence of shiga toxin genes (*stx1* and *stx2*). The examined coagulase positive *Staph. aureus* showed the presence of different enterotoxin genes *sea*, *seb*, *sec*, *sed* and *see*. Thus it is necessary to adopt a regime of good, safe and healthy production of such chicken meat products with cleaning and disinfection and hygienic packaging in order to ensure safe products for consumers.

**Key words:** Prevalence, Virulent Genes, Foodborne Bacteria, Chicken, Meat Products.

### INTRODUCTION

Poultry meat and its products are very popular food throughout the world, it considered as cheap, good delicious and nutritious, source of protein with good flavour and easily digestion. Ready to eat food can be described as the status of food being ready for immediate consumption at the point of sale, it may be raw or cooked, and can be consumed without further treatment Tsang (2002). The importance of food as a vehicle for transmission of several diseases has been documented, especially in developing countries where the hygienic standards are not strictly followed or enforced Harakeh *et al.* (2005). *Staph. aureus* produces a wide variety of toxins including staphylococcal enterotoxins (SEs; SEA to SEE, SEG to SEI, SER to SET) with demonstrated emetic activity, SEs are a major cause of food poisoning, which typically occurs after ingestion of different foods, particularly chicken meat products, contaminated with *Staph. aureus* by improper handling and subsequent storage at elevated temperatures

Tharwat and Elabbasy (2014). Salmonellosis was one of the most commonly zoonotic disease accounting for 133,258 confirmed humancases Osek and Wiczorek (2010). *Salmonella* often present in fresh tissues due to defects during slaughtering process of poultry and carcass manipulation Lee *et al.* (1998) as well as Cebedo *et al.* (2008) concluded that *S. spp.* are pathogenic bacteria that can contaminate food products during or after processing. Hamilton *et al.* (2009) mentioned that *E.coli* was isolated from 60% of examined poultry from butcher shops with mean counts of 0.70 log<sub>10</sub> cfu/g. and 16% from poultry sold in supermarket samples with mean counts of 0.51 log<sub>10</sub> cfu/g. Matossian and Kingcott (1979) detected food poisoning outbreak from donar kebab (a product similar to shawerma). *Staphylococcus spp.*, *E.coli* and *S. spp.* were isolated from raw chicken products and chicken shawerma Kaneko *et al.* (1999) and Pelczar *et al.*, 2006). Shiga toxin (Stx)-producing *E. coli* (STX-EC), also known as Verotoxin-producing *E. coli* which associated with infantile diarrhea, haemorrhagic colitis, thrombocyticpurpura, and hemolytic uremic syndrome in humans Griffin and Tauxe (1991). The aim of this study was to assess the presence of these bacteria in some chicken meat products and the risk of contamination on the consumer.

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## MATERIALS AND METHODS

### 1- Collection of samples:

Two hundred samples of chilled chicken meat products (50 samples each of chicken luncheon, chicken burger, chicken sausage and chicken shawerma) at  $\pm 4^{\circ}$  C were collected aseptically from different shops (small grocery and large supermarkets) from Mansoura city, Dakahlia province and transferred to the laboratory in an insulated ice-box without delay.

### 2- Bacteriological examination:

**2.1- Preparation of food homogenate:** according to technique recommended by ISO, 6887-2, (2003) 25 g. of each sample was removed by a sterile scissors and forceps and stomached using Seward stomacher 80 biomaster England with 225 ml sterile buffered peptone water (0.1%) to give a homogenate of 1/10 dilution from which ten fold serial dilutions were prepared and subjected to the following bacteriological examination.

**2.2- Total *E.coli* count:** according to technique recommended by FDA (2002a).

**2.3- *Staphylococcus aureus* count:** FDA (2002b). using Baird-Parker agar plates, incubated at  $35^{\circ}$  C for 48 hr. The suspected *Staphaureus* colonies were isolated, purified and confirmed by coagulase test and the total count was calculated.

**2.4- Isolation of *E. coli*** according to technique recommended by ISO, 16649/2, (2001)

**2.5- Isolation of *Salmonellae* ISO, 6579 (2002):** by enrichment in peptone water at ( $37^{\circ}$  C for 24hr) then selection enrichment in Tetrathionate ( $37^{\circ}$  C for 24hr) and rappaportvasiliades at  $41.5^{\circ}$  C for 18 hr., plating on XLD, MaCconkey's and Hektoneentreic agar at  $37^{\circ}$ C for 24 hr. The presumptive colonies were confirmed biochemically and serologically.

### 3- Detection of virulence genes in *Staphaureus*, *E. coli* and *Salmonella* using PCR:

Carried out in Reference Lab for Quality Control on Poultry Production, Animal Health Research Institute, Dokki-Egypt.

#### 3.1- DNA extraction:

DNA extraction from positive samples were(6 *Staph. aureus*), (4*E. coli*) and (1 *Salmonella*) performed using the QIAamp DNA Mini kit (Qiagen, Germany, GmbH) with modifications from the manufacturer's recommendations. Briefly, 200  $\mu$ l of the sample suspension was incubated with 10  $\mu$ l of proteinase K and 200  $\mu$ l of lysis buffer at  $56^{\circ}$ C for 10 min. After incubation, 200  $\mu$ l of 100% ethanol was added to the lysate. The sample was then washed and centrifuged following the manufacturer's recommendations. Nucleic acid was eluted with 100  $\mu$ l of elution buffer provided in the kit.

#### 3.2- Oligonucleotide Primers:

The used Primers used were supplied from Metabion (Germany) are listed in Table (I) and Table (II).

#### 3.3- PCR amplification:

For uniplex PCR, primers were utilized in a 25-  $\mu$ l reaction containing 12.5  $\mu$ l of Emerald Amp Max PCR Master Mix (Takara, Japan), 1  $\mu$ l of each primer of 20 pmol concentrations, 4.5  $\mu$ l of water, and 6  $\mu$ l of DNA template. For stx1, stx2 duplex PCR, primers were utilized in a 50-  $\mu$ l reaction containing 25  $\mu$ l of Emerald Amp Max PCR Master Mix (Takara, Japan), 1  $\mu$ l of each primer of 20 pmol concentration, 13  $\mu$ l of water, and 8  $\mu$ l of DNA template. The reaction was performed in an appliedbiosystem 2720.

#### 3.4- Analysis of the PCR Products:

The products of PCR were separated by electrophoresis on 1.5% agarose gel (Applichem, Germany, GmbH) in 1x TBE buffer at room temperature using gradients of 5V/cm. For gel analysis, 20  $\mu$ l of the uniplex PCR products and 30  $\mu$ l of the duplex PCR products were loaded in each gel slot. Generuler 100 bp ladder (Fermentas, Thermo Scientific, Germany) was used to determine the fragment sizes. The gel was photographed by a gel documentation system (Alpha Innotech, Biometra) and the data was analyzed through computer software.

**Table I:** primer sequence for *Staph. aureus* enterotoxins genes used in multiplex PCR (Mehrotra *et al.*, 2000).

Primer pairs	Nucleotide sequence(5'→3')	Amplicon size (bp)
<i>sea</i> Forward Reverse	5' GGTTATCAATGTGCGGGTGG 3' 5' CGGCACTTTTTTCTCTTCGG 3'	102 bp
<i>seb</i> Forward Reverse	5' GTATGGTGGTGTAACTGAGC 3' 5' CCAAATAGTGACGAGTTAGG 3'	164 bp
<i>sec</i> Forward Reverse	5' AGATGAAGTAGTTGATGTGTATGG 3' 5' CACACTTTTAGAATCAACCG 3'	451 bp
<i>sed</i> Forward Reverse	5' CCAATAATAGGAGAAAATAAAAAGG 3' 5' ATTGGTATTTTTTTTCGTTC 3'	278 bp
<i>see</i> Forward Reverse	5' AGGTTTTTTTCACAGGTCATCC 3' 5' CTTTTTTTTCTTCGGTCAATC 3'	209bp

**Table II:** Primers sequences, target genes, amplicon sizes and cycling conditions.

Target gene	Primers sequences	Amplified segment (bp)	Primary denaturation	Amplification (35 cycles)			Final extension	Reference
				Secondary denaturation	Annealing	Extension		
<i>stn</i>	TTG TGT CGC TAT CAC TGG CAA CC	617	94°C 5 min.	94°C 30 sec.	59°C 45 sec.	72°C 45 sec.	72°C 10 min.	<b>Murugkar <i>et al.</i>, 2003</b>
	ATT CGT AAC CCG CTC TCG TCC							
<i>invA</i>	GTGAAATTATCGC CACGTTTCGGGCAA	284	94°C 5 min.	94°C 30 sec.	55°C 30 sec.	72°C 30 sec.	72°C 7 min	<b>Oliveira <i>et al.</i>, 2003</b>
	TCATCGCACCGTC AAAGGAACC							
<i>Stx1</i>	ACACTGGATGATC TCAGTGG	614	94°C 5 min.	94°C 30 sec.	58°C 45 sec.	72°C 45 sec.	72°C 10 min.	<b>Dipineto <i>et al.</i>, 2006</b>
	CTGAATCCCCCTC CATTATG							
<i>Stx2</i>	CCATGACAACGGA CAGCAGTT	779	94°C 5 min.	94°C 30 sec.	58°C 45 sec.	72°C 45 sec.	72°C 10 min.	<b>Dipineto <i>et al.</i>, 2006</b>
	CCTGTCAACTGAG CAGCACTTG							

**Statistical analysis:**

The results are expressed as mean  $\pm$  standard Error (SE). Data were statistically analyzed using statistical analysis systems. (SAS version 9.1, SAS Institute, Inc., 2003).

**RESULTS**

The achieved results of *Staph. aureus* in Tables (1 & 2) for Chicken luncheon, Chicken burger, Chicken sausage and Chicken shawerma were 3.2 $\pm$ 1.6, 3.5 $\pm$ 1.5, 3.6 $\pm$ 1.4 and 3.7 $\pm$ 1.3 log<sub>10</sub>cfu/g. with incidence rate 6%, 2%, 2% and 2 %, respectively. The results of *E. coli* in Tables (1,2) for Chicken luncheon, Chicken burger, Chicken sausage and Chicken shawerma were 3.4 $\pm$ 1.8, 3.1 $\pm$ 1.3, 3.7 $\pm$ 2.1 and 3.8 $\pm$ 1.5 log<sub>10</sub> cfu/g. with incidence rate 2%, 4%, 0% and 2%, respectively, serologically the isolated *E. coli* indicates presence of the enterotoxigenic strains *E. coli* O127:H6 in

chicken luncheon and *E. coli* O125:H21 and *E. coli* O127:H6 in chicken burger. *Salmonella* were not detected in Chicken luncheon, Chicken burger and Chicken sausage and detected in 2% of the examined chicken shawerma was *S. typhimurium*.

By PCR the results showed the presence of enterotoxin producing genes (A, C, D and E) in *Staph. Aureus* the three isolates of *Staph. aureus* isolated from luncheon showed the presence of enterotoxin gene 1<sup>st</sup> (A, E) , 2<sup>nd</sup> (A, D) and 3<sup>rd</sup> (B, D). The isolate of *Staph. aureus* isolated from burger showed presence of enterotoxin genes (A and E), the sausage isolate showed the presence of enterotoxin gene (A) while shawerma isolate showed presence of enterotoxin genes (A) .The virulence genes of shiga toxin (*stx1* and *stx2*) were examined using PCR in the four *E. coli* isolates the results were positive for these genes.

**Table 1:** Statistical analytical results of *Staph. Aureus* and *E. coli* in the examined samples expressed as  $\log_{10}\text{cfu/g}$ . (n=50).

Microbial count $\log_{10}\text{cfu/gm} \pm \text{S.E.}$	Chicken luncheon	Chicken burger	Chicken sausage	Chicken shawerma
<i>STAPH.aureus</i>	3.2±1.6	3.5±1.5	3.6±1.4	3.7±1.3
<i>E. coli</i>	3.4±1.8	3.1±1.3	3.7±2.1	3.8±1.5

**Table 2:** The incidence, Serotyping and virulence gene of isolated, *Staph. Aureus*, *E. coli* and *S. spp.* from the examined samples (N= 50 of each).

samples	Chicken luncheon		Chicken burger		Chicken sausage		Chicken shawerma	
	NO %	Strains & virulence gene	NO %	Strains & virulence gene	NO %	Strains & virulence gene	NO %	Strains & virulence gene
<i>Staph. aureus</i>	Cp 3(6%)	1 <sup>st</sup> <i>sea, see</i> 2 <sup>nd</sup> <i>sea, sed</i> 3 <sup>rd</sup> <i>seb, sed</i>	Cp 1(2%)	<i>sea, sed</i>	Cp 1(2%)	<i>sea</i>	Cp 1(2%)	<i>sea</i>
<i>E. coli</i>	1 (2%)	ETEC O127:H6 <i>Stx2</i>	2 (4%)	1 <sup>st</sup> ETEC O125:H21 <i>Stx1 and stx2</i>	ND -	-	1 (2%)	ETEC O125:H21 <i>Stx1 and stx2</i>
<i>S. SPP.</i>	ND	-	ND	-	ND	-	1 (2%)	<i>S. typhimurium</i> <i>inv A, and stm,</i>

ND. = not determined

No. = number of positive samples, C p =coagulase positive

## DISCUSSION

*Staph. aureus*, *E. coli* and *Salmonella* are the major causes of food borne infection and intoxication and their presence in food constitute an important hygienic problem for food processors, food handlers and consumers Bergadol (1989). The enterotoxigenesis generally is not lethal and the elderly are more susceptible than the younger individuals, where the amount of *STAPH. aureus* enterotoxins required for intoxication about 94-184  $\mu\text{g}$  Erol and Iseri, (2004).

The achieved results of *Staph. aureus* in Tables (1 & 2) for Chicken luncheon, Chicken burger, Chicken sausage and Chicken shawerma were 3.2±1.6, 3.5±1.5, 3.6±1.4 and 3.7±1.3  $\log_{10}\text{cfu/g}$ . with incidence rate 6%, 2%, 2% and 2 % respectively. The results nearly similar Saleh *et al.* (2010) who mentioned that *Staph. Aureus* count were  $1.14 \times 10^3 \pm 3.32 \times 10^2$ ,  $2.17 \times 10^3 \pm 4.31 \times 10^2$  and  $2.2 \times 10^3 \pm 4.45 \times 10^2/\text{g}$ . with different incidence of 4%, 12% and 16% for luncheon, beef-burger and sausage respectively higher percentage were reported by Amal, (2004) 15% and 25% in *Staph. aureus* for luncheon and fresh sausage; Fatim, (2004) could isolate *Staph. Aureus* from luncheon in percentage of 16%; and Soutos *et al.* (2003) in percentage of 19.4% in luncheon; Mousa, (1993) reported that *S. aureus* count was  $2.3 \times 10^4 \text{cfu/g}$ . for luncheon;

Ahmed, (1992) 6.6% in sausage. EL-Mossalami *et al.* (2009) detected *Staph. aureus* in 92%, 80% and 88% with mean values of  $3.25 \pm 6 \times 10^3$ ,  $2.8 \pm 1.4 \times 10^2$  and  $4.1 \pm 2 \times 10^3 \text{cfu/g}$ . in sausage, beef burger and shawerma respectively; Armany *et al.* (2016) could isolate *S. aureus* in percentage of 24% and 20% in raw sausage and luncheon; Shawish and AL-Humam (2016) were 12%, 22% and 30% in beef luncheon, beef burger and beef sausage; AL-Ghamdi, (2012) *S. aureus* count in chicken luncheon and chicken burger were  $1.47 \times 10^6$  and  $1.2 \times 10^7 \text{cfu/g}$  respectively. Ibrahim, (2009) detected *Staph. aureus* in 22.85% and 31.85% in luncheon, and sausage and EL-Khatieb (1997) (29%) in sausage. the percentage of coagulase positive *Staph. aureus* strains isolated from Chicken luncheon, Chicken burger, Chicken sausage and Chicken shawerma were 6%, 2%, 2% and 2% respectively as in Table (2). Chomvarin *et al.* (2006); Oh, *et al.* (2007) and Chiang *et al.* (2008) concluded that the occurrence of enterotoxigenic *Staph. aureus* in ready to eat food products has been reported in various parts all over the world; Shalaby and Zaki, (2008) could isolate 4, 5 and 3 enterotoxigenic strains of *Staph. aureus* from beef burger, sausage and shawerma respectively and Motten *et al.* (2011) found Coagulase positive *Staph. aureus* in luncheon by 7%, 7% and 5% from the collected samples from three supermarkets. Eldaly *et al.* (2014) showed that the isolation percentages of *Staph. aureus* in the

examined samples of luncheon, burger, and sausage were 15%, 10%, and 20% respectively.

The results of *E. coli* in Tables (1,2) for Chicken luncheon, Chicken burger, Chicken sausage and Chicken shawerma were  $3.4 \pm 1.8$ ,  $3.1 \pm 1.3$ ,  $3.7 \pm 2.1$  and  $3.8 \pm 1.5$   $\log_{10}$  cfu/g. with incidence rate 2%, 4%, 0% and 2% respectively, serologically the isolated *E. coli* indicates presence of the enterotoxigenic strains *E. coli* O127:H6 in chicken luncheon and *E. coli* O125:H21 and *E. coli* O127:H6 in chicken burger. These results nearly similar to Samaha *et al.* (2012) were 8% in chicken luncheon; Ibrahim (2009) were 5.71% in luncheon; Fawzy (2004) were 8% in luncheon and Amal (2004) were 5 and 25% in luncheon and fresh sausage. Meanwhile, higher results were recorded by Armany *et al.* (2016) 20% and 24% in raw sausage and luncheon respectively and Mousa (1993) were 14% in luncheon; Ibrahim, (2009) were 42.85% in sausage and Fathi *et al.* (1992) in luncheon and sausage which were 41.67% and 20% which may be due to post processing contamination or inefficient cooking and improper handling.

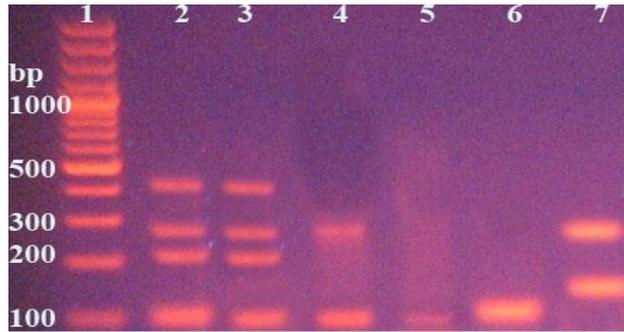
*Salmonella* were not detected in Chicken luncheon, Chicken burger and Chicken sausage and detected in 2% of the examined chicken shawerma was *S. typhimurium*. EL Jakee *et al.* (2014) detect *Salmonella* in burger, sausage and poultry products by 10, 35 and 25% respectively, the isolated *Salmonella* were *S. enteritidis* and *S. typhimurium* and Samaha *et al.* (2012) could isolate 8% *Salmonella* in chicken luncheon, Amal (2004); Ibrahim (2009) and Saleh *et al.* (2010) can not find *Salmonella* in luncheon while in sausage Mousa (1993); Saleh *et al.* (2010); Kozacinski *et al.* (2008); Ibrahim (2009); and Tudor *et al.* (2010) can't found *S. spp.* in fermented sausage. Amal (2004) found *salmonella* by 5% in sausage. The health hazard from *Salmonella* must not be underestimated. The fact that *Salmonella* was detected in samples from supermarkets, where chicken are displayed under refrigeration, shows that the spread of infection was not only confined to seemingly unhygienic environments FAO, (2013). It was suggested that to prevent contamination by *Salmonella* control measures must be taken at all stages of the food chain, from agricultural production, to processing, manufacturing and preparation of foods in both commercial establishments and at home WHO (2013).

PCR was applied to evaluate the presence of virulence genes in the isolated *Staph. aureus*, *E. coli* and *Salmonella*. *Staph. aureus* is one causes of food poisoning, its pathogenicity result from possession of

virulence genes that produce different toxins which result in self-limiting sever illness. For this reason the virulence genes of 6 isolated coagulase positive *Staph. aureus* were examined by PCR and the results showed the presence of enterotoxin producing genes (A, C, D and E) in *Staph. aureus* the three isolates of *Staph. aureus* isolated from luncheon showed the presence of enterotoxin gene 1<sup>st</sup>(A, E) , 2<sup>nd</sup> (A, D) and 3<sup>rd</sup> (B, D). The isolate of *Staph. aureus* isolated from burger showed presence of enterotoxin genes (A and E), the sausage isolate showed the presence of enterotoxin gene (A) while shawerma isolate showed presence of enterotoxin genes(A) as shown in Table (I) (Photo No. 1). The result agree with Eldaly *et al.* (2014) who found that luncheon samples harbored *seb* gene s while burger samples harbored *sed* gene also Tharwat and Elabbasy (2014) reported that SEA enterotoxin gene was the predominant enterotoxin genes which were detected in examined chicken burger and chicken luncheon.

*Staph. aureus* enterotoxin were analyzed from ready to eat products including pork ham, chicken cold cuts, pork sausage, salami and pork luncheon meat in a study conducted by Fijalkowski *et al.* (2016), this study reported that the most prevalent enterotoxin genes were *sei* (36%), *seln* (32%) and encoding exfoliative toxin A (37%). Another study conducted by Puah *et al.* (2016) revealed an incidence of (96.2%) virulence genes from *Staph. aureus* isolated from 200 food samples. A total of 30.8% of the isolates carried *SE* gene which cause food poisoning meanwhile the most common enterotoxin genes found were *seg* (11.5%) and *egc* (5.8%). On the other hands *Inv A* and *stn* virulence genes in the isolated *S. Typhimurium* were positive. *InvA* gene was amplified and detected at 284 bp while *stn* gene detected and amplified at 617 bp. In Korea, Li *et al.* (2006) could detect 17 virulence genes from isolated *Salmonella* using PCR assays, 14 genes assayed (82.4%) out of these 17 genes included *invA* gene.

The virulence genes of shiga toxin (*stx1* and *stx2*) were examined using PCR in the four *E. coli* isolates the results were positive for these genes Table (2) (Photo No. 2), these results were nearly similar to Balague *et al.* (2006) who collected 500 food samples from shops selling ready to eat foods in Argentina and *E. coli* virulence genes were examined by multiplex PCR (*stx1*, *stx2*, *eae A*, *cnf1*, *cnf2*, *ein v*, *Lt1*, *ST1* and *ST11*), ten *E. coli* isolates showed the presence of *stx1*, *stx2* genes while other genes were negative. Another study carried by Bohaychuck *et al.* (2006) reported shiga toxin producing *E. coli* O22: H8 from beef samples in Alberta and Canada.



**Photo No. (1):** Agarose gel electrophoresis of *Staph. aureus* PCR products using enterotoxins *Staphylococcus* primer.

**Lane "1":** 100 bp DNA ladder

**Lane "2 ":** positive amplification of 102 bp for enterotoxin A, 209 bp for enterotoxin E, 278 bp for enterotoxin D and 451 bp for enterotoxin C

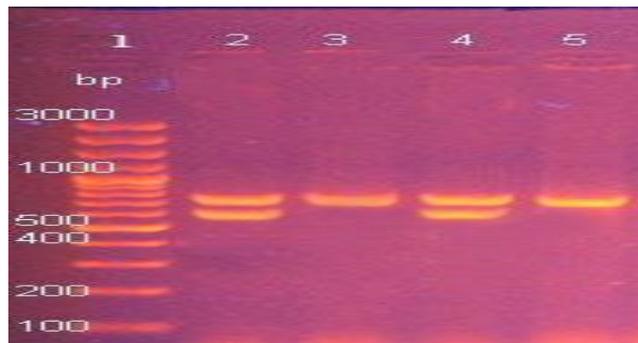
**Lane "3":** positive amplification of 102 bp for enterotoxin A, 209 bp for enterotoxin E, 278 bp for enterotoxin D and 451 bp for enterotoxin C

**Lane "4":** positive amplification of 102 bp for enterotoxin A and 278 bp for enterotoxin D

**Lane "5":** positive amplification of 102 bp for enterotoxin A

**Lane "6":** positive amplification of 102 bp for enterotoxin A

**Lane "7":** positive amplification of 164 bp for enterotoxin B and 278 bp for enterotoxin D



**Photo No. (2):** Agarose gel electrophoresis of *E. coli* PCR products using *stx1* and *stx2* primers.

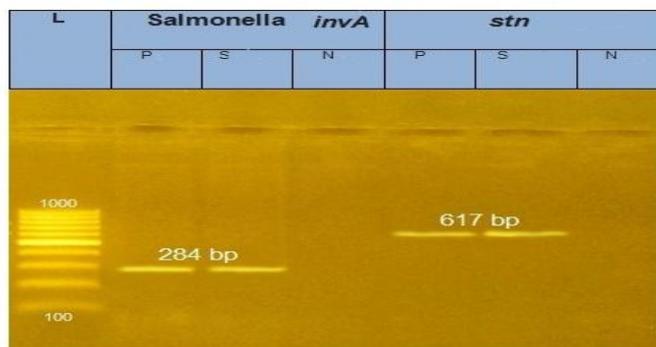
**Lane "1":** 100 bp DNA ladder

**Lane "2 ":** positive amplification of 614 bp for *stx1* gene and 779 bp for *stx2*.

**Lane "3":** positive amplification of 779 bp for *stx2*.

**Lane "4":** positive amplification of 614 bp for *stx1* gene and 779 bp for *stx2*.

**Lane "5":** positive amplification of 779 bp for *stx2*.



**Photo No. (3):** Agarose gel electrophoresis of *Salmonella* and PCR products using *invA*, and *stn*, primers

**L=** 100 bp DNA ladder.

**N=** negative control.

**P=** positive control (give amplification at 617pb for *stn* gene, 284 bp for *invA*, 614 bp

Sample of *S. Typhimurium* showed 284 bp amplification for *invA* gene and 617 pb for *stn* gene.

## CONCLUSION

This study confirms that chicken meat products may serve as a source of foodborne pathogens and accordingly a potential public health hazard. Corrective action needs to be employed to minimize the risk of consuming this type of fast food, such action must aim to minimizing the bacterial contamination during the production of chicken meat products (cleaning, cutting, seasoning and stacking), its cooking and serving. Regular surveillance by the public health regulatory bodies will ensure compliance with WHO and ISO standards for food safety.

Also, handling, storage and processing steps are major avenue for the cross contamination of the major materials used for the preparation of such product. Personal hygiene and processing practice of the food vendors are major factors.

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### مدى تواجد جينات الضراوة ببعض البكتيريا المنقولة بالغذاء في بعض منتجات لحوم الدواجن

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تم جمع ٢٠٠ عينة من لانشون وبيرجر وسجق وشاورما الدجاج بواقع ٥٠ عينة لكل منها. حيث تم عمل فحص بكتيري لكل من الاستاف اورييس والايشيرشيا كولاي وكذا معرفة مدى تواجد ميكروب السالمونيلا حيث كانت نسب العزل لميكروب الاستاف اورييس كالتالي ٦% و ٢% و ٢% و ٢% بينما نسب العزل للايشيرشيا كولاي كالتالي ٢% و ٤% و ٠% و ٢% على الترتيب وتم عزل عترة السالمونيلا تيفيموريم من شاورما الدجاج بنسبة ٢%. حيث تبين من هذه الدراسة ان عينات الانشون والسجق والهامبورجر كانت خالية من السالمونيلا عند الفحص البكتيريولوجي. تم عزل ٦ معزولات من ميكروب الاستاف اورييس (الموجبة لتجلط البلازما). تنتج المكورات العنقودية الذهبية مجموعة واسعة من السموم المعوية والتي تقوم بإثارة مراكز طالق في المخ وتشكل أحد الأسباب الرئيسية للتسمم الغذائي، والذي يحدث عادة بعد تناول الأطعمة المختلفة، لا سيما منتجات لحوم الدجاج الملوثة بالمكورات العنقودية الذهبية عن طريق سوء التعامل والتخزين في درجات حرارة مرتفعة بالإضافة لما تسببه كل من ميكروب السالمونيلا والايشيرشيا كولاي من حالات الإسهال الحاد لذلك تم تي تؤثر على قدرة الميكروب على احداث حالات مرضية عند تناول الاطعمة الملوثة بهذه الميكروبات. عند فحص السالمونيلا فحص جينات الضراوة لكل منهما. واجراء اختبار تفاعل البلمرة المتسلسل لتحديد وجود جينات الضراوة في الميكروبات المعزولة وال تيفيموريم المعزولة اظهرت وجود جيني invAstn. وبفحص معزولات ميكروب الايشيرشيا كولاي المعزولة تواجد بها جيني stx1 , stx2 كما أثبتت تواجد معظم جينات الضراوة لميكروب الستاف اورييس المعزول من العينات. وقد نقشت الأهمية الصحية للمعزولات وكذلك كيفية الإقلال من تواجدها باتباع نظم إدارة سلامة الغذاء أثناء التصنيع.