



Food Poisoning Bacteria Contaminating Beef at Abattoir Level

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ABSTRACT

It is very important to achieve the safety and sanitary quality of meat, so we should estimate the bacterial load in beef sample at abattoir level. The study was conducted on 90 beef samples (25 g of each) collected from 3 abattoirs in menofia governorate statistically, the results of aerobic plate count (APC), Enterobacteriaceae count and Staphylococci count, were not significantly different ($P>0.05$) between the three slaughterhouses A, B and C. The prevalence of the Enteropathogenic *E. coli* isolated from the beef samples in A, B and C were 16.66 %, 26.66%, and 20 %, respectively. Serologically, the serotypes of *E. coli* were O26:H11, O114:H4, O172, O128: H172, O 125: H 21, O163:H2, O91: H21, O78, O124, O111:H2, and O158. The prevalence of *Salmonella* in the three abattoirs were 6.66%, 23.33 %, and 13.33 %, respectively, and the identified serotypes were *Salmonella Infantis*, *Salmonella Typhimurium*, *Salmonella* Enteritidis, *Salmonella Tsevie*, *Salmonella Kentucky*, *Salmonella Heidelberg*, and *Salmonella Tamale*. Also, *Staphylococcus aureus* was isolated from 34.44% (31/90) beef samples, with the following percentages from the three abattoirs, 26.66%, 43.33%, and 33.33%, respectively. Conclusively, the results were higher than those recommended by Egyptian Organization of Standardization (EOS), therefore good slaughtering and hygienic practices recommended by general organization for veterinary services (GOVS) must be followed.

Keywords: beef, bacteriological profile, *E. coli*, *salmonellae*, *S. aureus*.

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1.INTRODUCTION:

The intact tissues of healthy slaughtered animals are mostly sterile, but the meat may be contaminated during slaughtering, handling, processing and storage from hands, workers, clothes, knives, hide, gut, fecal material on feet or from the environment.

Microbial contamination of the carcass results in spoilage of meat, reduced shelf-life of meat and public health hazards (Phillips *et al.*, 2006) either due to presence of spoilage bacteria responsible for unfavorable changes or pathogenic bacteria leading to harmful effects

as food infection or intoxication among consumers (Eley, 1992).

Fecal matter was a major source of contamination and could reach carcasses through direct deposition, as well as by indirect contact between contaminated and clean carcasses, equipment, workers, installations and air (Borch and Arinder, 2002).

The food borne pathogens are responsible to impose a substantial burden of infection in the developed countries, while the impact in case of developing countries is higher. It reduces markedly social and economic productivity of the countries. Amongst the food borne pathogens, *E. coli*, *Salmonella* and *Staphylococcus aureus* are the most common and frequent pathogens responsible for food poisoning and food related infections (Pires *et al.*, 2012).

The aerobic plate count indicates the level of microorganisms in a product and provide general estimate of the quality, shelf life and post heat processing contamination (Maturin and Peeler, 1998).

Enterobacteriaceae are group of bacteria that is used to assess the general hygienic status of a food product (HPA, 2004). So, the object of the current study was to evaluate the bacteriological quality of beef at abattoir level.

2. MATERIALS AND METHODS:

2.1. Samples:

A total of 90 fresh beef samples (25 g of each) were collected from 3 different abattoirs namely A, B and C located in Menofia governorate. The samples were transferred in an ice box directly within an hour to the laboratory with a minimum delay for bacteriological examination.

2.2. Determination of APC, Enterobacteriaceae and total Staphylococci counts.

The technique recommended by (ICMSF, 1996) was carried out.

2.3. Screening of Enteropathogenic *E. coli*:

The *E. coli* strains in the meat samples were isolated using the method of the Bacteriological Analytical Manual (APHA, 2001) One ml of homogenized meat sample inoculated into MacConkey broth for enrichment of the *E. coli* strains, and then incubated at 37° C for 18 to 24 h. The enriched cultures were streaked onto Eosin Methylene Blue (EMB) agar in two plates and incubated at 37 °C for 24 h. Presumptive *E. coli* colonies on EMB plates, which were round and had a metallic-green color with a dark or purple center, were further confirmed by inoculating the colonies EMB plates and incubated at 44 °C for 24 h. Then presumptive *E. coli* colonies picked up and kept in Semi-solid nutrient agar for biochemical and serological identification.

2.4. Screening of salmonella organism according to ISO 6579:

(2002) protocol (International Organization for Standardization (ISO) 2002), The homogenate specified for isolation of salmonella was incubated at 37°C for 22 h for pre-enrichment. After resuscitation, 0.1 mL was inoculated into 10 ml Rappaport-Vassiliadis medium and incubated at 42°C for 24 h. After enrichment, a loopful of each enriched sample was streaked onto xylose lysine desoxycholate agar and incubated at 37°C for 24 h. the purified suspected colonies were selected and streaked onto slope nutrient agar for biochemical and serological identification.

2.5. Screening of *S. aureus* according to (APHA, 2001):

Accurately, 0.1 ml from each of previously prepared serial dilutions was spread over duplicated plates of Baird Parker agar using a sterile glass spreader. The inoculated and control plates were incubated at 37°C for 48 hours. The developed colonies were enumerated, and the total Staphylococci count /g was calculated. Also, the colonies were picked up and purified on nutrient agar slopes for further identification. Films were prepared from a pure culture of the isolated microorganism stained with Gram's stain and then examined microscopically. Staphylococci appeared as Gram positive cocci resembling grape like clusters. The purified suspected colonies were picked up with a sterile loop for biochemical and serological identification.

3. RESULTS:

Table (1) indicated that the mean values of APC (cfu/cm²) of the tested beef samples in Shibin-Elkom, Ashmon and Menouf abattoirs were $1.33 \times 10^7 \pm 5.64 \times 10^6$, $1.85 \times 10^7 \pm 5.14 \times 10^6$ and $1.29 \times 10^7 \pm 3.85 \times 10^6$ respectively. On the other side, the mean values of total enterobacteriaceae count (cfu/cm²) of the tested beef samples abattoirs were $1.15 \times 10^4 \pm 2.93 \times 10^3$, $1.49 \times 10^4 \pm 3.74 \times 10^3$ and $1.80 \times 10^4 \pm 5.15 \times 10^3$ respectively. Furthermore, total Staphylococci count (cfu/cm²) of the tested beef samples were $1.75 \times 10^4 \pm 4.91 \times 10^3$, $2.19 \times 10^4 \pm 5.82 \times 10^3$ and $2.31 \times 10^4 \pm 5.74 \times 10^3$ respectively.

Table (2) showed that Enteropathogenic *E. coli* was serologically identified from the

examined samples of beef as A, B and C. abattoir were 16.66 % 26.66 % and 20 %, respectively. To clarify, In a abattoir EPEC were O₁₁₄ H₄ and O₁₇₂ at percentage of 3.33% and 3.33%, while EHEC was O₂₆H₁₁ at percentage of 6.66% finally ETEC was O₁₂₈H₁₇₂ at percentage of 3.33% (Table, 9). In B abattoir EPEC were O₁₆₃H₂ and O₇₈ at percentage of 3.33% and 3.33%, EHEC were O₂₆H₁₁ and O₉₁H₂₁ at percentage of 6.66% and 3.33%, ETEC were O₁₂₅H₂₁ and O₁₂₄ at percentage of 6.66% and 3.33% (Table, 9). Also, EPEC O₁₅₈ was isolated from C abattoir at percentage of 3.33%, EHEC were O₂₆H₁₁, O₉₁H₂₁ and O₁₁₁H₂ at percentage of and 3.33, 3.33% and 6.66%, ETEC was O₁₂₈H₁₇₂ at percentage of 3.33%.

Table (3) indicated that *Salmonella* organisms were isolated from 6.66%, 23.33 % and 13.33 % for A, B and C. respectively. *Salmonellae* could be identified serologically as *S. Infantis* 3.33% *S. Typhimurium* 3.33% for A abattoir, *S. Typhimurium* 3.33%. *S. Enteritidis* 10% *S. Tsevie* 3.33% *S. Kentucky* 3.33% *S. Heidelberg* 3.33% for B abattoir and *S. Typhimurium* 6.66% *S. Enteritidis* 3.33% and *S. Tamale* (3.33%) for C abattoir with percentage of 6.66% for A abattoir, 23.33% for B abattoir and 3.33% for C abattoir.

Table (4) showed that *S. aureus* organism was isolated from 26.66%, 43.33% and 33.33% from A, B and C abattoirs, respectively with total no. of 31 and percentage of 34.44% of the three abattoirs.

Table (1): average counts of Aerobic Plate (APC/g), Enterobacteriaceae and Staphylococci of the examined beef samples (n=30)

Bacterial group	Abattoirs		
	A	B	C
APC	$1.33 \times 10^7 \pm 5.64 \times 10^6$	$1.85 \times 10^7 \pm 5.14 \times 10^6$	$1.29 \times 10^7 \pm 3.85 \times 10^6$
Enterobacteriaceae	$1.15 \times 10^4 \pm 2.93 \times 10^3$	$1.49 \times 10^4 \pm 3.74 \times 10^3$	$1.80 \times 10^4 \pm 5.15 \times 10^3$
Staphylococci	$1.75 \times 10^4 \pm 4.91 \times 10^3$	$2.19 \times 10^4 \pm 5.82 \times 10^3$	$2.31 \times 10^4 \pm 5.74 \times 10^3$

Table (2): Incidence and serotyping of Enteropathogenic *E. coli* isolated from the examined beef samples (n=30).

<i>E. Coli</i> Strains	A		B		C		Serotyping
	abattoirs						
	No.	%	No.	%	No.	%	
O ₂₆ : H ₁₁	2	6.66	2	6.66	1	3.33	EHEC
O ₁₁₄ : H ₄	1	3.33	-	-	-	-	EPEC
O ₁₇₂	1	3.33	-	-	-	-	EPEC
O ₁₂₈ : H ₁₇₂	1	3.33	-	-	1	3.33	ETEC
O ₁₂₅ : H ₂₁	-	-	2	6.66	-	-	ETEC
O ₁₆₃ : H ₂	-	-	1	3.33	-	-	EPEC
O ₉₁ : H ₂₁	-	-	1	3.33	1	3.33	EHEC
O ₇₈	-	-	1	3.33	-	-	EPEC
O ₁₂₄	-	-	1	3.33	-	-	ETEC
O ₁₁₁ : H ₂	-	-	-	-	2	6.66	EHEC
O ₁₅₈	-	-	-	-	1	3.33	EPEC
Total	5	16.66	8	26.66	6	20	

Table (3): Incidence and serotyping of *salmonella* isolated from the examined beef samples (n=30).

abattoirs <i>Salmonella</i>	A		B		C		Group	Antigenic Srtucture	
	No.	%	No.	%	No.	%		O	H
<i>S. Infantis</i>	1	3.33	-	-	-	-	C ₁	6,7,14	r:1,5
<i>S. Typhimurium</i>	1	3.33	1	3.33	2	6.66	B	1,4,5,12	i:1, 2
<i>S. Enteritidis</i>	-	-	3	10	1	3.33	D ₁	1,9,12	g, m:
<i>S. Tsevie</i>	-	-	1	3.33	-	-	B	4,5	i: e, n, z ₁₅
<i>S. Kentucky</i>	-	-	1	3.33	-	-	C ₃	8,20	i: z ₆
<i>S. Heidelberg</i>	-	-	1	3.33	-	-	B	4,5,12	r:1,2
<i>S. Tamale</i>	-	-	-	-	1	3.33	C ₃	8,20	Z ₂₉ : e, n, z ₁₅
Total	2	6.66	7	23.3	4	13.3	-	-	-

Table (4): Incidence and serotyping of *Staphylococcus aureus* isolated from the examined beef samples (n=30).

Abattoir	No.	%
A	8	26.67
B	13	43.33
C	10	33.33
Total (90)	31	34.44

4. DISCUSSION:

A complete ignorance on the part of the meat handlers/ butchers in hygienic handling of carcasses during slaughter and retailing processes might be the main factors for producing meat with high microbial load. Hot and humid climate of this coastal area would have contributed in increasing the microbial load (Mukhopadhyay *et al.* 2009).

Lower aerobic Plate Count was recorded by El-Dally (1994) who studied that the mean value of total APC in shoulder muscle of beef was 4.7×10^3 Murray *et al.* (2001) who studied the mean value of total APC were $5.6 \times 10^2 \pm 0.44 \times 10^3$ cfu /g and in beef carcasses.

El-Taher (2009) who noticed that, the mean value of APC was 8.17×10^4 in raw meat. Salama (2013) who examined that mean value of APC/g of examined cattle meat samples from slaughterhouses of Elbagour, Menouf and Shibin-Elkoom were $1.92 \times 10^5 \pm 0.36 \times 10^5$, $8.58 \times 10^4 \pm 2.01 \times 10^4$ and $3.14 \times 10^4 \pm 0.69 \times 10^4$, respectively. Higher count was recorded by Paul and Sylvia (2014) who recorded that mean APC were 1.64×10^9 cfu/g for abattoir and 1.92×10^9 cfu/g for butcheries.

Enterobacteriaceae have an epidemiological importance as some of their members are pathogenic and may cause serious infections and food poisoning outbreaks to human being. Furthermore, the

total Enterobacteriaceae count can be taken as indicator of possible enteric contamination in the absence of coliform organisms (Mosupye and Van Holy, 2000).

Lower total Enterobacteriaceae count was recorded by El-Dally (1994) who found that the mean value of total Enterobacteriaceae count in shoulder muscle of beef was 5.67×10^2 cfu/cm². Murray *et al.* (2001) who revealed that total Enterobacteriaceae counts were $0.11 \times 10 \pm 0.20 \times 10$ cfu/g in beef carcasses. while higher count was recorded by Feizullah and Daskalov (2010) who revealed that the values of total Enterobacteriaceae count varied between 2×10 to 1.5×10^3 cfu/cm² for the smaller factory and 1.9×10 to 1.1×10^6 cfu/cm² for the high capacity slaughterhouse.

Staphylococcal food borne disease is one of the most common diseases worldwide resulting from the contamination of food by *staphylococcus aureus* enterotoxins. Although several SEs have been identified; SEA, a highly stable SE, is considered the most cause of SFD worldwide. Presence of pathogens in food products imposes potential hazard for consumers and cause grave economic loss and loss in human productivity via food borne disease. (Kadariya *et al.* 2014).

Lower count was recorded by by El-Taher (2009) who noticed that Staphylococci count were 4.16×10^3 for raw meat. Salama (2013) who revealed that mean value of Staphylococcal counts/g of examined meat samples from slaughterhouses of Elbagour, Menouf and Shibin-Elkom were $7.28 \times 10^3 \pm 1.69 \times 10^3$, $3.95 \times 10^3 \pm 0.84 \times 10^3$ /g and $1.07 \times 10^3 \pm 0.26 \times 10^3$, respectively.

Millions of people suffer from preventable food borne diseases every year. Food borne diseases are the growing public health problem and imply a great impact

worldwide. Pathogenic *E. coli* are responsible for three major types of clinical infections like enteric and diarrheal diseases, urinary tract infections, sepsis and meningitis. (Maheux *et al.* 2009).

Actually, *E. coli* was previously isolated from fresh meat by El-Dally (1994) who said that *E.coli* was found in one sample, El-Taher (2009) *E.coli* were isolated from 20% from raw meat , , Salama (2013) who reported that *E.coli* from cattle meat samples from slaughterhouses of Elbagour (25%), Menouf (20%) and Shibin Elkom (10%) and Enteropathogenic *E.coli* organisms were isolated from 30%, 20% and 15% of the examined samples of camel meat at Elbagour, Menouf, and Shibin Elkom abattoirs, respectively. , Abd El- Salam-azza (2014) revealed that *E.coli* species were recorded with lower percentages (12.8%).

The presence of potentially pathogenic Salmonella serotypes at the slaughtering stages is an evidence of the circulation of this pathogen in the food environment; its presence could increase consumers' risks of infection if proper food handling and preparation techniques are not followed, (Narváez-Bravo *et al.*, 2013).

Salmonella organisms were previously isolated from fresh meat by Majagaiya *et al.* (2008) who detected that a total of 250 raw meat samples for salmonella during September 2008 to December 2008 in Nepal. Salmonella was found to be positive in 23(9.2 %) of 250 examined samples, 4 samples of buffalo (8.00%) were found to be positive for salmonella. Different salmonella spp. was isolated from different types of meat samples belonging to Sero group D and E., Salama (2013) who isolated Salmonella from cattle meat samples from slaughterhouses of Elbagour (25%), Menouf (20%) and Shibin

Elkom (10%) , Abd El- Salam-Azza (2014) who said that Salmonella species were recorded with lower percentages (6%,) .

Food intoxication caused by *S. aureus* occurred by eating food containing ready formed Staphylococcus enterotoxin produced by enterotoxin producing *S. aureus*, today there are over twenty types of enterotoxins has been discovered and differentiated into Staph. enterotoxin A to Staph. enterotoxin IV (Hennekinne *et al.* 2012).

S. aureus organism were previously isolated from fresh meat by El-Taher (2009) who said that *S. aureus* were isolated from 26.6%, 33.3% and 53.3% of raw meat, raw chicken meat and kofta. respectively, Datta *et al.* (2012) who isolated 35 of *S. aureus* and all these isolates were tested for their sensitivity against common antibiotics used in Bangladesh.

5. CONCLUSION:

One can have concluded from the achieved results in this study that the examined beef samples were contaminated with different types of bacteria at different Monifyea abattoirs (Shibin-Elkom, Ashmon, Menouf) Furthermore, the contamination of such examined samples may be acquired during slaughtering, handling as well as preparation of their carcasses.

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