

Detection of Shiga-Like Toxin Producing *Escherichia coli* in Food of Animal Origin by Street Vendors at Luxor City

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Abstract

A total of 100 samples of ready-to-eat (RTE) meat sandwiches (50 cooked liver and, 50 chicken shawarma sandwiches) were randomly collected from street vendor carts at Luxor city. Samples were examined for shiga-like toxin producing *Escherichia coli* (STEC). The detected prevalence rates of *Escherichia coli* (*E. coli*) were (16/50) 32% and (2/50) 4% from cooked liver sandwiches and, chicken shawarma sandwiches, respectively. Serotyping of *E. coli* isolated from cooked liver sandwiches revealed the isolation of the following strains; O₁₇:H₁₈, O₂₆:H₁₁, O₈₆, O₉₁:H₂₁, O₁₁₁:H₂, O₁₁₃:H₄, O₁₁₉:H₆, O₁₂₁:H₇, O₁₂₈:H₂, O₁₄₆:H₂₁. On the other hand, serotyping of isolates from chicken shawarma sandwiches revealed the presence of only 2 *E. coli* serotypes (O₉₁:H₂₁ and O₁₂₁:H₇). Investigating the virulence genetic attributes for STEC by PCR for isolates from cooked liver sandwiches revealed that 12/16, 11/16 and 4/16 of *E. coli* isolates were positive for *stx1* gene, *stx2* gene, and *eaeA* gene, respectively. While, the two *E. coli* isolates from chicken shawarma sandwiches revealed that one of them was carried for both *stx1* and *stx2* genes and the other was carriers for *stx2* genes. On the other hand, only 4/16 of *E. coli* isolates (O₂₆:H₁₁ and O₁₁₁:H₂) were carried for *stx1*, *stx2* and *eaeA* genes. It could be concluded that the cooked liver sandwiches and shawarma chicken sandwiches prepared in street vendor carts were subjects of inadequate hygienic measures during preparation, and processing that may lead to contamination by shiga-toxin producing *E. coli* strains consequently may impose a potential public health risk.

Keywords: *Shiga, Like Toxin producing Escherichia coli, Food, Animal Origin, Street Vendors, Cooked liver, chicken, Shawarma.*

Introduction

In Egypt, Ready-to-eat foods are commonly sold in restaurants however a great part of these foods are vended by street vendors where foods are not effectively protected from dust and flies. Moreover, the safe food storage temperatures are difficult to maintain. Liver is considered one of the highly nutritive edible offal which contains vitamin A, vitamin B, vitamin C, vitamin D, Iron, Zinc and copper and mostly eaten as takeaway sandwiches (Salma *et al.*, 2015). Shawarma is a Middle Eastern Arabic style sandwich usually composed of shaved

lamb, goat, chicken, turkey, beef or a mixture of meats (Odu and Akano, 2012). Basically, it is a wrap of shredded meat (beef, lamb, or marinated chicken) prepared by alternately stacking strips of fat and pieces of seasoned meat on a rotating vertical skewer. The meat is roasted from the outside, while most of the inside remains raw.

There is an increase in the consumption of ready-to-eat fast food because of a change in social patterns characterized by increased mobility, large numbers of itinerant workers and less family centered activities.

Vendors were often of poor education level, unlicensed, untrained in food hygiene, technology and worked under crude unsanitary conditions as they often used stands and carts of crude and inefficient construction, running water was not easily accessible (**Abdalla *et al.*, 2008**). The hands are the most important vehicles for the transfer of organisms from feces, nose, skin, or other sites to food (**WHO, 1989**). Waste water was usually discarded in streets and garbage was discarded nearby providing harborage for insects and rodents (**Muinde and Kuria, 2005**).

Microbiological quality problems of ready-to-eat cooked liver and shawarma sandwiches depend greatly on low initial quality of raw liver and meat type and other ingredients, inefficient cooking process and improper sanitary practices for personnel and for cooking/processing utensils (**Kayaardi *et al.*, 2006**). Cross-contamination during preparation has been traced back to the use of uncooked vegetables and unhygienic handling (**Reij and Aantrekker, 2004**).

True food poisoning or food intoxication caused by eating food that contains a toxin due to bacterial growth in food while food infection is the second type of food-borne illness.

Enteric *E. coli* are part of the natural flora of many animals. Human infections occur through consumption of contaminated food products, drinking water contaminated with animal or human waste, or through direct person-to-person spread from poor hygiene (**Berger *et al.*, 2010**). *E. coli* considered as reflection of environmental contamination during slaughter processing and product handling (**Kanpelancher, 1981**). Enteric *E. coli* infections are traditionally divided into 6 pathotypes based on their pathogenicity profiles (virulence factors, clinical disease and phylogenetic profile): Enteropathogenic *E. coli* (EPEC), Enterohamorrhagic *E. coli* (EHEC), Enteroinvasive *E. coli* (EIEC, including *Shigella* spp), Enteroaggregative *E. coli* (EAEC), Enterotoxigenic *E. coli* (ETEC) and Diffusely Adherent *E. coli* (DAEC) (**Kaper *et al.*, 2004**).

Shiga toxin-producing *Escherichia coli* (STEC) or verotoxin-producing *Escherichia coli* remains a major cause of foodborne-related gas-

trointestinal diseases in humans (**Wani *et al.*, 2004**), particularly since these infections may result in life-threatening sequel such as hemorrhagic colitis (HC), hemolytic uremic syndrome (HUS), and thrombotic thrombocytopenia purpura (TTP). Enterotoxigenic *E. coli* (ETEC) is reported to be the most commonly isolated bacterial enteropathogenic in children under 5 year of age in developing countries, accounting for approximately 20% of cases, equivalent to several hundred million cases of diarrhea and several tens of thousands of deaths each year (**Qadri *et al.*, 2005**). ETEC is also the most common cause of travelers' diarrhea accounting for 10–60% of infections depending on the region visited (**Gascón *et al.*, 1998**).

The pathogenicity of these bacteria is mainly mediated by shiga toxins (*Stx1*, *Stx2* and their variants) encoded by *stx1* and *stx2* genes. Within human disease-associated strains, those producing Shiga toxin type 2 appear to be more commonly responsible for serious complications such as HUS than those producing only Shiga toxin type 1. In addition, a subset of STEC strains considered to be highly virulent for humans has the capacity to produce attaching and effacing lesions on intestinal mucosa, a property encoded on a pathogenicity island termed the locus for enterocyte effacement (LEE). LEE encodes a type III secretion system and *E. coli* secreted proteins, which deliver effector molecules to the host cell and disrupt the host cytoskeleton. LEE also carries *eaeA*, which encodes an outer membrane protein (intimin) required for intimate attachment to epithelial cells. The *eaeA* gene has been used as a convenient diagnostic marker for LEE-positive STECs (**Wani *et al.*, 2004**; **Best *et al.*, 2005**; **Paton and Paton, 2005**; **Dhanashree and Mallya, 2008**; **Feng, 2013** & **Mohammadi *et al.*, 2013**). The real-time multiplex PCR is robust and effective for the rapid and reliable detection of the seven predominant STEC serogroups of major public health (O₂₆, O₄₅, O₁₀₃, O₁₁₁, O₁₂₁, O₁₄₅, and O₁₅₇) and the detection of their virulence genes (**Anklam *et al.*, 2012** and **Díaz-Sánchez *et al.*, 2012**).

Therefore, the present study was conducted to evaluate the prevalence of Shiga like-toxin producing *E. coli* in cooked liver sandwiches

and Shawarma chicken sandwiches sold in Luxor city and their significance as public health hazards.

Materials and Methods

Samples Collection: A total of 100 ready-to-eat meat sandwiches (50 samples of cooked liver sandwiches and 50 samples shawarma sandwiches) were purchased from different street vendor carts in Luxor City. All collected samples were immediately transferred in ice-box containers, aseptically handled and moved promptly to Reference Laboratory for veterinary Quality control on Poultry Production, Luxor Branch / Animal Health Research Institute for bacteriological examination, isolation and identification of shiga-like toxin producing *E. coli*.

Preparation of samples: Twenty-five grams (± 0.5) of each sample were weighed and transferred into a sterile stomacher bag containing 225 ml 0.1% (W/V) sterile buffered peptone water were homogenized using a lab blender (Seward stomacher lab system 400 R \UK) for 2 minutes to obtain the original homogenate fluid of a dilution rate 10^{-1} . The preparation is carried out according to (ISO 6887-2: 2017)

Isolation of *E. coli*: was carried out according to the method described by Lee and Arp (1998). All isolates were detected in RLQP-Luxor branch/Animal Health Research Institute.

Identification of Enteropathogenic *E. coli*:

Suspected isolates of *E. coli* were identified according to MacFaddin (2000).

Serological identification of *E. coli* isolates:

The isolates were serologically identified according to Kok *et al.* (1996) by using rapid diagnostic *E. coli* antisera sets (DENKA SEIKEN Co., Japan) for diagnosis of the Enteropathogenic types.

Detection of Virulence genes of STEC by Polymerase Chain Reaction (PCR):

Primer sequences of *E. coli* used for PCR identification system:

Application of PCR for identification of shiga toxins (*stx1* & *stx2*) and intimin (*eaeA*) genes was performed by using primers (Pharmacia Biotech) as shown in the following table

Target gene	Oligonucleotide sequence (5' → 3')	Product size (bp)	References
<i>stx1</i> (F)	5' ACACTGGATGATCTCAGTGG 3'	614	Dhanashree and Mallya (2008)
<i>stx1</i> (R)	5' CTGAATCCCCCTCCATTATG 3'		
<i>stx2</i> (F)	5' CCATGACAACGGACAGCAGTT ' 3	779	
<i>stx2</i> (R)	5' CCTGTCAACTGAGCAGCAC-TTTG 3'		
<i>eaeA</i> (F)	5' GTGGCGAATACTGGCGAGACT ' 3	890	Mazaheri <i>et al.</i> , (2014)
<i>eaeA</i> (R)	5' CCCCATTCCTTTTTACCGTCG 3'		

DNA Extraction using QIA amp kit (Shah *et al.*, 2009): For genomic DNA extraction, single cloning of the bacteria was picked up from selective agar plate onto nutrient agar plates and incubated at 37 °C for 26 hours. After overnight culture on nutrient agar plates, one or two colonies were suspended in 20 ml of sterile distilled water, and the suspension was then heated at 100°C for 20 minutes. Accurately, 50-200 µl of the culture were placed in Eppendorf tube and frozen at -40 °C for further testing.

DNA amplification reaction of *E. coli* (Fagan *et al.*, 1999): The amplification was performed on a Thermal Cycler (Master cycler, Eppendorf, Hamburg, Germany). PCR assays were carried out in 1 ml of nucleic acid template prepared by using reference EHEC isolates (approximately 30 ng of DNA), 10 mM Tris-HCl (pH 8.4), 10 mM KCl, 3 mM MgCl₂; 2 mM concentrations of each primer, 0.2 mM concentrations of each 2'-deoxynucleoside 5'-

triphosphate, and 4 U of Ampli Taq DNA polymerase (Perkin-Elmer). The reference strains were *E. coli* O157:H7 Sakai (positive for *stx1*, *stx2* and *eaeA*) and *E. coli* K12DH5a (a non-pathogenic negative control strain) that does not possess any virulence gene. The amplified DNA fragments were analyzed by 2% of agarose gel electrophoresis (Applichem, Germany, GmbH) in 1x TBE buffer stained with ethidium bromide and captured as well as visualized on UV transilluminator. A 100 bp plus DNA Ladder (Qiagen, Germany, GmbH) was used to determine the fragment size.

Cycling conditions of the different primers during PCR in the following table:

Target gene	Primary denaturation	Secondary denaturation	Annealing	Extension	Final extension
<i>stx1</i>	95°C 3 min.	95°C 20 sec.	58°C 20 sec.	72°C 1.5 min.	72°C 5 min.
<i>stx2</i>	95°C 3 min.	95°C 20 sec.	58°C 20 sec.	72°C 1.5 min.	
<i>eaeA</i>	95°C 3 min.	95°C 20 sec.	58°C 20 sec.	72°C 1.5 min.	

Results and Discussion

E. coli virulence factors are mostly encoded on mobile elements that can move between organisms, and some organisms can have characteristics of more than one pathotype. (ISU/CFSPH, 2016). The presence of *E. coli* in food of animal origin is considered as indicator of mis-preparation, mis-handling, mis-storage or unhygienic service (Tebbut, 1999). Although,

E. coli is readily killed by temperature above 55°C, serious incidents occurred in such products, which reflects high level of abuse, cross contamination between raw and cooked foods (Varnam and Evans, 1991).

Table (1). Prevalence rate of *E. coli* isolated from the examined ready-to-eat meat sandwiches samples

Ready-to-eat meat sandwiches	No. of Samples	Positive Samples	
		No.	%
Cooked liver	50	16	32
Chicken Shawarma	50	2	4

Results in Table (1) revealed that the prevalence rate of *E. coli* was (32 % and, 4 %) in the examined cooked liver sandwich samples and, chicken shawarma sandwich samples, respectively. The current results of cooked liver samples were nearly similar to those reported by **Doaa (2015)** (30 %) and **El-zekaty et al., (2016)** (32 %); higher prevalence rates were recorded by **Dalia et al., (2013)** (42%) and **Shaltout et al., (2017)** (40%). Furthermore, lower prevalence rates were recorded by **Salma et al. (2015)** (10 %) and **El-shenawy et al., (2016)** (20 %). Prevalence rate of *E. coli* in the examined chicken shawarma samples were lower than that recorded by **Eman and Sherifa (2012)** (20 %) and **Hassanin et al., (2014)**

(33.3 %).

E. coli are serotyped based on the O (somatic lipopolysaccharide), H (flagellar) and K (capsular) antigens. A number of serotypes are known to contain EHEC. Some well-known organisms involved in human disease include *E. coli* O₁₅₇:H₇, *E. coli* O₁₅₇: H- (also known as *E. coli* O₁₅₇: NM, for "nonmotile"), and members of serogroups O₂₆, O₅₅, O₉₁, O₁₀₃, O₁₁₁, O₁₂₁ and O₁₄₅. Additional serogroups that have been reported in human clinical cases are O₄₅, O₈₀, O₁₀₄, O₁₁₃, O₁₁₇, O₁₁₈, O₁₂₈ and others (**ISU/CFSPH, 2016**).

Table (2). Serological identification of *E. coli* isolated from examined ready-to-eat meat sandwiches samples (N. = 18):

Serotypes	Prevalence rate of <i>E. coli</i> in examined sandwiches			
	Cooked liver		Chicken Shawarma	
	No.	(1) %	No.	(1) %
O ₁₇ :H ₁₈	1	5.6	0	0
O ₂₆ :H ₁₁	4	22.1	0	0
O ₈₆	1	5.6	0	0
O ₉₁ :H ₂₁	2	11	1	5.6
O ₁₁₁ :H ₂	1	5.6	0	0
O ₁₁₃ :H ₄	1	5.6	0	0
O ₁₁₉ :H ₆	1	5.6	0	0
O ₁₂₁ :H ₇	1	5.6	1	5.6
O ₁₂₈ :H ₂	3	16.5	0	0
O ₁₄₆ :H ₂₁	1	5.6	0	0
Total (10 serotypes)	16		2	

(1) No. of isolates /No. of examined strains

Results in Table (2) revealed that the identified *E. coli* serotypes were O₁₇:H₁₈ (5.6%), O₂₆:H₁₁ (22.1%), O₈₆ (5.6%), O₉₁:H₂₁ (11%), O₁₁₁:H₂ (5.6%), O₁₁₃:H₄ (5.6%), O₁₁₉:H₆ (5.6%), O₁₂₁:H₇ (5.6%), O₁₂₈:H₂ (16.5%), O₁₄₆:H₂₁ (5.6%) from the examined cooked liver sandwich samples; while the identified *E. coli* serotypes from the examined chicken shawarma sandwich samples were O₉₁:H₂₁ (5.6%), and O₁₂₁:H₇ (5.6%), respectively.

Similar results for cooked liver sandwiches were reported by many researchers as that reported by **El-zekaty *et al.*, (2016)** who identified O₂₆ (12%), O₈₆ (4%) and O₁₁₉ (8%).

Results were higher in studies by **Salma *et al.*, (2015)** who found that the incidence of *E. coli* were serotyped to O₂₆ (67%) in examined cooked liver sandwiches; and by **Doaa (2015)** who reported that the incidence of *E. coli* O₁₁₁ in cooked liver sandwiches was 10 % (3/30). On the other hand, **Morshdy *et al.* (2018)** reported that incidence of *E. coli* O₂₆, O₁₁₁, O₁₁₃, O₁₁₉ and O₁₂₇ in examined chicken shawarma sandwiches were 3(6%), 3(6%), 2(4%), 4(8%) and 3(6%), respectively.

Table (3). Occurrence of virulence genes of Shiga toxin-producing *E. coli* isolated from the examined ready to eat sandwiches:

Type of RTE samples	Serotypes	No. of isolates	Detection rate of Shiga-toxin virulence genes		
			* <i>stx1</i>	** <i>stx2</i>	*** <i>eaeA</i>
			No.	No.	No.
Cooked Liver (50)	O ₁₇ :H ₁₈	1	0	1	0
	O ₂₆ :H ₁₁	4	4	3	3
	O ₈₆	1	0	1	0
	O ₉₁ :H ₂₁	2	2	2	0
	O ₁₁₁ :H ₂	1	1	1	1
	O ₁₁₃ :H ₄	1	1	0	0
	O ₁₁₉ :H ₆	1	1	1	0
	O ₁₂₁ :H ₇	1	0	1	0
	O ₁₂₈ :H ₂	3	3	0	0
	O ₁₄₆ :H ₂₁	1	0	1	0
Total		16	12	11	4
Chicken Shawarma (50)	O ₉₁ :H ₂₁	1	1	1	0
	O ₁₂₁ :H ₇	1	0	1	0
Total		2	1	2	0

* *stx1*: Shiga- toxin 1 gene, ** *stx2*: Shiga- toxin 2 gene, *** *eaeA*: intimin gene

Results in Table (3) and Figure (1) revealed that most of the studied isolates produced shiga toxins (shiga-toxin 1 (*stx1*) and shiga-toxin 2 (*stx2*) and only two serotypes were produced intimin gene (*eaeA*). From cooked liver sandwich samples, out of 12/16 of *E. coli* serotypes (O₂₆:H₁₁, O₉₁:H₂₁, O₁₁₁:H₂, O₁₁₃:H₄, O₁₁₉:H₆, O₁₂₁:H₇ and O₁₂₈:H₂) carried *stx1*; out of 11/16 of *E.coli* serotypes (O₁₇:H₁₈, O₂₆:H₁₁, O₈₆, O₉₁:H₂₁, O₁₁₁:H₂, O₁₁₉:H₆, O₁₂₁:H₇, and O₁₄₆:H₂₁) carried *stx2*, while ; out of 4/16 of *E. coli* serotypes (O₂₆:H₁₁ and O₁₁₁:H₂) carried *eaeA* gene. While, from chicken shawarma sandwich samples, were isolated two *E. coli* sero-

types; O₉₁:H₂₁ which carried *stx1* and *stx2* genes and O₁₂₁:H₇ that had *stx2* gene only. On the other hand, only 4/16 of *E. coli* isolates (O₂₆:H₁₁ and O₁₁₁:H₂) were carriers *stx1*, *stx2* and *eaeA* genes.

Morshody *et al.* (2018) reported that shiga toxins genes were detected in chicken shawarma sandwiches samples. *Stx1* was detected in 3 *E. coli* isolates including O₂₆:H₁₁, O₁₁₉:H₆ and O₁₂₇:H₆ while only one isolate was positive for *Stx2* (O₁₁₉:H₆). In addition, four isolates including O₂₆:H₁₁, O₁₂₇:H₆, O₁₁₃:H₄, and O₁₁₁:H₂ were all positive for *eaeA* gene.

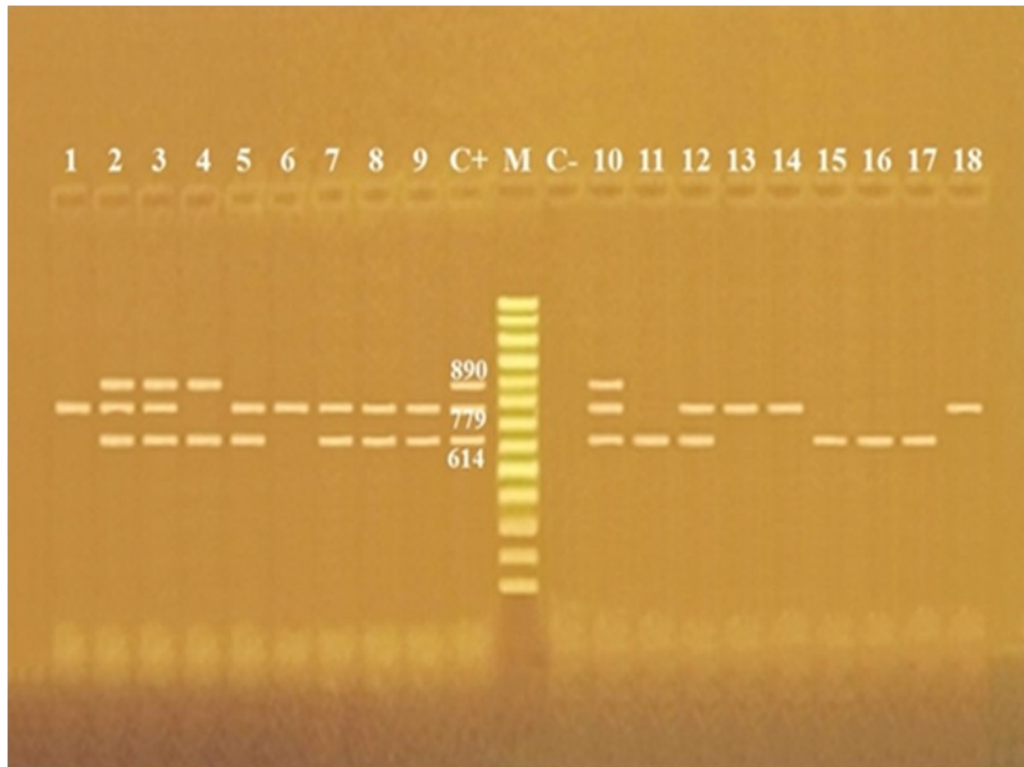


Figure (1): Agarose gel electrophoresis of multiplex PCR for *stx1* (614 bp), *stx2* (779 bp) and *eaeA* (890 bp) virulence genes for characterization of Enteropathogenic *E. coli* (n=18)

Lane M: 100 bp ladder. **Lane C+:** Control positive *E. coli* for *stx1*, *stx2* and *eaeA* genes. **Lane C-:** Control negative. **Lanes 1** (O17), **6** (O86), **13**, **14** (O121) & **18** (O146): Positive *E. coli* for *stx2* gene. **Lanes 2, 3** (O26) & **10** (O111): Positive *E. coli* for *stx1*, *stx2* and *eaeA* genes. **Lane 4** (O26): Positive *E. coli* for *stx1* and *eaeA* genes. **Lanes 5** (O26), **7**, **8**, **9** (O91) & **12** (O119): Positive *E. coli* for *stx1* and *stx2* genes. **Lanes 11** (O113) & **15**, **16**, **17** (O128): Positive *E. coli* for *stx1* gene.

Note: lanes 9 & 14 were from chicken shawarma sandwiches and other lanes were from cooked liver sandwiches.

The production of Shiga toxins (*stx*) is the unifying feature of all non-O157 STEC and O157 STEC. Various types of *stx* are produced, but they fall into two main types: Shiga toxin 1 (*stx1*) and Shiga toxin 2 (*stx2*) (Paton and Paton, 1998). Production of *stx* alone is usually insufficient for STEC strains to cause disease. Studies have shown that accessory virulence factors such as intimin, which are encoded by *eaeA* gene, are usually needed for STEC to be pathogenic (Paton *et al.*, 2004). The majority of *stx* genes are bacteriophage encoded, thus allowing a level of interstrain and possibly interspecies mobility. The presence of *stx* is thought to be the primary factor responsible for

intestinal manifestations (bloody diarrhoea) and systemic complications (HUS). The toxin uptake mechanisms in the intestine have yet to be fully elucidated for all STEC. (WHO, 1998) Shiga toxin-producing *E. coli* (STEC), also known as verotoxin-producing *E. coli* (VTEC), infections are public health concerns because of the severe illnesses they cause, such as hemorrhagic colitis and hemolytic uremic syndrome (HUS). STEC constitute a heterogeneous group of bacteria abundant in the reservoir and in the environment. (Smith *et al.*, 2014).

According to EFSA (2013) Serotypes responsible for haemorrhagic colitis (HC) and haemo-

lytic uraemic syndrome (HUS), O₁₅₇:H₇ and O₁₅₇:NM, were assigned to seropathotype A. Seropathotype B strains (O₂₆:H₁₁, O₁₀₃:H₂, O₁₁₁:NM, O₁₂₁:H₁₉ and O₁₄₅:NM) have a strong association with outbreaks and HUS but less commonly than those caused by seropathotype A. Seropathotype C serotypes (O₉₁:H₂₁, O₁₀₄:H₂₁, O₁₁₃:H₂₁, O₅:NM, O₁₂₁:NM and O₁₆₅:H₂₅) are associated with sporadic HUS cases but not with epidemics. **Rehamand Mohamed (2018)** found *E. coli* serovars O₁₁₁:H₂ that are carriers for *stx1*, *stx2* and *eaeA* genes, O₁₁₉:H₄ that are carriers of *stx1* and *stx2* genes and O₁₂₈:H₂ that are also carriers of *stx1* gene from stool samples of children with diarrheal symptoms in hospitals.

Food was recognized as a major vehicle of transmission of O157 STEC and was likely to play the same role for non-O157 STEC, but there is limited information on the occurrence of non-O157 STEC in the food supply. Meat and meat products, dairy products and any foods cross-contaminated by animal products during preparation may be contaminated (**WHO, 1998**). On the other hand, complete elimination of pathogens from raw materials (**Eisel *et al.*, 1997**) and food processing environment (**Tompkin, 2012**) is difficult, particularly when many food pathogens were known to be able to attach on food contact surfaces (**Jessen and Lammert, 2009**).

Conclusion

From the results it could be concluded that the hygienic conditions of some processed RTE food as cooked liver and chicken shawarma sandwiches offered by some street vendors in Luxor city demonstrated unsatisfactory microbiological quality with reference to the relevant Egyptian standards in a way that may impose a high risk for consumers.

Recommendations

Hygienic awareness should be applied for personnel who involved in handling of food at street vendor shops

Routine microbiological examination should be adopted in meat product, butcher's shops, groceries and other outlets that offer food of animal origin.

One health approach shall be apply for official regular monitor campaigns on food and food

products of animal origins and street vendors.

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