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# Bacteriological assessment of raw meats sold in Lagos, Nigeria

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Paper Information	A B S T R A C T		
	The high nutritive value of meat, having both essential macro- and micro-		
Received: 17 January, 2016	nutrients, makes it an important part of a balanced diet for most people.		
	However, meat is also a suitable medium for growth of microorganisms.		
Accepted: 28 May, 2016	This study evaluated the bacteriological quality of raw meat sold in Lagos,		
	Nigeria. Ten fold dilutions of twenty meat samples were plated using the		
Published: 20 August, 2016	spread plate technique. Total viable bacterial count (TVBC), total		
	Enterobacteriaceae count (TEC), total coliform count (TCC) and		
	Escherichia coli (ECC) were determined using plate count agar, violet red		
	bile glucose agar, Mac Conkey agar and ethylene methylene blue (EMB)		
	agar, respectively. Enterococci spp were counted on Slanetz Bartley		
	medium. Staphylococcus aureus and Micrococci were enumerated on		
	Mannitol salt agar (MSA) and Baird Parker agar, respectively. Salmonella-		
	Shigella agar was used for the isolation of Salmonella spp. Biochemical		
	tests were performed for further identification of isolates. TVBC ranged		
	from 1.44 x $10^4$ cfu/g to 4.38 x $10^4$ cfu/g; TEC ranged from 1.02 x $10^3$		
	cfu/g to 2.45 x 10 <sup>°</sup> cfu/g; TCC had a rage of $1.24 \times 10^{\circ}$ cfu/g to 2.76 x 10 <sup>°</sup>		
	cfu/g. TSAC ranged from $1.25 \times 10^2$ to $2.17 \times 10^2$ cfu/g. Salmonella counts		
	ranged from nil to $1.20 \times 10^2$ cfu/g. S. aureus had the highest percentage		
	occurrence while P. aeruginosa had lowest. This study revealed that raw		
	meats sold in some parts of Lagos are of poor bacteriological quality.		
	Hygienic practices of meat sellers need to be improved upon while organizing public enlightenment programmes is also crucial.		
-			
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Key words: Bacteria; food-borne; quality; safety; public health; food handling; sanitation.

**Abbreviations:** ECC = Escherichia coli count; EMB = Ethylene methylene blue; MSA = Mannitol salt agar; TEC = Total Enterobacteriaceae count; TCC = Total coliform count; TSAC = Total Staphylococcus aureus count; TVBC = Total viable bacterial count.

#### Introduction

Food security is a complex issue, where animal proteins such as meats, meat products, fish and fishery products are generally regarded as high risk commodity in respect of pathogen contents, natural toxins and other possible contaminants and adulterants (Yousuf et al., 2008). Food borne infections and illnesses is a major international health problem with consequent economic reduction. It is a major cause of illness and death worldwide (Adak et al., 2005). Foodborne infections lead to diarrheal diseases which usually have long-term effects on children's growth as well as on their physical and cognitive development, as well as the death of many children and (WHO, 2004).

Meat is considered the most nutritive source of protein consumed by humans. It is a perishable food. It contains sufficient nutrient needed to support the growth of microorganisms (Magnus, 1981). Water, protein and fat, phosphorus, iron and vitamins are also contained in meat and which are the chief constituents. Carcass is the major primary unit of meat and represents the ideal meat after head, hide, intestine and blood. Lean flesh, fat flesh and edible glands or organs such as heart, liver, kidney tongue and brain are the edible parts of a carcass. The preservation of meat as a perishable food usually is accomplished by a combination of preservation methods which greatly lengthen the keeping quality of the meat.

Meat is considered spoilt when it is unfit for human consumption. It is subjected to changes by its own enzyme, by microbial action and its fat may be oxidized chemically. Microorganisms grow on meat causing visual, textural and organoleptic change when they release metabolites (Jackson and Mcgowan, 2001).

Bacterial gastrointestinal infections continue to cause illness and death and contribute to economic loss in most parts of the world, including high-income countries that have developed surveillance and control programs (Ternhag et al., 2008). The possible sources of these bacteria are likely to come from the skin of the animal from which the meat was obtained. Other potential sources of microbial contaminations are the equipment used for each operation that is performed until the final product is eaten, the clothing and hands of personnel and the physical facilities themselves art all implicated (Rombouts and Nouts, 1994).

A number of foods have been reported to have high incidence of bacteria in Nigeria (Okonko et al. 2009a, b; Clarence et al., 2009; Bello and Osho, 2012; Bello et al., 2013; Bello et al., 2014). However, there is limited information on the microbiological risk associated with fresh meat retailed within such a highly populous community like Ikorodu. This study was carried out to determine the bacteriological quality of meat vended in Ikorodu, Lagos, Nigeria.

#### **Materials And Methods**

## Sample Collection

Twenty meat samples were purchased from stalls across Ikorodu markets. These were collected into separate sterile plastic bags and transported to the laboratory immediately after collection in ice-chest and tested upon arrival. Samples that could not be transported immediately were stored at 4<sup>o</sup>C for no longer than 4 hours (Elmali and Yaman, 2005).

## **Bacteriological Analysis**

## Preparation of samples for analysis

A ten-gram sample was weighed, introduced into a mixer with a sterile spatula under aseptic conditions, and then homogenized by adding 90 ml peptone (water 0.1%). One mil portion of each homogenate was used to prepare ten-fold dilutions up to  $10^{-6}$  with peptone water (Agaoglu et al., 2000).

## **Total Viable Bacterial Count (TVBC)**

Spread plate technique was used to inoculate agar plates. Aerobic mesophiles were determined using plate count agar (Oxoid CM 325); plates were incubated at  $30^{\circ}$ C for 24 to 48 hours.

## **Total Enterobacteriaceae Count (TEC)**

Enterobacteriaceae were isolated and enumerated on Violet Red Bile Glucose Agar. Plates were incubated at 37<sup>o</sup>C for 24 to 48 hours. Pink-red colour colonies with precipitation were taken into consideration.

## Total Coliform Count (TCC) and Escherichia coli

Coliform and Escherichia coli were enumerated on Mac Conkey agar and ethylene methylene blue (EMB) agar, respectively. Plates were incubated at  $37^{\circ}$ C for 24-48 hours. Pink red colonies with precipitation on Mac Conkey agar were enumerated as coliforms while colonies with greenish metallic sheen on EMB agar were counted as E. coli. Indole, methyl Red, Voges-Proskaner and Citrate (IMViC) tests were performed on colonies that showed shiny-metallic green to identify E. coli.

## Enterococci spp

Enterococci spp were counted on Slanetz Bartley medium after incubating aerobically at 37<sup>o</sup>C for 24-48 hours. The red colonies grown on this medium were taken into considerations.

## Total Staphylococcus aureus Count and Micrococci

Staphylococcus aureus and Micrococci were enumerated on Mannitol salt agar (MSA) and Baird Parker agar, respectively. Plates were incubated at 37°C for 24 to 48 hours. Yellow colonies on MSA were regarded as Staphylococcus aureus while small brown-black colonies without zones on Baird Parker agar were considered as Micrococcus spp. Catalase and coagulase tests were used for identification of Staphylococcus aureus (Addo et al., 2007).

#### **Bacillus cereus**

Bacillus cereus was isolated on Mannitol egg-yolk polymyxin (MYP) agar aerobically at 30<sup>o</sup>C for 24 to 48 hours. Typical colonies of Bacillus cereus were rough and dry with a bright pink background surrounded by an egg yolk precipitate. These were selected for further examinations (Gram stain, catalase test, motility test, nitrate reduction, VP reaction etc).

#### Salmonella spp

A twenty-five-gram sample was incubated in 225 ml buffered peptone water at 37<sup>o</sup>C for 24 hours. Subsequently, 0.1ml was inoculated into Rapport Vassiliadis broth and incubated at 43<sup>o</sup>C for 24 to 48 hours. Streak plates were prepared on Salmonella-Shigella agar and incubated at 24 to 48 hours. Pink-red colonies with black centers were

inoculated onto triple-sugar iron agar and lysin iron agar. Biochemical tests were performed for the identification of Salmonella spp.

#### **Results And Discussion**

Table 1 shows the bacteriological quality of raw meats sold in Ikorodu, Lagos, Nigeria. Twenty samples were analyzed. Each sample showed varying bacteriological qualities. The highest TVBC was encountered in sample 1 with count of 4.38 x  $10^4$  cfu/g while the lowest count was encountered in sample 18 with counts of 1.44 x  $10^4$  cfu/g. TEC ranged from 1.02 x  $10^3$  to 2.45 x  $10^3$  cfu/g as obtained in samples 11 and 1, respectively. Sample 1 had the highest TCC while the lowest occurred in sample 9 with counts of 2.76 x  $10^3$  cfu/g and 1.24 x  $10^3$  cfu/g, respectively. TSAC ranged from 1.25 x  $10^2$  to 2.17 x  $10^2$  cfu/g. Salmonella did not occur in some of the samples. Salmonella counts ranged from nil to 1.20 x  $10^2$  cfu/g as in sample 1.4.9, 11, 13, 18 and 19 were free of Salmonella.

Table 2 shows the occurrence of individual bacterial species in the different samples. Staphylococcus aureus was found to be present in all twenty meat samples while the occurrence of other bacterial species varied in the different samples. The percentage occurrence of bacterial species from raw meat samples sold in Ikorodu, Lagos, Nigeria is shown in Figure 1. S. aureus had the highest percentage occurrence of 19.23% while P. aeruginosa had the lowest (2.56%). Micrococcus spp was next to S. aureus with percentage occurrence of 16.67%; followed by Enterococcus spp (15.38%) and Klebsiela spp (11.54%). Salmonella spp and E. coli had same perentage occurrence of 14.1% and Proteus spp (6.41%).

Table 1: Bacteriological quality of raw meats sold in Ikorodu, Lagos, Nigeria.
Desterial scents in CEU/s

Sample	Bacterial counts in CFU/g						
	TVBC	TEC	TCC	TSAC	Salmonella		
					Count		
1	$4.38 \ge 10^4$	$2.45 \times 10^3$	$2.76 \times 10^3$	$2.17 \text{ x } 10^2$	-		
2	$4.10 \ge 10^4$	$2.40 \times 10^3$	$2.48 \times 10^3$	$2.00 \text{ x } 10^2$	$3.00 \ge 10^2$		
3	3.97 x 10 <sup>4</sup>	$1.44 \text{ x } 10^3$	2.45 x 10 <sup>3</sup>	1.49 x 10 <sup>2</sup>	$2.47 \ge 10^2$		
4	$4.20 \ge 10^4$	1.95 x 10 <sup>3</sup>	2.25 x 10 <sup>3</sup>	$1.27 \text{ x } 10^2$	-		
5	$3.50 \ge 10^4$	$1.20 \ge 10^3$	$2.14 \text{ x } 10^3$	$1.46 \ge 10^2$	$1.44 \ge 10^2$		
6	$2.97 \text{ x } 10^4$	$1.12 \text{ x } 10^3$	$1.95 \times 10^3$	1.96 x 10 <sup>2</sup>	$2.40 \ge 10^2$		
7	$3.00 \ge 10^4$	$1.50 \ge 10^3$	$1.90 \ge 10^3$	$1.44 \text{ x } 10^2$	$2.72 \times 10^2$		
8	$2.48 \times 10^4$	$1.45 \times 10^3$	$1.43 \times 10^3$	$1.44 \text{ x } 10^2$	$2.65 \ge 10^2$		
9	1.83 x 10 <sup>4</sup>	$1.15 \ge 10^3$	$1.24 \text{ x } 10^3$	$1.27 \text{ x } 10^2$	-		
10	$2.00 \ge 10^4$	$1.25 \times 10^3$	$1.97 \text{ x } 10^3$	$1.37 \ge 10^2$	$2.00 \ge 10^2$		
11	2.45 x 10 <sup>4</sup>	$1.02 \text{ x } 10^3$	$1.65 \ge 10^3$	1.39 x 10 <sup>2</sup>	-		
12	2.94 x 10 <sup>4</sup>	$1.11 \text{ x } 10^3$	$1.52 \times 10^3$	$1.40 \ge 10^2$	$1.20 \ge 10^2$		
13	$1.48 \ge 10^4$	$1.42 \text{ x } 10^3$	1.59 x 10 <sup>3</sup>	$1.26 \ge 10^2$	-		
14	$2.24 \text{ x } 10^4$	$1.48 \ge 10^3$	$1.60 \ge 10^3$	$1.25 \ge 10^2$	$1.85 \ge 10^2$		
15	$2.47 \times 10^4$	$1.54 \times 10^3$	$1.40 \ge 10^3$	$2.40 \times 10^2$	$1.22 \times 10^2$		
Sample		Bacterial counts i	n CFU/g				
	TVBC	TEC	TCC	TSAC	Salmonella		
					Count		
16	$3.00 \ge 10^4$	$1.04 \text{ x } 10^3$	$1.35 \times 10^3$	$2.22 \text{ x } 10^2$	$1.27 \ge 10^2$		
17	$2.35 \times 10^4$	$1.49 \times 10^3$	$1.29 \times 10^3$	1.96 x 10 <sup>2</sup>	$1.55 \ge 10^2$		
18	$1.44 \ge 10^4$	$1.64 \ge 10^3$	$1.54 \text{ x } 10^3$	$2.62 \text{ x } 10^2$	-		
19	$1.75 \times 10^4$	$1.60 \ge 10^3$	$2.23 \times 10^3$	$1.45 \times 10^2$	-		
20	$2.32 \times 10^4$	$1.95 \times 10^3$	$2.54 \times 10^3$	$2.57 \times 10^2$	$2.25 \times 10^2$		

 Table 2: Occurrence of bacterial species in different meat samples purchased in Ikorodu, Lagos, Nigeria

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1Staphylococcus aureus, Escherichia coli, Klebsiella spp, Enterococcus spp,2Staphylococcus aureus, Klebsiella spp, Salmonella spp, Micrococcus spp3Staphylococcus aureus, Micrococcus spp, Escherichia coli, Salmonella4Escherichia coli, Klebsiella spp, Enterococcus spp, Staphylococcus aureus Proteus spp5Staphylococcus aureus, E. coli, Klebsiella spp, Enterococcus spp, Micrococcus spp6Staphylococcus aureus, Micrococcus spp, Escherichia coli, Klebsiella spp, Salmonella7Psuedomonas aeroginosa, Staphylococcus aureus, Salmonella, Micrococcus spp ,8Staphylococcus aureus, Enterococcus spp, Salmonella, Proteus ssp9Enterococcus spp, Micrococcus spp, Staphylococcus aureus10Staphylococcus aureus, Klebsiella spp, Salmonella, Micrococcus spp ,11Staphylococcus aureus, Klebsiella spp, Salmonella, Micrococcus spp ,12Micrococcus aureus, Klebsiella spp, Salmonella, Micrococcus spp ,13Staphylococcus aureus, Enterococcus spp ,14Micrococcus aureus, E. coli, Klebsiella spp, Enterococcus spp ,13Staphylococcus aureus , E. coli, Klebsiella spp, Escherichia coli, Staphylococcus aureus13Staphylococcus aureus , E. coli, Klebsiella spp, Salmonella14Micrococcus spp, Staphylococcus aureus , Klebsiella spp, Salmonella15Staphylococcus aureus , Enterococcus spp, Escherichia coli16Escherichia coli, Enterococcus spp, Salmonella spp, Proteus spp17Staphylococcus aureus, Micrococcus spp, Escherichia coli, Salmonella18Psuedomonas aeroginosa, Staphylococcus aureus, Klebsiella spp, Micrococcus spp <th>_</th> <th>Bampie</th> <th></th>	_	Bampie	
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14Micrococcus spp, Staphylococcus aureus , Klebsiella spp, Salmonella15Staphylococcus aureus, Enterococcus spp, Escherichia coli16Escherichia coli, Enterococcus spp, Salmonella spp, Proteus spp17Staphylococcus aureus, Micrococcus spp, Escherichia coli, Salmonella18Psuedomonas aeroginosa, Staphylococcus aureus, Klebsiella spp, Micrococcus spp		13	Staphylococcus aureus, E. coli, Klebsiella spp, Enterococcus spp,
<ol> <li>Staphylococcus aureus, Enterococcus spp, Escherichia coli</li> <li>Escherichia coli, Enterococcus spp, Salmonella spp, Proteus spp</li> <li>Staphylococcus aureus, Micrococcus spp, Escherichia coli, Salmonella</li> <li>Psuedomonas aeroginosa, Staphylococcus aureus, Klebsiella spp, Micrococcus spp</li> </ol>		14	Micrococcus spp, Staphylococcus aureus, Klebsiella spp, Salmonella
<ol> <li>Escherichia coli, Enterococcus spp, Salmonella spp, Proteus spp</li> <li>Staphylococcus aureus, Micrococcus spp, Escherichia coli, Salmonella</li> <li>Psuedomonas aeroginosa, Staphylococcus aureus, Klebsiella spp, Micrococcus spp</li> </ol>		15	Staphylococcus aureus, "Enterococcus spp, Escherichia coli
<ol> <li>Staphylococcus aureus, Micrococcus spp, Escherichia coli, Salmonella</li> <li>Psuedomonas aeroginosa, Staphylococcus aureus, Klebsiella spp, Micrococcus spp</li> </ol>		16	Escherichia coli, Enterococcus spp, Salmonella spp, Proteus spp
18 Psuedomonas aeroginosa, Staphylococcus aureus, Klebsiella spp, Micrococcus spp		17	Staphylococcus aureus, Micrococcus spp, Escherichia coli, Salmonella
		18	Psuedomonas aeroginosa, Staphylococcus aureus, Klebsiella spp, Micrococcus spp
19 Staphylococcus aureus, Micrococcus spp, Escherichia coli, Enterococcus spp		19	Staphylococcus aureus, Micrococcus spp, Escherichia coli, Enterococcus spp
20 Micrococcus spp, Enterococcus spp, Proteus ssp, Salmonella spp, Staphylococcus aureus		20	Micrococcus spp, Enterococcus spp, Proteus ssp, Salmonella spp, Staphylococcus aureus



Parameter	S.aureus	Salmonella	Micrococcus	Enterococcus	E.coli	Proteus	Klebsiella	P.auroginosa
		Spp	spp	spp		spp	spp	, e
Gram's reaction	+	_	+	+	_	_	_	_
Catalase test	+	-	+	_	+	-	+	+
Coagulate test	+	-	-	+	_	-	_	_
Citrate Utilization	_	_	+	+	+	+	+	+
test								
Oxidate test	_	_	-	+	_		+	+
Urease test	+	_	-	N/A	_	+	+	N/A
Indole test	_	_	-	_	+	+	_	_
Glucose	+	+	+	+	+	+	+	N/A
Lactose	+	+	-	+	+	+	N/A	N/A
Sucrose	+	_	+	+	+	+	+	N/A
Mannitol	+	+	_	+	+	_	_	N/A
Maltose	+		+	+	+	+	_	N/A
Cellular	Cocci	Rod	Cocci	Cocci	Straight	Rod	Rod	Rod
morphology					rods			
Growth in Mannitol	Bright	N/A	N/A	N/A	N/A	NA	N/A	N/A
salt	yellow							
Growth in	N/A	N/A	N/A	Pink	Red/Pink	NA	Mucoid	Pale
MacConkey agar								
Growth in blood	Creamy	N/A	N/A	Creamy	Circular	NA	Large	Greenish
agar	white						white	

<b>T</b> 1 1 2 D 1 1 1 1		C1		C				<b>N T · · ·</b>
Table 3. Biochemical	charactaristics	of bootomol	1 10 Olator	trom row m	ant cold in	lzorodu	0000	NIGOTI
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Keys: + = Positive; - = Negative; N/A = Not applicable

A total of 82 isolates comprising of 8 different genera of bacteria were obtained in this study. The percentage occurrence of bacterial species from raw meat samples varied from one market to the other. This showed that all markets contributed equally to the microbial diversity reported in this study. The bacteria isolates were identified as S. aureus, P. aeruginosa, Micrococcus spp, Enterococcus spp, Klebsiela spp, Salmonella spp, E. coli and Proteus spp by comparing their morphological and biochemical characteristics with standard reference organisms.

The presence of these organisms in meat could be attributed to the fact that meat contains abundant nutrients required for the growth of bacteria. The high total viable counts recorded showed the microbial diversity on meats sold in these markets, and as a reflection of the environmental (Clarence et al., 2009). Some of the microorganisms isolated from fresh meat samples in this study have been earlier found in foods form other places (Enabulele and Uraih, 2009; Sobukola et al., 2009; Clarence et al., 2009; Oyeleke, 2009; Okonko et al., 2009). Nkanga and Uraih (1981) reported high prevalence rate of S. aureus in meat samples from traditional market in Benin City, Nigeria. E. coli and S. aureus are normal flora in human and animals, their presence in foods are indications of contaminations with faecal matters and excessive human handling (Clarence et al., 2009; Bello et al., 2013a).

The isolation of Enterococcus spp may be as a result of poor environmental conditions due to dust and contamination of the water used during slaughtering. This organism might have contaminated the meats during the process of handling by meat sellers. This is also in accordance to the assertion of Okonko et al. (2009a, b) that improper handling and improper hygiene might lead to the contamination of ready-to-eat foods and this might eventually affects the health of the consumers (Bello et al., 2013a, b; Bello et al., 2014).

However, the processors/handlers/sellers should observe strict hygienic measures so that they may not serve as source of chance inoculation of microorganisms and fecal contamination of fresh meats and other meat products. The presence of indicator and other organisms examined in this study is of special concern and perhaps the greatest danger associated with fresh meats used for food preparation, eating purposes and for other human consumption is contamination by human excrement. It demonstrates a potential health risk as the organism is pathogenic and causes complications in children.

#### Conclusion

It was revealed in this study that raw meat sold in Ikorodu, Lagos, Nigeria are of poor bacteriological quality. It showed that fresh meats are often contaminated with pathogenic bacteria and their presence meat foods should receive particular attention, because their presence indicate public health hazard and give warning signal for possible occurrence of food borne intoxication.

#### Recommendations

Meat handlers and sellers should be educated on the adverse effect of lack of proper personal and environmental hygiene and sanitation. Fresh meats to be used for consumption purposes should be adequately cooked before use and NAFDAC should ensure and enforce strict compliance of the recommended food standards as regards the production and sales of processed and packaged meat products. Veterinary doctors should inspect the animals to be slaughter before the meat is sold to the general public. Good manufacturing practices should be strictly adhered to by butchers and those selling the meat. The water used in washing the meat should be sterile, also the equipment must be washed properly before use. Further regulatory and educational efforts are needed to improve the safety of produce items. Continued progress on the part of regulators and industry to improve food safety are dependent on local, state, and federal agencies' ability to conduct epidemiologic and laboratory investigations that identify the offending agents and link them with specific foods; this should be adopted and executed in developing countries that are yet to adopt them.

#### References

- Adak GK, Meakins SM, Yip H, Lopman BA, O'Brien SJ. 2005.Disease risks from foods, England and Wales, 1996–2000. Emerging Infectious Diseases, [cited 2009 August 18]. Available from HYPERLINK "http://www.cdc.gov/ncidod/EID/vol11no03/04-0191.htm"
- Addo KK, Mensah GI, Bonsu C, Akyeh ML. 2007. Food and its preparation conditions in hotels in Accra Ghana: A concern for food safety: African. Journal of Food, Agricultural. and Nutritive. Development. 7(5): 1 12
- Agaoglu S, Yavuz MT, Berktas M, Guducuoghu H. 2000. Detection of scherichia coli 0157:h7 in retail ground beef, raw ground beef patties and raw meat balls sold in Van. Eastern Journal of Medicine 5 (2): 73-75.
- Bello OO, Bello TK, Fashola MO, Oluwadun A. 2014. Microbiological quality of some locally-produced fruit juices in Ogun State, Southwestern Nigeria. E3 Journal of Microbiology Research 2(1): 001-008.
- Bello OO, Bello TK, Bankole SA. 2013b. Occurrence of antibiotic-resistant Staphylococcus aureus in some street-vended foods in Ogun State, Nigeria. Journal of Advances in Biology 1(1): 21-28.
- Bello OO, Osho A. 2012. Antimicrobial Effects of Spices on Spoilage Organisms of Moin-Moin. Advances in Bioresearch. 3 (2): 60 65.
- Bello TK, Bello OO, Egberongbe HO, Azeez IA, Oluwadun A. 2013. Antibiotics resistance profile of Escherichia coli and Enterobacter aerogenes isolated from well waters in Ago-Iwoye, southwestern Nigeria. Journal of Advances in Biology 2(2): 135-144.

El-Gohany AH. 1994. Sausage and minced meat as a source of food poisoning microorganisms to man. Int-Veterinary-Medicine Journal, 30: 146-215.

Enabulele SA, Uraih N. 2009. Enterohaemorrhagic Escherichia coli 0157:H7 Prevalence in meat and vegetables sold in Benin City, Nigeria. African Journal of Microbiology Research, 3(5): 276-279.

- Jackson D, Mcgowan CH. 2001. Diet management effects on carcass attributes and meat quality of young goats. Small Ruminant Research, 28(1):93-98.
- Magnus P. 1981. Meat Composition. Food Science and Technology, 4th edition. Gohumunary Pub. London, pp. 108-215.
- Nkanga EJ, Uraih N. 1981. Prevalence of Staphylococcus aureus in meat samples from traditional market in Benin City Nigerian and possible control by use of condiments. Journal of Food Protection 44: 4 8.
- Okonko IO, Donbraye E, Babatunde SOI. 2009b. Microbiological Quality of Seafood processors and water used in two different sea processing plants in Nigeria EJEAFche, 2009b; 8(8): 621-629.
- Okonko IO, Ogun AA, Adejoye OD, Ogunjobi AA, Nkang AO, Adebayo-Tayo BC. 2009a. Hazards analysis critical control points (HACCP) and Microbiology qualities of Sea-foods as affected by Handler's Hygiene in Ibadan and Lagos, Nigeria. African Journal of Food Science, 3(1):035-050

- Oyeleke SB. Microbial assessment of some commercially prepared yoghurt retailed in Minna, Niger State. African Journal of Microbiology Research, 2009; 3(5): 245-248
- Rombouts FM, Nout R. 1994. Food Microbiology and Hygiene. Encyclopedia of Human Biology Academic Press, III: 661-665. Sobukola OP, Awonorin OS, Idowu AM, Bamiro OF. 2009. Microbial profile and critical control points during processing of 'robo' snack from melon seed (Citrullus lunatus thumb) in Abeokuta, Nigeria. African Journal of Biotechnology, 8(10):2385-2388
- Ternhag A, Törner A, Svensson Å, Ekdahl K, Giesecke J. 2008. Short- and long-term effects of bacterial gastrointestinal infections. Emerging Infectious Diseases, [cited 2009 August 18]. Available from http://www.cdc.gov/EID/content/14/1/143.htm
- WHO 2004. Food and health in Europe: a new basis for action, Text editing: Mary Stewart Burgher, WHO regional publications. European series, No. 96
- Yousuf AHM, Ahmed MK, Yeasmin S, Ahsan N, Rahman MM, Islam MM., 2008. Prevalence of Microbial Load in Shrimp, Penaeus monodon and Prawn, Macrobrachium rosenbergii from Bangladesh. World Journal of Agricultural Sciences, 4 (S): 852-855.
- Zheng J, Keys CE, Zhao S, Meng J, Brown EW. 2007. Enhanced subtyping scheme for Salmonella enteritidis. Emerging Infectious Diseases, [cited 2009 August 18]. Available from http://www.cdc.gov/EID/content/13/12/1932.htm