



Isolation, Identification and Antibiogram of Bacteria Isolated from Raw Cow Milk in Dutsin-Ma, Katsina State

Abdullahi K. ^{a*}, Innocent W. J. ^b, Hafsat S. B. ^b,
Habibu M. ^c and Yusuf Abdurrahman ^c

^a Environmental Unit, Department of Microbiology, Federal University Dutsin-Ma, Katsina State, Nigeria.

^b Medical Unit, Department of Microbiology, Federal University Dutsin-Ma, Katsina State, Nigeria.

^c Food and Industrial Unit, Department of Microbiology, Federal University Dutsin-Ma, Katsina State, Nigeria.

Authors' contributions

This work was carried out in collaboration among all authors. All authors read and approved the final manuscript.

Article Information

DOI: 10.9734/SAJRM/2024/v18i1340

Open Peer Review History:

This journal follows the Advanced Open Peer Review policy. Identity of the Reviewers, Editor(s) and additional Reviewers, peer review comments, different versions of the manuscript, comments of the editors, etc are available here: <https://www.sdiarticle5.com/review-history/111083>

Original Research Article

Received: 26/10/2023

Accepted: 01/01/2024

Published: 12/01/2024

ABSTRACT

The study was conducted to isolate and identify the bacteria, to know the sources of contamination of milk and antibiotic sensitivity of bacteria obtained from Dutsin-Ma, Katsina State. A total of 45 samples were collected from different locations in Dutsin-Ma such as Wednesday market, opposite the FUDMA takeoff site and Hospital road. All these samples were analyzed by culturing in different media such as *Salmonella-Shigella* agar, Eosin Methylene Blue agar, Mannitol Salt Agar, Nutrient agar, Cetrimide agar, and MacConkey agar. Biochemical tests were performed to identify the

*Corresponding author: Email: kabdullahi@fudutsinma.edu.ng;

organism. Among 45 samples, 20 (37.7%) were *Staphylococcus* spp. Similarly, 11 (20.8%), 6 (11.3%), 4(7.5%) and 12(22.6%) were found positive for *Escherichia coli*, *Pseudomonas aeruginosa*, *Klebsiella* and *Salmonella* spp. and. respectively. Results of the antibiotic sensitivity test represent that, out of ten antibiotics *Staphylococcus* sp. were very sensitive against Gentamicin (95%), Ciprofloxacin (90%), Streptomycin (70%), and highly resistant against Zinnacef (60%), Ampiclox (70%), Amoxicillin (50%). *Salmonella* sp. were highly sensitive to Pefloxacin (83.3%), Sparfloxacin (83.3%), and Ciprofloxacin (95%), but resistant against Augmentin (83.3%), Streptomycin (75%), Sulfamethoxazole (66.6%). *Klebsiella* spp. were highly sensitive to Pefloxacin (50%), Sparfloxacin, Chloramphenicol (75%), and Ciprofloxacin (75%), but resistant to Gentamicin (100%), Streptomycin (100%), Sulfamethoxazole (75%), Augmentin (75%). *Escherichia coli* were highly sensitive to Gentamicin (72.7%), Ciprofloxacin (100%), Ofloxacin (90.9%), Sparfloxacin (72.7%), but highly resistant to Sulphamethoxazole (72.2%), Whereas, *Pseudomonas aeruginosa* were highly sensitive Ciprofloxacin (100%), Chloramphenicol (66.6%), Ofloxacin (66.6%), but highly resistant against Streptomycin (100%), Augmentin 83.35%, Perfloxacin (83.3%). Data from this study suggested that raw milk contaminated with drug-resistant bacteria may cause public health hazards.

Keywords: Raw; milk; antibiotics; resistant.

1. INTRODUCTION

Raw milk is obtained from cows at homes in the Fulani hamlets and villages where the shelf-life and safety of the products are not considered Brunelle et al. [13]. Milk is a major component in the human diet of the vegetarian class, but it also serves as a very good medium for the growth of many Microorganisms including pathogenic bacteria [Ruegg, 2003]. Milk is a highly valuable Food, but raw milk contains and favours the growth of many microorganisms AJ, [9]. Foodborne illnesses are an important challenge to public health and cause significant economic problems in many countries [WHO. 2015]. The crucial goal of all food safety programs is to prevent food products contaminated by potential pathogens from reaching the consumer Kearns, [27,31]. Milk is an excellent medium for bacterial growth, which not only spoils the milk and associated products but also can cause infections in consumers [37-40]. Abdulkadir et al. [7]. Because of the specific production, it is not possible to fully avoid contamination of milk with microorganisms; therefore, the microbial contamination of milk is an important tool in determining its quality [1]. According to Adesokan et al. [8] During the normal milking operation however, milk is subjected to contamination from many different sources including (1) the udder and body of cows, (2) dust from the air, (3) litter and floor (4) flies, insects and rodents (5) water supply (6) hands and clothes of the milkier (7) utensils, bottles (8) atmosphere, etc. [2]. Thus, milk and dairy products prepared from milk could be an important source of food-borne pathogens [3].

Milk can be contaminated by various types of microorganisms such as *Streptococcus*, *Staphylococcus*, *Micrococcus*, *Escherichia*, *Bacillus*, *Salmonella* and *Pseudomonas* sp [4]. Huge numbers of microbes can get access to milk and various milk products including *Escherichia coli*, which is an indicator of milk contamination, constituting a public health hazard [5,30,34,35,36]. The diseases transmissible to humans through the consumption of spoiled milk like brucellosis, tuberculosis, salmonellosis, listeriosis, *Escherichia coli* infections and many others were described extensively in 1962 by [Kapla et al., 1962]. Antimicrobial development and eventual clinical adoption is one of the most significant issues in medical history, with engineered medicines having saved millions of lives against diseases that would have been lethal [14,19]. Nonetheless, due to the development of multidrug resistance (MDR) in these pathogens, treating infectious diseases is becoming increasingly difficult. Bhaskar. [10]. The present study was undertaken aiming to isolate and identify the bacteria and determine the antibiotic resistance of bacteria from raw cows in Dustin-Ma, Katsina State, Nigeria.

Bacterial contamination in raw milk has been a great threat to the economic and human health which causes mild disease to life-threatening illness. Bramley [11,12,24,25,26]. Raw milk can carry dangerous bacteria such as *Salmonella*, *E. coli*, *Listeria*, *Campylobacter*, and others that cause foodborne illness, often called "food poisoning." These bacteria can seriously injure the health of anyone who drinks raw milk or eats

products made from raw milk [6,7,23]. However, the bacteria in raw milk can be especially dangerous to people with weakened immune systems (such as transplant patients and individuals with HIV/AIDS, cancer, and diabetes), children, older adults, and pregnant women. In fact, the Centers for Disease Control finds that foodborne illness from raw milk especially affects children and teenagers [43 and 44]. Hence, there is need for this study.

Bacterial contamination of raw milk is of grave public health concern, especially when they are Antibiotic-resistant bacteria (ARB). Dewi, [15]. This is because, in addition to being a human pathogen, ARB are of health concern that can be disseminated from contaminated milk via mobile genetic elements like plasmids and transposons Falegan *et al.* [16]. Therefore, resulting in complicated, untreatable, and prolonged infections in humans, leading to higher healthcare costs and sometimes death [Ezekiel et al., 2019]. The aim of this research is to isolate, identify and determine antibiotic-resistant bacteria from raw cows in Dustin-Ma, Katsina State, Nigeria.

2. MATERIALS AND METHODS

2.1 Sample Collection

A total of 45 cow milk samples were collected randomly (using random sample collection) from 3 different locations in Dutsin-ma local government area of Katsina state, Nigeria. Nine samples were taken per week from each location for five (5) weeks. The samples were collected in sterile universal bottles and transported to the Microbiology Laboratory of the Department of Microbiology, Federal University, Dutsin-Ma for microbiological analysis. According to Gonzalez et al. [21].

2.2 Isolation of Bacteria

Serial dilutions of 10^{-1} , 10^{-2} , 10^{-3} , 10^{-4} and 10^{-5} were prepared. Aliquots of 1ml from each dilution were inoculated into sterile nutrient agar and MacConkey agar plates using the pour plate method and incubated at 37°C for 24 hours. Plate counts were recorded in cfu/ml. Colonial appearances such as size, shape, consistency, colour, elevation and differential characteristics such as pigmentation and the isolate was further sub-cultured in eosin methylene blue, *Salmonella Shigella* and mannitol salt agar and Gram

Staining were done to further identify the isolates [Cheesbrough,2003].

2.3 Gram Staining Technique

A smear of the suspected colony from overnight culture plates was made on a clean, grease-free slide. The smear was heat-fixed on a slide by passing the slide briefly over the Bunsen burner flame. The smear was then covered with a crystal violet stain for 1 minute. The stain was removed and rinsed with tap water. Afterwards, Lugol's iodine was added for 1 minute while decolonization was carried out by the addition of acetone for a few seconds. The slide was quickly washed with distilled water and counter-stained with Safranin for 1 minute. It was then finally flooded with water blot-dried and examined under the microscope using the oil-immersion objective lens. Suspected *Staphylococcus aureus* isolates were Gram-positive cocci (appearing purples) and arranged in clusters, *Klebsiella* spp., *Escherichia coli*, *Pseudomonas aerogenosa* and *Salmonella* spp. isolate was a Gram-negative rod (appearing pink) (Cheesbrough, 2003).

2.4 Biochemical Characterization and Identification

2.4.1 Catalase test

A drop of 3% hydrogen peroxide solution was placed on a clean, grease-free glass slide. A loopful of overnight colonies of the test organism was thereafter emulsified on the hydrogen peroxide. Observed bubble formation was regarded as positive and no bubble formation was regarded as negative (Cheesbrough, 2003).

2.4.2 Coagulase test

One loopful of the colony was emulsified on a clean grease-free glass slide. Ten microliter of citrated human plasma was added and observed for the presence of agglutination which indicates a positive reaction (Cheesbrough, 2003).

2.4.3 Urease test

The media was prepared based on the manufacturer's instructions then 40% of the urea solution was added and mixed well then poured in a tube and slanted then the isolate was inoculated and incubated for 18-24 hours. A positive test is demonstrated by an intense magenta to bright pink colour in 15-24hrs,

negative test shows no colour (Cheesbrough, 2003).

2.4.4 Triple sugar iron test

The media was prepared based on the manufacturer's instruction and poured into a test tube sterilized at 115°C for 30 minutes then it was allowed to set in the sloped form with a butt inoculation made by stabbing through the centre of the medium to the bottom of the tube with a straight inoculation needle then the surface of the agar slant was streaked with the isolated colony. A positive result was indicated by colour change, gas production and production of H₂S. Gehringer et al. [20].

2.4.5 Indole test

The media was prepared based on the manufacturer's instruction and the isolated colony was inoculated at 37°C for 24-28 hours then Kovac's reagent was added to the broth culture formation of pink to-red colour ("cherry-red ring) in the reagent layer on top of the medium within a second of adding the reagent indicates positive and no colour change indicated negative (Cheesbrough, 2003).

2.4.6 Methyl red test

The broth (MRVP) was prepared according to the manufacturer's instruction and the organism was inoculated and then incubated for 18-24hrs then methyl red reagent colour changed to red indicating positive and no colour change indicated negative (Cheesbrough, 2003).

2.4.7 Voges-Proskauer test

The broth (MRVP) was prepared according to the manufacturer's instruction and the organism was inoculated and then incubated for 18-24hrs then VPI and VPIL reagent colour change and ring formation indicated a positive lack of colour change indicated negative (Cheesbrough, 2003).

2.4.8 Citrate utilization test

The media Simmons agar was prepared based on the manufacturer's instruction and poured into a tube slanted then the organism was inoculated and incubated for 18-24hrs at 37°C positive result is indicated by color change from green to blue (Cheesbrough, 2003).

2.5 Antimicrobial Susceptibility Testing

Isolates were screened for phenotypic resistance and susceptibility to gram-positive and gram-negative discs. The procedure includes inoculation of stock cultures stored at 4°C on nutrient agar slants into 10ml of nutrient broth which was then incubated overnight at 37°C. Thereafter, serial dilution of 10¹ into sterile distilled water was carried out. Afterwards, 1 ml of the culture solution was transferred into sterile petri dishes. Thereafter, sterile Mueller Hinton agar that had been cooled to 55°C in a water bath was poured into each and allowed to solidify. Antibiotic sensitivity disc was later placed on one of the solidified plates sterilely and both plates were incubated at 37°C in an incubator overnight. Zone of inhibition seen around the antibiotic disc the following day were measured while the length was categorized as resistant, intermediate and sensitive after comparing with the Clinical Laboratory Standard Institute standard for each bacteria isolate (CLSI, 2021). Multidrug-resistant isolates were selected based on their resistance to ≥ 3 classes of antibiotics. Galton et al. [18]

2.6 Statistical Analysis

Microsoft Office Excel 2016 was used for the data analysis. The fungi isolated were recorded as frequency and prevalence. Two-Factor Without Replication Analysis of Variance (ANOVA) was used to compute and arrive at a statistical decision and $p < 0.05$.

3. RESULTS

Table 1. Demonstrated the mean bacterial counts of raw cow milk collected from the sample locations, in which Wednesday market of Dutsin-Ma is the highest with (8.5×10⁶ cfu/ml), while the least mean count was shown by Hospital Road with (1.04×10⁶ cfu/ml).

Table 2. shows the morphological characteristics of all the bacterial isolates of row cow milk using both the all-purpose and selective media used for the study.

Table 3. Demonstrated the microscopic and biochemical characteristics of five bacterial species associated with the row cow milk.

Table 4. Presents the distributions of all the bacterial species isolated from the study sites, which include *Staphylococcus aureus*,

Salmonella species, Escherichia coli, Pseudomonas aeruginosa, and Klebsiella species.

Table 5. shows the percentage of occurrence of bacterial species isolated from the three sample locations. Staphylococcus aureus showed the highest percentage of occurrence (37.2%) while Klebsiella spp. showed the lowest prevalence (7.5%).

Table 6 shows the antibiotic susceptibility profiles of gram-positive bacteria on different antibiotics.

Table 7 shows the antibiotic susceptibility profiles of gram-negative bacteria isolated from raw cow milk on different antibiotics

Fig. 1 shows the distribution of Isolates of Raw Cow Milk from Different Locations.

Fig. 2 shows the percentage of Occurrence of Bacteria Species.

Fig. 3 shows the percentage of occurrence of bacteria species.

Fig. 4 presents antibiotics resistance profile of *Staphylococcus aureus*.

Fig. 5 describes antibiotics Resistance profile of *Klebsiella spp.*

Fig. 6 shows the antibiotics resistance profile of *Escherichia coli*.

Fig. 7 shows the antibiotics resistance profile of *Pseudomonas aeruginosa*.

Table 1. Mean value of bacteria load

S/No	Locations	Mean bacterial count	Percentage %
1.	Wednesday market	8.5 X 10 ⁶	48.5
2.	Front of school	7.99 X 10 ⁶	45.6
3.	Hospital road	1.04 X 10 ⁶	5.9
	Total	1.753 X 10 ⁷	100

Table 2. Morphological characteristics of bacteria isolated from cow milk in dustin-ma

Media	Morphology	Bacteria
SSA	Pink with black dot	<i>Salmonella</i>
EMB	Green with a metallic sheen	<i>Escherichia coli</i>
	Pink no sheen mucoid	<i>Klebsiella spp</i>
MSA	Round Transparent	<i>Staphylococcus aureus</i>
Cetrimide	Greenish-blue	<i>Pseudomonas aeruginosa</i>

Key; SSA – Salmonella Shigella Agar, EMB – Eosin methylene blue, MSA – Mannitol salt agar

Table 3. Gram staining and biochemical test result

Bacteria	Gram Stain	Catalase	Oxidase	Urease	Citrate	Tsi	Indole	Mr	Vp	Coagulase
<i>Escherichia Coli</i>	-				-	+	+	+	-	
<i>S. Aureus</i>	+	+		+			-	+	+	+
<i>Salmonella spp.</i>	-			-	-	+	-	+	-	
<i>Klebsiella spp.</i>	-	+			+		-	-	+	+
<i>P. aeruginosa</i>	-	+	+		+		-	-	-	

- = negative + =positive, VP = Voge's Proskauer, MR = Methyl Red, TSI = Triple sugar ion test

Table 4. Distribution of isolates of raw cow milk from different locations

S/N	Isolate	Locations			Total
		Wednesday Market	Front of School	Hospital Road	
1.	<i>Staphylococcus aureus</i>	7	8	5	20
2.	<i>Escherichia coli</i>	4	3	4	11
3.	<i>Pseudomonas aeruginosa</i>	2	1	3	6
4.	<i>Klebsiella Spp.</i>	2	-	2	4
5.	<i>Salmonella Spp.</i>	5	4	3	12
	Total	20	16	17	53

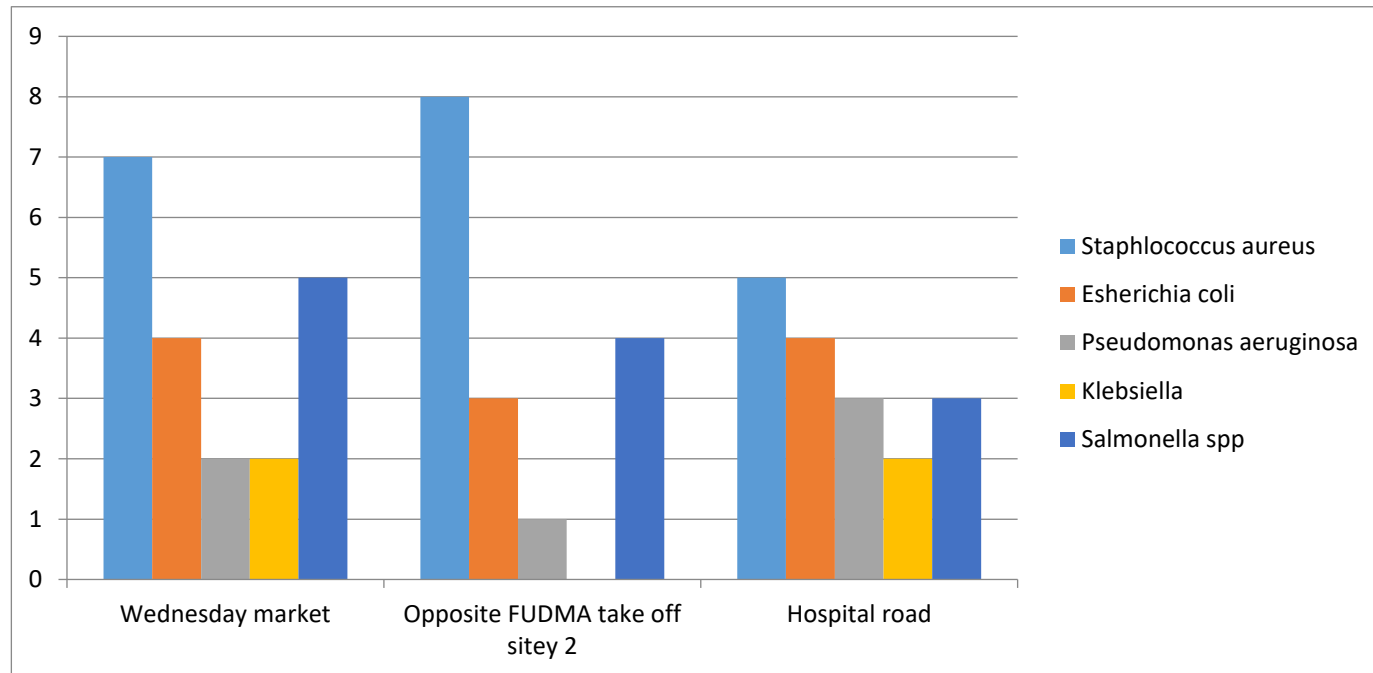


Fig. 1. Distribution of isolates of raw cow milk from different locations

Table 5. Percentage of occurrence of bacteria species

S/N	Isolate	Wednesday Market	Hospital Road	Front of Fudma Take off	Percentage (%)
1.	<i>Staphylococcus Aureus</i>	7	5	8	37.7
2.	<i>Escherichia Coli</i>	4	4	3	20.8
3.	<i>Pseudomonas</i>	2	3	1	11.3
4.	<i>Klebsiella Spp.</i>	2	2	-	7.5
5.	<i>Salmonella Spp.</i>	5	3	4	22.6
	Total	20	17	16	100

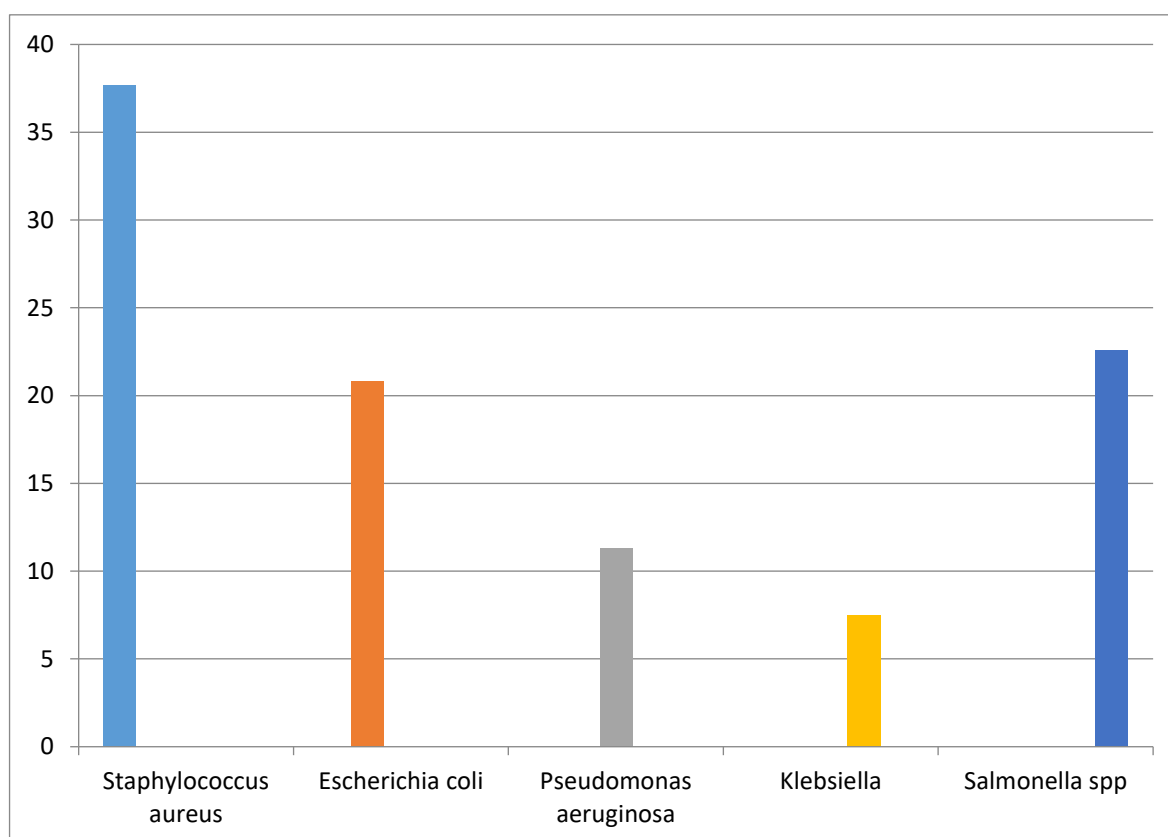


Fig. 2. Percentage of occurrence of bacteria species

Table 6. Antibiotics susceptibility pattern of gram positive bacteria

Antibiotics profile (%)	Isolates <i>Staphylococcus aureus</i> (n=20)		
	Susceptible	Intermediate	Resistance
Zinnacef	4(20)	4(20)	12(60)
Rocephin	10(20)	4(20)	5(25)
Streptomycin	14(70)	0(00)	4(20)
Erythromycin	12(60)	0(00)	8(40)
Gentamicin	19(95)	0(00)	1(5)
Ampiclox	4(20)	2(10)	14(70)
Sulfamethoxazole	12(60)	0(00)	8(40)
Amoxicillin	8(40)	2(10)	10(50)
Pefloxacin	10(50)	6(30)	4(20)
Ciprofloxacin	18(90)	2(10)	0(00)

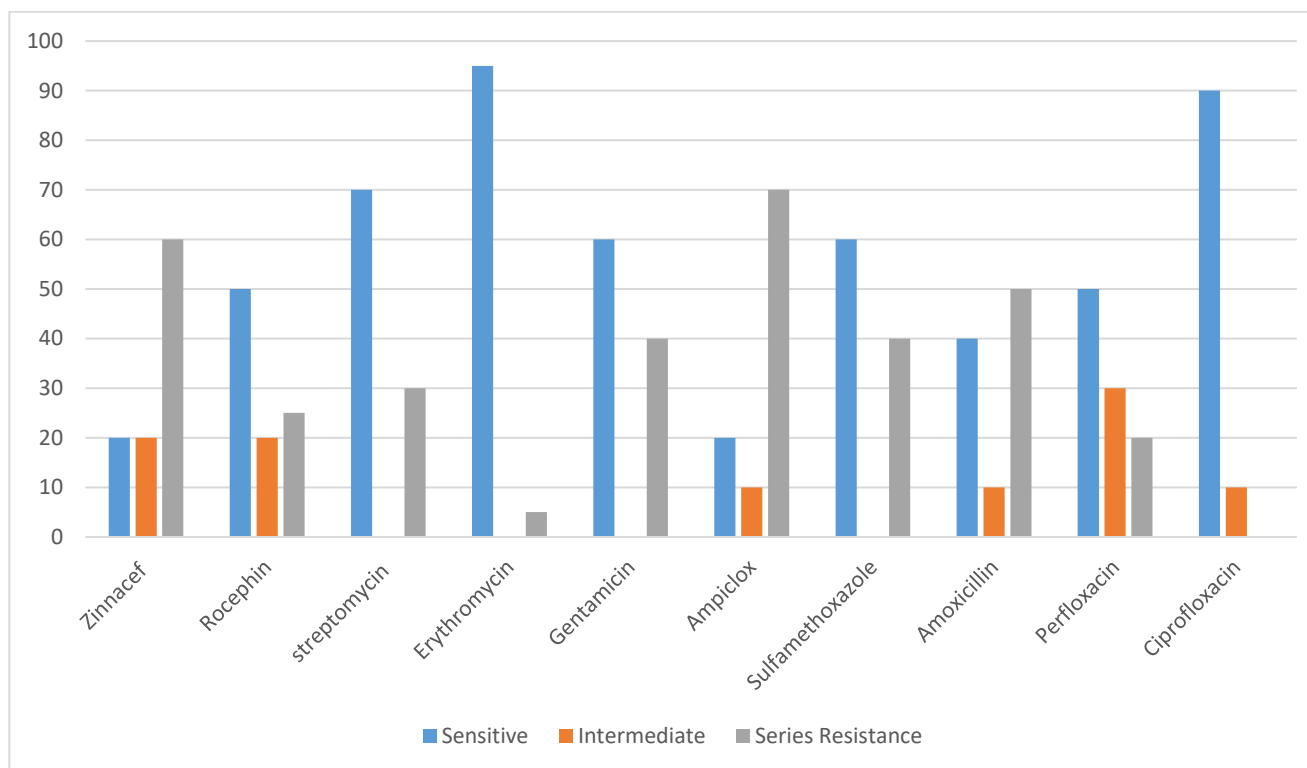


Fig. 3. Antibiotics resistance profile of *Staphylococcus aureus*

Table 7. Antibiotics susceptibility pattern of gram-negative bacteria from raw cow milk

Antibiotics profile (%)	Isolates											
	<i>Salmonella</i> (n=12)			<i>Klebsiella</i> (n=4)			<i>Escherichia coli</i> (n=11)			<i>Pseudomonas aeruginosa</i> (n=6)		
	S	I	R	S	I	R	S	I	R	S	I	R
AM	4(33.3)	3(25)	5(41.6)	1(25)	1(25)	2(50)	4(36.3)	2(18.2)	5(45.5)	2(33.3)	0	4(66.6)
PEF	10(83.3)	0	2(16.6)	2(50)	1(25)	1(25)	4(36.3)	5(45.5)	3(27.3)	1(16.6)	0	5(83.3)
SP	10(83.3)	1(8.3)	1(8.3)	1(25)	3(75)	0	8(72.7)	2(18.2)	1(9.09)	2(33.3)	3(50)	1(16.6)
CH	7(58.3)	3(25)	2(16.6)	3(75)	1(25)	0	6(54.5)	0	5(45.5)	4(66.6)	0	3(33.3)
GEN	6(50)	2(16.6)	4(33.3)	0	0	4(100)	8(72.7)	1(9.09)	3(27.3)	2(33.3)	1(16.6)	3(33.3)
AU	2(16.6)	0	10(83.3)	1(25)	0	3(75)	4(36.3)	2(18.2)	5(45.5)	1(16.6)	0	5(83.3)
OFX	7(58.3)	2(16.6)	3(25)	1(25)	3(75)	0	10(90.9)	1(9.09)	1(9.09)	4(66.6)	2(33.3)	0
S	2(16.6)	1(8.3)	9(75)	0	0	4(100)	2(18.2)	4(36.3)	5(45.5)	0	0	6(100)
SXT	1(8.3)	3(25)	8(66.6)	0	1(25)	3(75)	2(18.2)	1(9.09)	8(72.7)	1(16.6)	2(33.3)	3(33.3)
CPX	11(95)	0	0	3(75)	1(25)	0	11(100)	0	0	6(100)	0	0

KEY S=Sensitive, I= Intermediate, R=Resistant, AM=Amoxicillin, PEF=Pefloxacin, SP=Sparfloxacin, CH=Chloramphenicol, CN=Gentamicin, AU=Augmentin, OFX=Ofloxacin, S=Streptomycin, SXT= Sulfamethoxazole, CPX=Ciprofloxacin

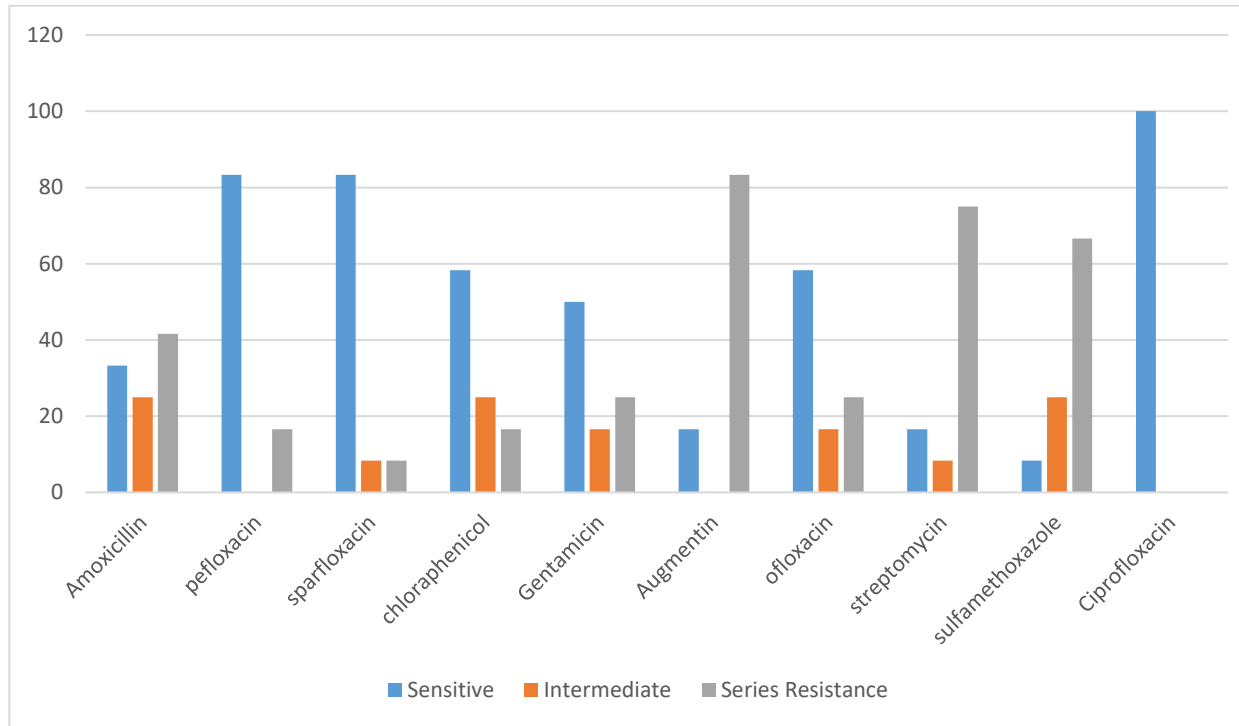


Fig. 4. Antibiotics resistance profile of *Salmonella*

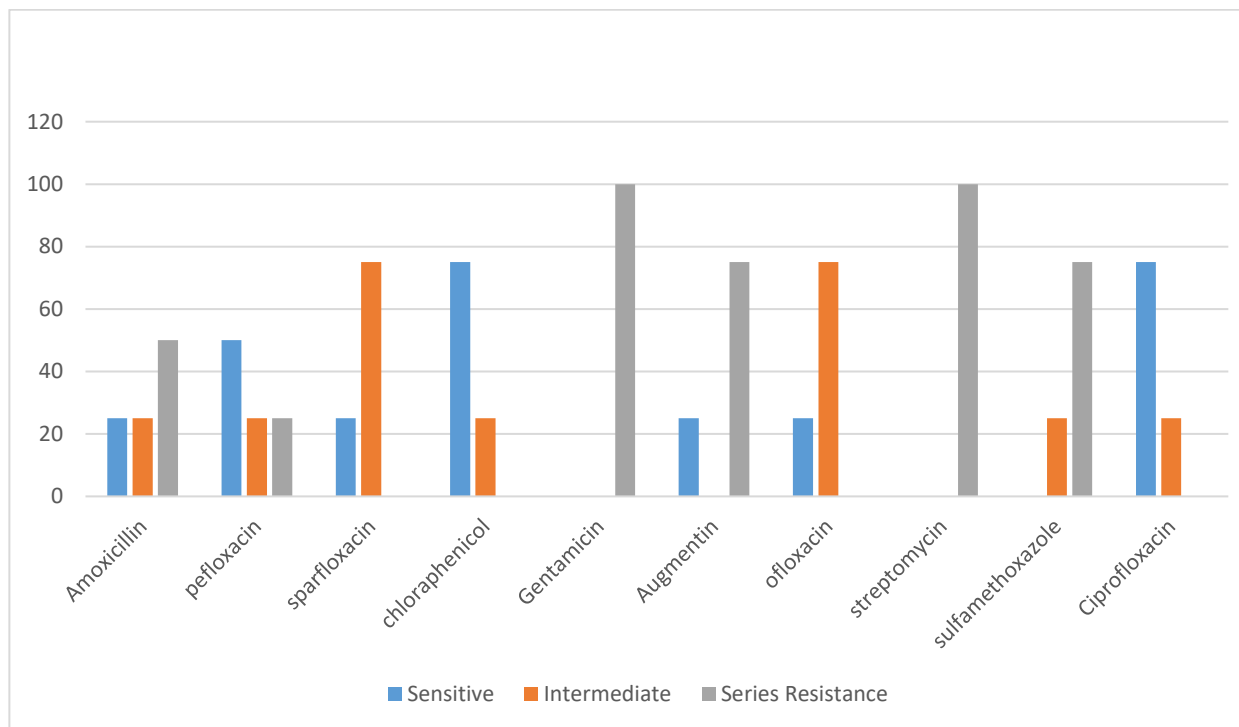


Fig. 5. Antibiotics resistance profile of *Klebsiella* spp

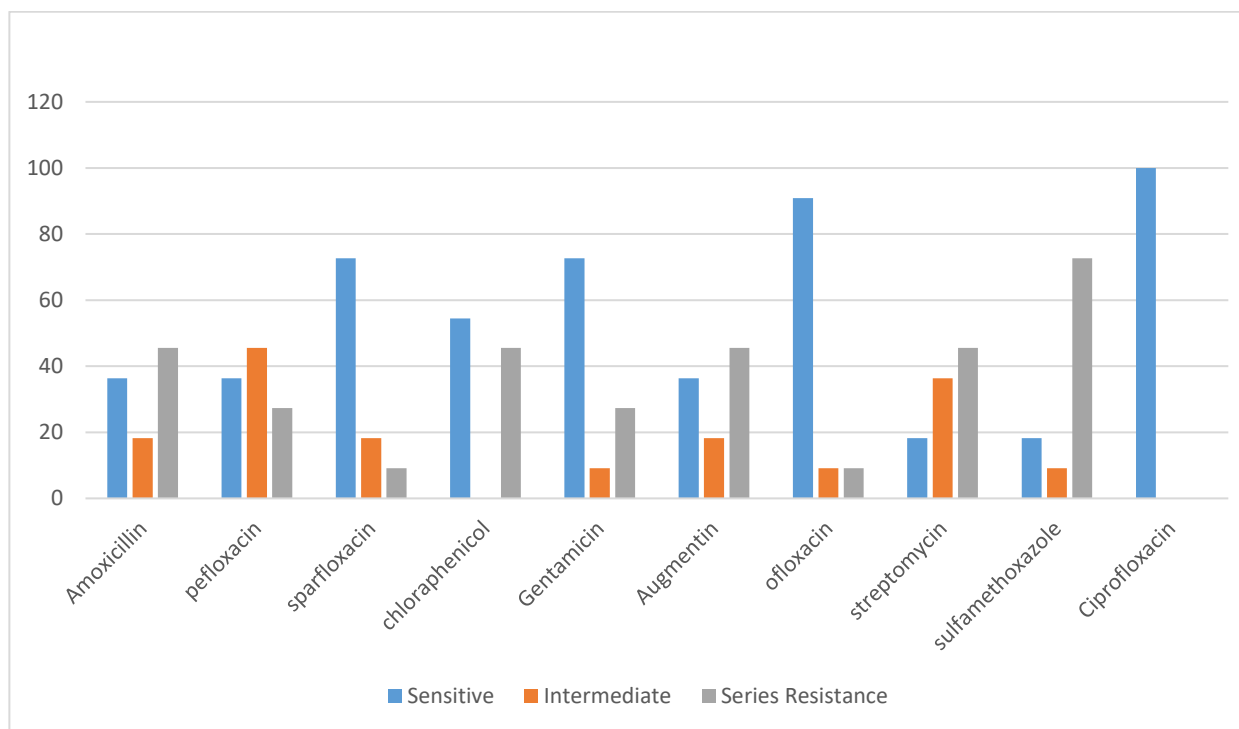


Fig. 6. Antibiotics resistance profile of *Escherichia coli*

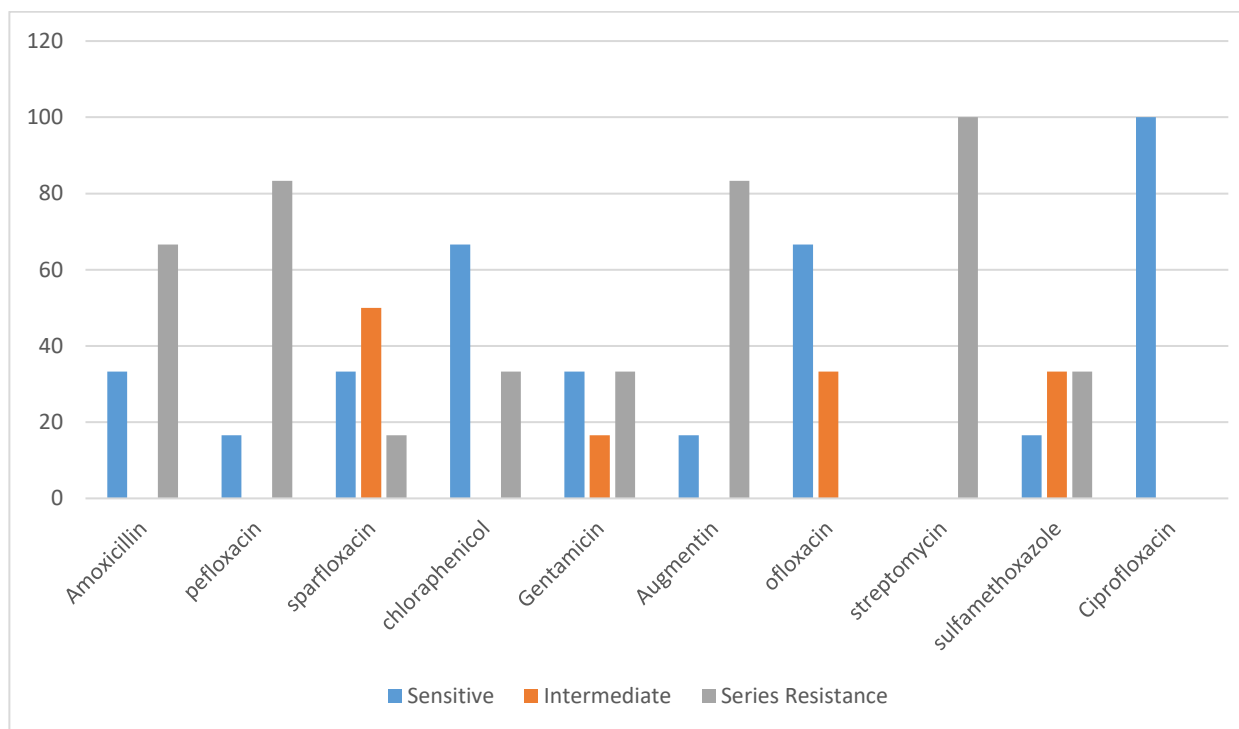


Fig. 7. Antibiotics resistance profile of *Pseudomonas aeruginosa*

4. DISCUSSION

The highest mean value of bacteria load of raw cow milk sample from Wednesday market was 8.5×10^6 CFU/ml (48.5%), whereas the moderate mean value of bacteria load from FUDMA take off campus was 7.99×10^6 CFU/ml (45.6%) While the lowest mean value was obtained from Wednesday Market with 1.04×10^6 CFU/ml (5.9%).

Morphological characteristics of the bacteria on Eosin Methylene Blue, Salmonella Shigella, Cetrimide and mannitol salt agar. On Eosin Methylene Blue agar *Escherichia coli* and *Klebsiella* were identified, *Escherichia coli* appeared green with a metallic sheen, and *Klebsiella* pink with no sheen mucoid. On Cetrimide *Pseudomonas aeruginosa* appeared green. On Salmonella-Shigella agar *Salmonella* was identified which appeared colorless with a black dot, and on Mannitol salt agar *Staphylococcus aureus* was identified as round and transparent.

The distribution of the bacteria in different locations, Wednesday market 20 were isolated *Staphylococcus aureus* (7), *Salmonella* spp. (5), *Klebsiella* spp. (2), *Pseudomonas aeruginosa* (2) and *Escherichia coli* (4), in hospital road 17 were isolated with *Staphylococcus aureus* (5), *Salmonella* spp. (3), This agree with the finding of Jamila et al. [33], *Klebsiella* spp. (2), *pseudomonas aeruginosa* (3) and *Escherichia coli* (4), in front of FUDMA take off campus 16 were isolated with *Staphylococcus aureus* (8), *Salmonella* (4), *Klebsiella* (0), *Pseudomonas aeruginosa* (1) and *Escherichia coli* (3). *Klebsiella* spp. were not present in front of FUDMA take off campus.

The percentage of bacteria isolates. This study revealed that *Staphylococcus aureus* has the highest occurrence with 37.8% followed by *Salmonella* spp. with (22.6%), *Pseudomonas aeruginosa* (11.3%), *Escherichia coli* (20.8%) and *Klebsiella* spp. (7.5%). This finding does not agree with the findings of (Nwosu et al., 2017) who reported that *E. coli* (86.7%) and *Salmonella* spp. (86.7%), *Staphylococcus aureus* (80.0%), *Klebsiella* spp. (73.3%) and *Pseudomonas aeruginosa* (66.7%). Reta et al. (2016) reported a prevalence rate of (24.2%) for *Staphylococcus aureus* in cow milk consumed at Jigjiga City, Ethiopia and (a 20.8%) prevalence rate for *Escherichia coli* reported by Makat et al. (2014) isolated from locally processed Cow milk products in Nassarawa state.

The antibiotics resistant to gram-positive *Staphylococcus aureus*, Gentamicin, and Ciprofloxacin were sensitive at 95% and 90% respectively this does not correlate with the study of (Rokeya et al., 2019) which show sensitivity to ciprofloxacin 64% and gentamicin 93%. The organism was also found to be resistant to Ampiclox and Zinnacef with 70% and 60% respectively.

The antibiogram of gram-negative bacteria all the isolates were highly sensitive to ciprofloxacin *Salmonella* was found to be also sensitive to Perfloracin and Sparfloracin with 83.3% respectively, also resistant to Augmentin (83%) and In streptomycin (83.3%). In a study by Makat et al., (2014), *Salmonella* spp. was also sensitive to both Perfloracin (43%) and Sparfloracin (85.7%) and also resistant to Augmentin (14.4%) and Streptomycin (14.2%) (Makat et al., 2014). *Klebsiella* was sensitive to chloramphenicol (75%) and resistant to gentamicin and streptomycin with 100%. *Escherichia coli* was sensitive to Ofloxacin (90.9%) (but resistant to Sulfamethoxazole (72.7%). *Pseudomonas aeruginosa* was resistant to streptomycin (100%) and Augmentin (83.3%). This study does not correlate with the study of (Nwosu et al., 2017).

5. CONCLUSION

The highest mean value of bacteria load of raw cow milk sample from Wednesday market was 8.5×10^6 CFU/ml (48.5%), whereas the moderate mean value of bacteria load from FUDMA take off campus was 7.99×10^6 CFU/ml (45.6%) While the lowest mean value was obtained from Wednesday Market with 1.04×10^6 CFU/ml (5.9%).

The distribution of the bacteria in different locations, Wednesday market 20 were isolated *Staphylococcus aureus* (7), *Salmonella* spp. (5), *Klebsiella* spp. (2), *Pseudomonas aeruginosa* (2) and *Escherichia coli* (4), in hospital road 17 were isolated with *Staphylococcus aureus* (5), *Salmonella* spp. (3), *Klebsiella* spp. (2), *pseudomonas aeruginosa* (3) and *Escherichia coli* (4), in front of FUDMA take off campus 16 were isolated with *Staphylococcus aureus* (8), *Salmonella* (4), *Klebsiella* (0), *Pseudomonas aeruginosa* (1) and *Escherichia coli* (3). *Klebsiella* spp. were not present in front of FUDMA take off campus.

The percentage of bacteria isolates. This study revealed that *Staphylococcus aureus* has the highest occurrence with 37.8% followed by

Salmonella spp. with (22.6%), *Pseudomonas aerogenosa* (11.3%), *Escherichia coli* (20.8%) and *Klebsiella* spp. (7.5%). This agrees with the finding of McKinnon *et al.* and Thomas, [32 and 41].

The antibiotics resistant to gram-positive *staphylococcus aureus*, Gentamicin, and Ciprofloxacin were sensitive at 95% and 90% respectively. The antibiogram of gram-negative bacteria all the isolates were highly sensitive to ciprofloxacin *salmonella* was found to be also sensitive to Perfloxacin and Sparfloxacin with 83.3% respectively, also resistant to Augmentin (83%) and In streptomycin (83.3%). The results obtained show that there is the presence of pathogenic microorganisms that may be a potential source of food-borne infection that may result in food-borne diseases in the consumers of these sampled products

Thomas *et al.* [42]. The total viable bacteria count in all samples was above the standard, according to the Nigerian Agency for Food, Drugs Administration and Control NAFDAC (2009), the microbial load limited for total liable colony count is 1.0×10^2 cfu/ml.

This study also found high levels of resistance to commonly prescribed antibiotics augmentin, streptomycin, gentamicin and sulfamethoxazole in the bacterial isolates. This calls for the strengthening of regulations that cover the sale, distribution, dispensing, and prescription, of veterinary antibiotics. Khatun *et al.* and Kurweil *et al.* [28,29]. This is because antibiotic-resistant bacteria may cause complicated, untreatable, and prolonged infections in humans, leading to higher healthcare costs and sometimes death.

6. RECOMMENDATION

It is recommended that good sanitary measures should be taken by the people handling the cows, these measures should include proper handling of the cow, personal hygiene, treatment of udder infection of the cow, use of hygienic milking and processing equipment, and improved milk handling environment. It must be ensured that the cows are always in good health condition.

ACKNOWLEDGEMENT

We should like to thank all those who have assisted in data collection and analysis of this

research study. Thank you for your time and efforts. God bless all.

COMPETING INTERESTS

Authors have declared that no competing interests exist.

REFERENCES

1. Srinu B, Kumar AV, Kumar MS, Narayana BVL, Rao TM. Assessment of microbiological quality and associated health risks of raw milk sold in and around Hyderabad city. *International Journal of Pharma and Bio Sciences*. 2012;3(4): 609-614.
2. Ensminger ME. *Foods and nutrition encyclopedia*. 1994;2.1:2415.
3. Oliver SP, Jayarao BM, Almeida RA. Foodborne pathogens in milk and the dairy farm environment: food safety and public health implications. *Foodborne Pathogens & Disease*. 2005;2(2):115-129.
4. Mubarack HM, Doss A, Dhanabalan R, Balachander S. Microbial quality of raw milk samples collected from different villages of Coimbatore District, Tamilnadu, South India. *Indian Journal of Science and Technology*. 2010;3(1):61-63.
5. Virpari PK, Nayak JB, Brahmabhatt MN, Thaker HC. Study on isolation, molecular detection of virulence gene and antibiotic sensitivity pattern of *Escherichia coli* isolated from milk and milk products. *Veterinary World*. 2013;6(8).
6. Gajdács M, Albericio F. Antibiotic resistance: from the bench to patients. *Antibiotics*. 2019;8(3):129.
7. Abdulkadir M, Mugadi AG. Bacteriological examination of fura da nono (fermented milk; cereals mix) sold in some selected areas of Birnin Kebbi Metropolis. *ARNP Journal of Science and Technology*. 2012; 2:333-340.
8. Adesokan IA, Odetoyinbo BB, Olubamiwa AO. Biopreservative activity of lactic acid bacteria on suya produced from poultry meat. *African Journal of Biotechnology*. 2008;7(20).
9. AJ B. *Microbiology of raw milk*. Dairy microbiology. 1990;1:163-208.
10. Bhaskar SV. Foodborne diseases—disease burden. In *Food safety in the 21st century* Academic Press. 2017;1-10.
11. Bramley AJ. Sources of *Streptococcus uberis* in the dairy herd: I. Isolation from

- bovine faces and from straw bedding of cattle. *Journal of Dairy Research*. 1982; 49(3):369-373.
12. Bramley AJ, McKinnon CH, Staker RT, Simpkin DL. The effect of udder infection on the bacterial flora of the bulk milk of ten dairy herds. *Journal of Applied Bacteriology*. 1984;57(2): 317-323.
 13. Brunelle BW, Bearson BL, Bearson SM. Chloramphenicol and tetracycline decrease motility and increase invasion and attachment gene expression in specific isolates of multidrug-resistant *Salmonella enterica* serovar Typhimurium. *Frontiers in Microbiology*. 2015;5:801.
 14. Butler MT, Wang Q, Harshey RM. Cell density and mobility protect swarming bacteria against antibiotics. *Proceedings of the National Academy of Sciences*. 2010; 107(8):3776-3781.
 15. Dewi G. Investigating the Potential of Lemongrass Essential Oil Against Multidrug-Resistant *Salmonella Heidelberg* in Broiler Chickens (Doctoral dissertation, University of Minnesota); 2018.
 16. Falegan CR, Akere GA. Isolation of salmonella spp in 'wara'(local cheese) from three different locations in ado-ekiti, ekiti state, Nigeria. *The experiment*. 2014; 23(4):1628-1634.
 17. Fenlon DR, Logue DN, Gunn J, Wilson J. A study of mastitis bacteria and herdmanagement practices to identify their relationship to high somatic cell counts in bulk tank milk. *British Veterinary Journal*. 1995;151(1):17-25.
 18. Galton DM, Petersson LG, Merrill WG, Bandler DK, Shuster DE. Effects of premilking udder preparation on bacterial population, sediment, and iodine residue in milk. *Journal of dairy science*. 1984; 67(11):2580-2589.
 19. Gebreyes WA, Thakur S, Davies PR, Funk JA, Altier C. Trends in antimicrobial resistance, phage types and integrons among *Salmonella* serotypes from pigs, 1997–2000. *Journal of Antimicrobial Chemotherapy*. 2004;53(6):997-1003.
 20. Gehringer G. Multiplication of bacteria during farm storage, in factors influencing the bacteriological quality of raw milk. *IDF Bulletin*. 1980;120.
 21. Gonzalez RN, Jasper DE, Bushnell RB, Farver TB. Relationship between mastitis pathogen numbers in bulk tank milk and bovine udder infections in California dairy herds. *Journal of the American Veterinary Medical Association*. 1986;189(4):442-445.
 22. Harshey RM. Bacterial motility on a surface: Many ways to a common goal. *Annual Reviews in Microbiology*. 2003;57(1):249-273.
 23. Helms M, Vastrup P, Gerner-Smidt P, Mølbak K. Excess mortality associated with antimicrobial drug-resistant *Salmonella Typhimurium*. *Emerging infectious diseases*. 2002;8(5):490.
 24. Hogan JS, Smith KL, Hoblet KH, Todhunter DA, Schoenberger PS, Hueston WD, Conrad HR. Bacterial counts in bedding materials used on nine commercial dairies. *Journal of dairy science*. 1989;72(1):250-258.
 25. Jackson H, Clegg LFL. The microflora of raw bulk tank milk. *Canadian Journal of Microbiology*. 1966;12(3):429-432.
 26. Jeffrey DC. J. Wilson. Effect of mastitis-related bacteria on the total bacteria counts of bulk milk supplies. *J. Soc. Dairy Technol*. 1987;40(2):23.
 27. Kearns DB. A field guide to bacterial swarming motility. *Nature Reviews Microbiology*. 2010; 8(9):634-644.
 28. Khatun MM. Isolation, Identification and Antibiotic Sensitivity Test of Bacteria in Raw Cow Milk Obtained from Different Sources. *EC Microbiology*. 2020;16:32-41.
 29. Kurweil R, Busse M. Total count and microflora of freshly drawn milk. *Milchwissenschaft*. 1973;28:427.
 30. Lippolis JD, Brunelle BW, Reinhardt TA, Sacco RE, Thacker TC, Looft TP, Casey TA. Differential gene expression of three mastitis-causing *Escherichia coli* strains grown under planktonic, swimming, and swarming culture conditions. *Msystems*. 2016;1(4):10-1128.
 31. Mackenzie E. Thermotrophic and psychrotrophic organisms on poorly cleansed milking plants and farm bulk milk tanks. *Journal of Applied Microbiology*. 1973;36(3):457-463.
 32. McKinnon CH, Rowlands GJ, Bramley AJ. The effect of udder preparation before milking and contamination from the milking plant on bacterial numbers in bulk milk of eight dairy herds. *Journal of Dairy Research*. 1990;57(3):307-318.
 33. Miriam Jamila, Md Zulfekar Ali, Shaharin Sultan, Md Ariful Islam and Mst Minara Khatun. Isolation, Identification and Antibiotic Sensitivity Test of Bacteria in Raw Cow Milk Obtained from Different

- Sources. EC Microbiology. 2020;16.4: 32-41.
34. Olson JC. Jr, G. Mocquat. Milk and Milk Products. p. 470. In Microbial Ecology of Foods. 1980;II.
 35. Palmer J. Contamination of milk from the milking environment. Kieler Milchwirtschaftliche Forschungsberichte. 1981;33(4):307-316.
 36. Pankey JW. Premilking udder hygiene. Journal of dairy science. 1989;72(5):1308-1312.
 37. Partridge JD, Harshey RM. Swarming: flexible roaming plans. Journal of bacteriology. 2013; 195(5):909-918.
 38. Rehman MU, Rashid M, Sheikh JA, Bhat MA. Molecular epidemiology and antibiotic resistance pattern of enteropathogenic Escherichia coli isolated from bovines and their handlers in Jammu, India. Journal of Advanced Veterinary and Animal Research. 2014;1(4):177-181.
 39. Richardson GH. Standard methods for the examination of dairy products. (No Title); 1985.
 40. Smith K, Peter K, Daniela H, Melchior S. Food borne pathogenic microorganisms and natural toxins. Food drug Administration center food safety, Applied Nutrition. 2007;10:119-150.
 41. Thomas SB. the microflora of bulk collected milk. I; 1974.
 42. Thomas SB, Druce RG, King KP. The microflora of poorly cleansed farm dairy equipment. Journal of Applied Bacteriology. 1966;29(2):409-422.
 43. Tolle A. The Microflora of the Udder, P4 In Factors Influencing The Bacteriological Quality of Raw Milk. International Dairy Federation Bulletin; 1980.
 44. Zehner MM, Farnsworth RJ, Appleman RD, Larntz K, Springer JA. Growth of environmental mastitis pathogens in various bedding materials. Journal of dairy science. 1986;69(7):1932-1941.

© 2024 Abdullahi et al.; This is an Open Access article distributed under the terms of the Creative Commons Attribution License (<http://creativecommons.org/licenses/by/4.0>), which permits unrestricted use, distribution, and reproduction in any medium, provided the original work is properly cited.

Peer-review history:

The peer review history for this paper can be accessed here:
<https://www.sdiarticle5.com/review-history/111083>