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Prevalence of coagulase positive pathogenic *Staphylococcus aureus* in milk and dairy products

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ABSTRACT

Key words:

Staphylococcus aureus; coagulase; PCR

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erythromycin (\circ ^r%) and other strains were highly sensitive for gentamicin(\circ ^r%), norfloxacin (\circ ^r%), vancomycin (\circ ^r%) and linezolid (\circ \circ %). Then Polymerase chain reaction (PCR) was applied on positive strains of *S.aureus* to detect their virulence genes *staphylococcal* protein A (SpA) and coagulase gene (Coa) which were detected with percentage (Coa) of tested isolates, and beta-lactam resistance gene (Coa) was found in (Coa) of examined Coa strains.

\.INTRODUCTION

Staphylococcus aureus is one of the most common facultative pathogenic bacterium that has long been recognized as a challenge in both human and veterinary medicine[1], organism of bovine mastitis ,which is a ,complicated and economically common unbearable disease in dairy animals worldwide. Staphylococci can have a wide variety of virulence factors which allow the bacteria to avoid the immune system and contribute to increased severity of infections. Most of these factors have been initially founded in S. aureus and include surface proteins (Protein A, clumping factor, fibronectin binding proteins or iron regulated surface determinants), capsular polysaccharides, molecules involved in biofilm formation (for example polysaccharide intercellular adhesin) or toxins (pore-forming toxins, toxins that act as superantigens). Cell wall adhesins that recognize extracellular matrix proteins and Some enzymes (coagulase, staphylokinase and proteases) also contribute to immune evasion and host tissue penetration [4],[5],[0],[7].

Coagulase substance is seen as the fundamental danger factor that coagulation plasma and coats the bacterial cell, so prevent phagocytosis[\(^\)]through enables staphylococci to got into a fibrin meshwork, spread and go against opsonophagocytic instrument of host safe cells [A]. Production of coagulase is a crucial phenotypic feature and the major factor for identifying *S*. determinant aureus strains, the variability at the Υ' end coding region of the coa gene has been used for genotyping of S. aureus strains from humans and animals [9]. Staphylococcal protein A (spA) is a layer bound exoprotein of bacterial cell wall that consider a critical destructiveness factor which ruin opsonization by serum supplement phagocytosis and of Polymorphonuclear leukocytes through confining to FC locale of immunoglobulins[\`\\]. S.aureus cultivated a high protected against a wide grouping of hostile to contamination specialists which increase their danger and inconvenience in treatment [\\]. The most important antibiotic resistant genes of S.aureus strains was blaZ gene which coded for β lactamase important virulence factor that impair opsonization by serum complement and phagocytosis of Polymorphonuclear leukocytes through binding to FC region of immunoglobulins[17].

Presence of *S. aureus* in milk and dairy items even in low numbers has been viewed as a general wellbeing danger since it has been laid out that *Staphylococcus* enterotoxins continue for longer lengths in the polluted milk and dairy items, even after the microorganisms lose its viability['\"].

Improper food handling either in the home or food industry constitutes a major factor leading to *Staphylococcus* food poisoning outbreaks. Only few outbreaks can be traced directly to contamination during food processing [\frac{1}{2}].

The proper heat treatment followed by the refrigeration can minimize the chance of contamination with *S. aureus*. In our country it is commonly noticed that during heat treatment of milk, the temperature not rise up to the boiling point many a time or even if it reaches, consumers do not boil it enough. [\forall^o].

V. MATERIALS AND METHODS

7.1 Samples collection

A total of ' random samples of raw cow milk, raw buffalo milk, kareish cheese, baladi yogurt (' of each) were collected from different shops and supermarkets at Menofia and Gharbia Governorate. The collected

samples were transferred directly to the animal health research laboratory (Tanta branch) in an ice box under complete aseptic conditions for bacteriological examination.

Y.Y. Bacteriological examination[\\],[\\].

Samples were homogenized in peptone water, maintained for \hat \for \cdot C then samples were pre-enriched into nutrient broth and incubated at TY°C for Y f hrs. It was used for the growth and propagation of isolates before platting under aerobic condition. A loopful from incubated nutrient broth was streaked on Baired parker agar plates (oxoid) and incubated at TVoC for Y \(\xi - \xi \lambda \) hrs Positive samples (showed black shiny colonies with clear halo zone around colonies and opaque precipitation). Also, a loopful from incubated nutrient broth was streaked on Mannitol salt agar (Oxoid) and incubated at TV oC for YE - EA hrs Positive samples (yellow colonies and turned media to colorless). These colonies were kept in Brain Heart Infusion broth for biochemical identification **PCR** and examination

Y.Y.\. Morphological examination [\\].

The suspected *s.aureus* isolates were stained by Gram stain for morphological examination .

Y.Y.Y. Biochemical identification [\ 4].

Indol test, catalase test , urease test, oxidase test , coagulase test and $\beta-\text{hemolysis}$ test .

Y.Y. Antibiotic sensitivity test:

The obtained bacterial isolates were tested in vitro for their susceptibility to the following antimicrobial discs: pencillin (P) ' · IU, gentamicin (CN) ' · mcg, erythromycin (E) ' o mcg, vancomycin (VA) ' · ml , clindamycin(DA) ' mcg, norfloxacin (NOR) ' ·

mcg and linezolid (LNZ) r · mcg. According to $[^{r}$ ·] and the level of sensitivitywas deciphered agreeing $[^{r}$].

Y.Y.£. Molecular detection of Coagulase Positive S.aureus:

DNA was taken out from the bound S.aureus using QIAamp DNA limited scope unit. It was applied on \(\) sporadic limits. PCR Master Mix and cycling conditions of the foundations during PCR was prepared according to Emerald Amp GT PCR mastermix (Takara) pack. Oligonucleotide foundations used in PCR have express gathering and improve a specific thing as shown in (table, \(\)).

DNA tests were upgraded in an amount of Youl as follows: ۱۲.0 µl of Emerald Amp GT PCR expert mix, \ \ \mu \| of each and every preparation of γ· pmol centers, ٤.0 μl of water and γ μl of format DNA. The reaction was acted in a Biometra warm cycler. Temperature and time conditions of the starters during PCR were applied .Aliquots of upgraded PCR things were electrophoresed in 1.0 % agarose gel (ABgene) in 'x TBE pad at room temperature. For gel assessment, \o \mu l of PCR things were stacked in each gel opening. A \. bp DNA ladder (QIAGEN Inc, Valencia, CA, USA) was used to conclude the piece sizes. The gel was caught by a gel documentation structure and the data was inspected through PC programming.

Table ($^{\circ}$): Sequence and amplicon size of primers used for detection S. aureus virulence and resistance genes:

Target gene	Sequence	Amplified product	Reference	
S. aureus	AC GGAGTTACAAAGGACGAC	140. pp	F	
rrs rRNA	AGCTCAGCCTTAACGAGTAC	т тот ор	[۲ ۲]	
	ATA GAG ATG CTG GTA CAG G	Four different types of		
Coa		bands may be detected		
		۳۰، bp	F & # 3	
	GCT TCC GAT TGT TCG ATG C	٤٣٠ bp	[۲ ۳]	
		ov. bp		
		77. bp		
C	TCA ACA AAG AAC AAC AAA ATG C	777 h.c.	F¥ 4 7	
Spa	GCT TTC GGT GCT TGA GAT TC	YY7 bp	[
blaZ	TACAACTGTAATATCGGAGGG	۸۳۳ bp	[6 7]	
	CATTACACTCTTGGCGGTTTC]	[٢٥]	

Table (7): Cycling conditions of the primers during cPCR:

Target gene	Primary denaturation	Secondary denaturation	Annealing	Extension	No. of cycles	Final extension
S. aureus	٩٤°C	٩٤°C	oo°C	۸4.°C	٣0	۸4.°C
rrs rRNA	٥ min.	۳۰ sec.	٤٠ sec.	۱.۲ min.	, 5	۱۲ min.
Coa	٩٤°C	٩٤°C	∘o∘C	۸4.°C	30	۸4.°C

	° min.	۳۰ sec.	٤٠ sec.	٤٥ sec.		۱۰ min.
Spa	4 £ °C	٩٤°C	oo°C YY°C To		۸4.°C	
	° min.	۳۰ sec.	۳۰ sec.	۳۰ sec.	, 0	∨ min.
blaZ	9 £°C	٩٤°C	o, °C	۸4.°C	٣٥	۸4.°C
	٥ min.	۳۰ sec.	٤٠sec	o · sec	, 5	۱۰ in.

F.RESULTS

7.1. Incidence of *S. aureus* isolated from examined samples.

A total of of isolates of S. aureus were recovered from '·· Samples represented as Υ1/Υο (Λέ٪), 1/Υο (ΊΑ٪), Λ/Υο (ΥΥ٪), 1/Υο (Υέ٪) from of raw cow milk, raw buffalo milk, kareish cheese, baladi yogurt, respectively, (Table Υ).

$^{\gamma}$. Table ($^{\gamma}$): Incidence *S. aureus* isolated from examined samples:

Samples	Number of examined samples	Number of positive samples	% of positive samples
Raw cow milk	70	۲۱	٨٤%
Raw buffalo milk	۲0	١٧	٦٨٪
Kareish cheese	۲٥	٨	٣٢٪
Baladi yogurt	۲٥	٦	7 £ %
Total No	١	٥٢	٥٢٪

Y.T.Phenotypic Identification of Staphylococcus aureus:

In this study Staphylococcus aureus were confirmed through Gram staining, S.aureus is non-motile facultative anaerobic

Gram-positive cocci.graps likr cluster (Figure 1).

The isolated strains were cultured, on Baired-Parker agar: showed black shiny colonies with clear halo zone around colonies and opaque zone of precipitation figure ($^{\Upsilon}$), on Mannitol salt agar: showed (yellow colonies and turned media to colorless) figure ($^{\Upsilon}$).

Biochemically, Negative results were recorded on Indol test, Oxidase test and positive results were recorded on coagulase test (fig $^{\xi}$) and β – hemolysis test as show in (fig $^{\circ}$)

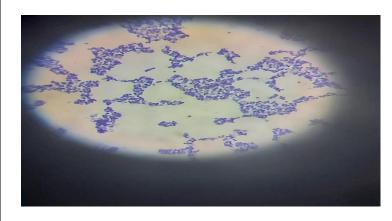


Fig (\) S.aureus under microscope after
Gram's staining appear as grape like cluster

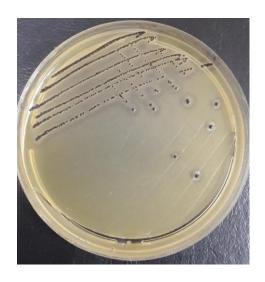


Fig (Y) Colonies of *S.aureus* on baired parker agar media.



Fig (*) S.aureus on mannitol salt agar.

r. r.Prevalence of Coagulase-positive



Samples that showed phenotypically positive growth on Baird–Parker medium investigated for their coagulase activity ,The prevalence of Coagulase-positive S.aureus was 15/71(77%), 7/10(51%), 7/10(77%), 7/10(77%) from raw cow milk, raw buffalo milk, kareish cheese, baladi yogurt , respectively.

Table(1): results of coagulase test:

sam	No,	No.	No.	% of	No.	% of
ples	of	of	of	coag	of	coag
pies	exa	s.au		ulas		ulas
			coag ulas		coag ulas	
	min	reu		e magit		e
	ed	S	e magit	posit	e	nega
	sam	isol	posit	ive	nega	tive
	ples	ates	ive	isola	tive	isola
			isola	tes	isola	tes
	.		tes	u u •/	tes	/• ر س
Ra	70		_	11%	Y	٣٤%
W		١	٤			
cow						
mil						
k						
Ra	40			٤١٪.	١.	٥٩٪
\mathbf{w}		٧				
buf						
falo						
mil						
k						
Kar	7			٧٥٪	۲	۲٥٪
eish						
che						
ese						
Bal	40			٥٠٪	٣	٥٠٪
adi						
yog						
urt						
Tot	١			_	7 7	_
al		۲	•			

Fig (\$\xi\$) coagulase test show that clot form when bacterial cell are incubated with plasma.

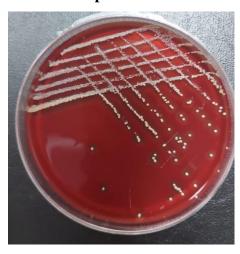


Fig (°) S.aureus showed β – hemolysis on blood agar media.

۳.٤. Antibiotic sensitivity test results:



Fig (7) Antibiotic sensitivity test of *S.aureus*.

Table (°): Invitro anti-microbial sensitivity test for isolated *S. aureus*

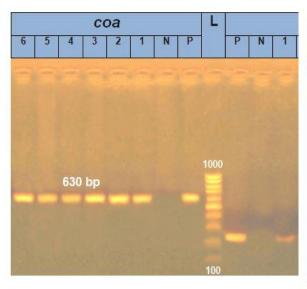
Antimicrobial disc	Code	Disk conce	Sensitive		ve Interm ediate		Resista nt		AA
		ntrati	No.	%	No	%	No	%	
		on			•		•		
	P	١٠IU	٣	۲.	•	٠.	17	٨	R
Penicillin						٠		•	
	DA	۲۰ μg	٦	٤٠	•	٠.	٩	7	R
Clindamycin						٠		•	
	LNZ	۳۰ μg	٩	۲.	•	٠.	۲	٤	S
Linezolid						•		٠	
	Е	۱٥ μg	۲	٤٠	١	۲.	٨	0	R
Erythromycin						٦		٣	
	CN	۱۰ μg	10	١.	•	٠.	•	٠.	S
Gentamicin		, -		•		•		٠	
	VA	۳۰ μg	١٣	٨٦	•	٠.	۲	١	S
Vancomycin						٠		٤	
	NOR	۱۰ μg	١٤	98	١	٦.	٠	٠.	S
Norfloxacin						٦		٠	

AA: Antibiogram activity

% Percentage in relation to isolates of *S. aureus* no=1°.

۳.٥. PCR results.

Detection of virulence and resistant genes of *S.aureus*. PCR using primers fragments for detection of two virulence genes, These genes were *staphylococcal spA* and coagulase (*coa*). It was applied on six isolates of *S. aureus*. all tested isolates harbored these genes('\.'.'/), Also these isolates were tested for detection of one resistance gene beta- lactam resistance gene (blaZ) by uniplex PCR. the results revealed that ("\"'/) of tested isolates harbored this gene.



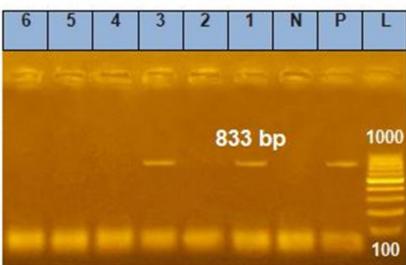


Fig (V): Agarose gel electrophoresis pattern of PCR for detection of staphylococcal spa and coagulase (coa) genes at TTT and TTT bp respectively.

L: Ladder from ' · · · bp to ' · · · · bp

Pos: Positive control: S.aureus ATCC ' o ٩ ٢ ".

Neg: Negative control: Field withdraw that
were attempted and attested to be negative
by PCR for the associated characteristics in

Way 1, 7, 7, 2, 0, 1: positive upgrade of staphylococcal virulance characteristics (spa and coagulase characteristics).

R.L.Q.P

Fig. ($^{\wedge}$): Agarose gel electrophoresis pattern of PCR for detection of *blaZ* gene of *S*.

aureus at $^{\wedge \forall \forall}$ bp.

L: Stepping stool from '.. bp to '... bp.

Pos: Positive control: S.aureus ATCC Yoqyw.

Neg: Negative control: Field confine that were tried and affirmed to be negative by PCR for the connected qualities in R.L.Q.P

Path ', ": Positive enhancement of blaZ quality at "" bp

4.DISCUSSION

S. aureus can convey heat stable enterotoxins achieving food intoxication in human with fluctuating power El-Makarem and Amer[77] along these lines, the ongoing audit was planned to perceive and depict coagulase positive S. aureus.

In this study of bounds of S. aureus (of %) were recovered from \... models from unrefined milk and its things, which nearly agreed with the eventual outcomes of Lílian et al., (Y·)) who reported higher event of S.aureus in rough milk (٦٨٪) But it went against Ahmadi [YV] and El-Sayed [Y9] who bound S.aureus at recurrence rate (15%). Moreover, this is higher prevalence at whatever point diverged from results of Zeinhom [7.] who uncovered that S. aureus was isolated from the rough milk tests with an overall inescapability of \rightarrow\.'\.(\rightarrow\.'\.\) and saw that as 17% of unrefined milk tests outperformed beyond what many would consider possible as demonstrated by the Egyptian Standards [^r].

The high incidence in raw milk could be attributed to environmental pollution, crosscontamination between the milk and each other and poor handling during transportation or in milk collection centers, besides, shedding of *S. aureus* from infected animals is another cause of contamination of milk and dairy food.

Positive isolates tested for their coagulase activity and the results were on, YAX, YEX, NYX for raw cow milk, raw buffalo milk, kareish cheese and baladi yogurt respectively.

According to microbiological standards set out by the Egyptian authority, w cheese must be free from *S. aureus* .

In this study the regularity of *S. aureus* in Karish chese was ($\mbox{\ref{thm:charge}}$), which nearly agreed with the eventual outcomes of Mousa [$\mbox{\ref{thm:charge}}$] and Elmaghraby [$\mbox{\ref{thm:charge}}$] who nitty gritty that inescapability of $\mbox{\ref{thm:charge}}$, and $\mbox{\ref{thm:charge}}$ in Karish cheese in Egypt, these results were seen as low when appeared differently in relation to other Egyptian examinations of El - Malt [$\mbox{\ref{thm:charge}}$] and Sallam [$\mbox{\ref{thm:charge}}$] that uncovered S. aureus transcendence speeds of $\mbox{\ref{thm:charge}}$, and $\mbox{\ref{thm:charge}}$, from Karish cheese.

Variable reported incidence rates could be influenced by many factors such as different techniques of cheese production, storage conditions, type of cheese either manufactured from raw or pasteurized milk, and also related to personal hygiene of workers and unhygienic measures during production.

The bacterium (*S. aureus*) is liable to inactivation and elimination by heating, but as the cheese manufacturer in Egypt use raw milk which is most likely not subject to heating $[^{ro}]$.

The prevalence of *s.aureus* in baladi yogurt was YE% then the positive isolates tested for coagulase activity and the coagulase positive isolates was YY%.

This consider low recurrence when diverged from delayed consequences of $[^{r_1}]$

who reported that % of positive *s.aureus* isolates was YY% and £A% for plain and fruit yogurt respectively with Prevalence of Coagulase-positive *S. aureus* in the examined fermented dairy product samples *FO% and YYW for plain and fruit yogurt respectively.

The spread of antibiotic-resistant pathogens continues to challenge sustainable treatment options, with severe public health consequences.

The most common mechanism of resistance to antibiotics of coagulase-positive *staphylococci* is the synthesis of beta lactamases that inactivate penicillinase sensitive antibiotics. Resistance to beta-lactam antibiotics in coagulase- positive staphylococci is increasing and following resistance in human isolates. Some studies show that penicillin disk diffusion method showes deviation in the case of sensitive beta-lactam antibiotics in relation to the dilution method. [*Y].

Fifteen *S. aureus* detaches were moreover investigated for their shortcoming to antimicrobials ,concerning the results of serum poison weakness testing, by far most of *S. aureus* strains separated were impenetrable to

penicillin G ($^{\wedge \cdot \%}$),clindamycin($^{\neg \cdot \%}$), erythromycin ($^{\circ r\%}$) sensitive showed most raised responsiveness towards gentamicin ($^{\circ r\%}$), followed by norfloxacin ($^{\circ r\%}$), vancomycin ($^{\wedge \cdot \%}$) and linezolid ($^{\neg \cdot \%}$).

higher security from penicillin scraped by Abera [$^{\text{MA}}$] and Shi [$^{\text{MA}}$] in a degree of ($^{\text{MA}}$) and ($^{\text{MA}}$) independently.

Practically identical results got by Begum [5.] who revealed that S. aureus was AY.AT% and TY.YE% impenetrable to Penicillin-G and Amoxicillin, independently;

In like manner nearly to eventual outcomes of Abdeen [\sharp \] The MRSA strains showed assurance from penicillin (\S \), cefoxitin (\S \), and erythromycin (\S \).

The development of PCR-based methods provides a promising option for the rapid identification of bacteria. With this method, identification of bacterial pathogens can be made in hours, rather than days, conventional cultural methods require. PCR can also improve the level of detection due to its high sensitivity. Theoretically, only a few cells of pathogen are necessary to yield a positive diagnosis [٤٢]. The molecular identification of the coa gene in S. aureus strains was performed through amplification of the coa gene at (7%) bp. The use of the coa gene to detect S. aureus strains from milk origin was previously reported in two studies o [10] and [ξ^{η}] which improved the coa quality from mastitic milk and dairy things at Tr. and yo. bp. Moreover, Javid [ξξ] perceived the coa quality at ολξ bp, ολο bp, yoy bp, and λ.γ bp.

The findings of the present investigation showed that \cdots ? the examined *S. aureus* isolates ($^{7/7}$) harbored *coa* gene. Similar result obtained by [77]

who reported that detection of the *coa* gene from milk samples could help to assess the microbiological safety of raw milk intended for direct use in the dairy industry. The *coa* gene was detected in all isolates with the primers used, even though one of them was considered coagulase-negative both by the tube coagulase test and by the clumping factor assay.

The SpA gene is a major important surface protein of bacterial cell wall, which binds with FC region of immunoglobulin G, so the decreasing in SpA on cell surface of S. aureus resulted in increasing number of free receptor sites for complement and enhance phagocytosis [$^{\xi \circ}$].

In the ongoing audit, $\sqrt{1}(1\cdot\cdot\%)$ of attempted *s.aureus* withdraws clutched SpA quality and it upgraded at $\sqrt{1}$ bp that like the examination of [$\frac{1}{2}$ $\frac{1}{2}$] that showed strengthening of SpA nature of *S.aureus* at $\sqrt{1}$ $\frac{1}{2}$ bp.and El-Sayed [$\sqrt{1}$ $\frac{1}{2}$] who recognized SpA quality Positive in $\sqrt{1}$ $\frac{1}{2}$ $\frac{1}$

[\mathfrak{s}_{1}] reported that SpA gene can be used for typing the isolates of S.aureus. in addition to [\mathfrak{s}_{2}] point by point that the revelation of genetic polymorphisms in the X district of the SpA quality can be used for making out of S.

aureus. In like manner Karahan [5] gathered that ID of SpA quality polymorphisms with Coa-PCR is proposed as a fair illustrative method for making out of *S. aureus* limits which gave critical results to practical control of staphylococcal mastitis.

Practically like eventual outcomes of El-Sayed [Υ^{q}] who declared that Spa quality and Coa quality were perceived in $({}^{q}\cdot {}^{\prime})$ and $({}^{\wedge}\cdot {}^{\prime})$ of the attempted *S.aureus* strains

The resistance to β -lactam antibiotics occur through hydrolyzing the β -lactam ring and convert to inactive form, so the application of PCR for detection of *blaZ* gene is recommended in veterinary laboratories for detection of the resistant strains of *S. aureus* [£ $^{\lambda}$]. In this study, $^{\gamma}$ ($^{\gamma}$ ($^{\gamma}$, $^{\gamma}$ %) limits of *S. aureus* presented to PCR were positive for blaZ quality and gave a single amplicon at $^{\lambda}$ $^{\gamma}$ $^{\gamma}$ bp. At any rate special result got by [£ $^{\eta}$] and [£ $^{\gamma}$] recognized a singular amplicon of $^{\gamma}$ $^{\gamma}$ $^{\gamma}$ bp of blaZ quality. [$^{\circ}$ $^{\circ}$]

noticed that all positive strains to *mecA* were also positive for gene and the presence of both genes was correlated to phenotypic betalactam resistance of S. aureus strains.

In this study, $\ ^{\uparrow}\ ^{\uparrow}$ of tested *S. aureus* isolates harbored *blaz* gene which disagreed with the result of $[\ ^{\circ}\ ^{\uparrow}]$

who found that all S. aureus isolates harbored *blaZ* gene associated with penicillin

•.CONCLUSIONS

Staphylococcal food poisoning is of major concern in public health programs worldwide. S. aureus may be present in milk and milk products as a result of milk collected from the animal suffering from disease condition and excreting S. aureus in milk or due to unhygienic during production, conditions processing, storage and handling of milk products, which are the main causes of food borne diseases. Results clearly indicated that milk milkbased products available in the market were contaminated with S. aureus, posing a high risk of food poisoning. Thus more hygienic preventive measures are required to reduce the bacterial contamination, so as to increase the wholesomeness of these milk and milk based products.

Study concluded that S. aureus was isolated from milk and milk products, and raw milk was highly contaminated with the bacterium than fermented milk products (cheese and yoghurt). However, none of the dairy products was devoid of the isolates.

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Publication consent

Each author has demonstrated their consent for the publication of the current manuscript.

Data and material availability:

All data of this study is provided.

Conflict of interests.

All authors have stated the absence of any conflicts of interest.

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Authors' contributions.

K.M.S: Conceptualization, Formal Analysis, Investigation, Supervision, Resources, Writing – original draft

A.R.S: Data collection, Formal Analysis, Project administration, Resources, Writing – review and editing.

M.M.Z: Conceptualization, Data curation, Formal Analysis, Resources, Supervision, , Writing – review and editing.

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