

Prevalence and Antibiotic Resistance of *Staphylococcus aureus* Isolated From Raw Milk and Dairy Products Collected From Alexandria, Egypt

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ABSTRACT

The aim of the present study was to determine the prevalence and antibiotic susceptibility of *Staphylococcus aureus* (*S. aureus*) isolated from raw milk and some dairy products collected from shops, grocery stores, street vendors at different markets in Alexandria, Egypt and identification of the genes responsible for resistance to some important antibiotics using polymerase chain reaction (PCR) technique. The results of the raw milk and dairy products examined, 22.5% of the samples were positive for *S. aureus*. It was isolated from 29% of raw milk samples, 12.5% of Kariesh cheese samples, 55% of White cheese samples, 10% of Ras cheese samples, 5.7% of yoghurt samples and 24.3% of cream samples. Antimicrobial susceptibility test revealed that all the isolates were susceptible to ciprofloxacin, co-trimoxazole, linezolid, rifampin and vancomycin. Resistance to penicillin G, tetracycline, cefoxitin, chloramphenicol, erythromycin and gentamycin were detected in 91.4%, 65.4%, 42%, 13.6%, 12.3% and 9.9% of the isolates, respectively. The *mecA* resistance gene was identified in 91.2% of cefoxitin resistant *S. aureus* isolates, *aacA-D* was detected in 87.5% gentamycin resistant isolates and *ermC* gene was detected in 100% of *S. aureus* isolates showed macrolides resistance whereas *ermA* could not be detected in any of them. The overall prevalence of methicillin resistant *S. aureus* (MRSA) in the tested raw milk and dairy product samples was 8.6%. The highest methicillin resistant *S. aureus* prevalence was reported in white cheese samples of 22.5% followed by raw milk, Ras cheese, cream, yoghurt and Kariesh cheese with prevalence of 11%, 7.5%, 7.14%, 2.9% and 2.5%, respectively.

Key words: *Staphylococcus aureus*, dairy products, prevalence, antibiotic resistance.

INTRODUCTION

Staphylococcus aureus is an important pathogen for both human and animals (Fetsch & Jöhler, 2018). It is normal inhabitant of the skin, nasal cavity and mucus membranes of humans, and may colonize the skin and upper respiratory tract of other mammals. It can cause various pyogenic and systemic infections, acute and chronic infections, and toxin-mediated syndromes disease as toxic shock syndrome, food poisoning and staphylococcal scalded skin syndrome (Hu *et al.*, 2018). *S. aureus* is one of the main causes of bovine mastitis; it is responsible for approximately 30% to 40% of all mastitis cases which can easily contaminate raw milk (Gwida & El-Gohary, 2013).

S. aureus is a major cause of food poisoning worldwide and one of the main pathogens associated with the consumption of raw milk and dairy products. *S. aureus* can contaminate milk through infected producing animals or from human source

during milking and handling through hand or arm lesions caused by *S. aureus* or by coughing and sneezing, during respiratory infections (El-Malt *et al.*, 2013).

The misuse and overuse of antimicrobial agents in veterinary practice resulted in the development of resistant bacteria which may spread to humans through the food supply (Aydin *et al.*, 2011). *S. aureus* is capable of acquiring antibiotic resistance determinants and therefore, often show resistance to multiple classes of antibiotics (Papadopoulos *et al.*, 2018). *Staphylococcus* developed resistance to a wide range of antimicrobial drugs, since the introduction of penicillin in 1942 until now (Akpaka *et al.*, 2017). *S. aureus* strains have been reported for resistance against various antibiotic classes such as beta-lactams, tetracyclines, aminoglycosides, fluoroquinolones, lincosamides, macrolides and streptogramins that are widely used in veterinary medicine (Kumar *et al.*, 2010).

There is a rising awareness about the antibiotic resistant bacteria and their effect on public health, thus studying antibiotic resistance in food animal products is essential for detecting changing patterns of resistance, implementing control measures on the use of antibiotics to eliminate the spread of antibiotic resistant bacteria. The aim of this study was to determine the prevalence and antibiotic resistance of *S. aureus* isolated from raw milk, cheese, yoghurt and cream samples collected from dairy shops, grocery stores, street vendors at different markets in Alexandria, Egypt and identification of the genes responsible for resistance to some important antibiotics.

MATERIALS AND METHODS

Materials

A total of 360 samples consist of 100 raw milk (cow and buffalo), 120 cheese (40 of each Kariesh cheese, white cheese and Ras cheese), 70 yoghurt samples and 70 cream samples were collected during the period from January 2016 to March 2017 at consumer level from dairy shops, grocery stores, street vendors at different markets in Alexandria governorate, Egypt. The samples were collected and transported in ice box to the laboratory.

Methods

Twenty five ml of raw milk or twenty five gram of the other dairy products were added to 225 ml of the recommended diluent according to ISO 6887-5: 2010 method. Saline peptone water was used for raw milk, sodium citrate diluent was used for Kariesh cheese, white cheese, and Ras cheese samples, di-potassium hydrogen phosphate diluent was used for yoghurt samples. For cream samples, 25 g of

cream was added to 225 ml of saline peptone water and warmed to 45 °C until the sample was completely melted. Saline peptone water was used for preparing decimal dilutions for all the samples.

Isolation and identification of *S. aureus*

Isolation of *S. aureus* was carried according to ISO 6888-1:1999 by plating 0.1 ml of each of the initial suspension on the surface of Baird-Parker agar (Lab M, UK) supplemented with egg yolk tellurite emulsion (50ml /L), and incubated at 37 °C for 48 hr . Typical coagulase positive colonies are black or grey, shining and convex and surrounded by a clear zone. Two isolates with typical *S. aureus* morphology per sample were selected and confirmed by biochemical tests (rabbit plasma coagulation, identified as Gram-positive cocci, catalase positive, oxidase negative, DNase positive, negative for indol production, can ferment lactose, sucrose, glucose and mannitol.)

Antibiotic susceptibility test

The antimicrobial susceptibility phenotypes of *S. aureus* isolates were determined using Kirby-Bauer disk diffusion assay according to the standards and interpretive criteria described by Clinical and Laboratory Standards Institute (CLSI,2012-CLSI, 2016). Only one confirmed isolate per sample was tested. The results were recorded based on CLSI guidelines. *S. aureus* isolates were tested against cefoxitin (FOX,30µg) , chloramphenicol (C,30µg) , ciprofloxacin (CIP, 5µg), clindamycin (DA,2 µg) , co-trimoxazole (SXT, 1.25/ 23.75 µg) , erythromycin (E,15 µg) , gentamicin (CN, 10 µg) , linezolid (LZ, 30 µg) , penicillin G (P, 10 U), rifampicin(RA, 5µg) , tetracycline (TE, 30 µg) and vancomycin (VA, 30 µg).

Table 1: Primers used for antibiotic resistance genes amplification

Gene	Primer name	Primer sequence	Size (bp)	Annealing Temp. °C
<i>mecA</i>	<i>mecA</i> F	AAAATCGATGGTAAAGGTTGGC	532	55
	<i>mecA</i> R	AGTTCTGCAGTACCGGATTTGC		
<i>ermA</i>	<i>ermA</i> F	AAGCGGTAAACCCCTCTGA	190	55
	<i>ermA</i> R	TTCGCAAATCCCTTCTCAAC		
<i>ermC</i>	<i>ermC</i> F	AATCGTCAATTCCTGCATGT	299	55
	<i>ermC</i> R	TAATCGTGGAATACGGGTTTG		
<i>aacA-D</i>	<i>aacA-D</i> F	TAATCCAAGAGCAATAAGGGC	227	55
	<i>aacA-D</i> R	GCCACACTATCATAACCACTA		

Detection of antimicrobial resistance genes

The presence of genes associated with resistance to methicillin (methicillin resistance *mecA* gene), erythromycin and clindamycin (erythromycin ribosome methylases *ermA*, *ermC* genes), gentamicin (aminoglycoside acetyltransferases *aacA-D* gene) were detected by polymerase chain reaction (PCR) (Techne thermal cycler PHC-3, Cambridgem, UK) and the set of primers (Kumar *et al.*, 2010) used for each gene is shown in Table (1). Bacterial DNA was extracted using QIAamp® DNA Mini Kit (QIAGEN, Germany) according to the manufacturer's instructions and used as DNA template for (PCR).

RESULTS AND DISCUSSION

Prevalence of *S. aureus*

Table (2) represents the prevalence of *S. aureus* in raw milk and some dairy products collected from Alexandria. Out of the 360 examined samples, 22.5% of samples were positive for *S. aureus*. White cheese samples had the highest *S. aureus* prevalence of 55% followed by the raw milk, cream, Kariesh cheese and Ras cheese samples with prevalence of 29%, 24.3 %, 12.5% and 10%, respectively. Whereas yoghurt samples had the lowest *S. aureus* prevalence of 5.7%.

Agban & Ahmed (2013) detected *S. aureus* in 28% of raw milk samples collected from Assiut governorate's rural areas. Elbagory *et al.*(2015) and Fadel & Ismail (2015) reported higher prevalence of 60% and 40% for *S. aureus* in raw milk samples collected from El-Gharbia, and Ismailia governorates, respectively. Whereas, lower prevalence of 16.3% were reported by Mansour *et al.*(2017) for *S. aureus* in raw milk samples from Cairo, Giza, Kafr El-Sheikh governorates.

For Kariesh cheese, similar results were found by Salem *et al.* (2016) who reported a prevalence of 12.6 % for samples collected from Alexandria. However, higher prevalence was reported by Agben & Ahmed (2013), Elbagory *et al.* (2015), Fadel & Ismail (2015) who reported a prevalence of 37%, 52.5% and 63.3 % for Kariesh cheese samples collected from Assiut, El-Gharbia, and Ismailia governorates, respectively.

Atef *et al.* (2017) reported a similar prevalence of 5% for *S. aureus* in yoghurt samples collected from both El-Minya and Alexandria governorates.

On the other hand, higher prevalence of 42% were reported by El-Ansary *et al.* (2014) in yoghurt samples collected from El-Behera governorate. Also, Ahmed *et al.* (2014) reported a prevalence of 86.4 % in yoghurt samples collected from Cairo, Giza and Monfiya governorates. Whereas Sadek *et al.* (2014) could not detect *S. aureus* from yoghurt samples in Assiut governorate.

In contrary, higher prevalence of 75.5 % and 31.4 % for *S. aureus* were reported by Nazem *et al.*(2015) and Meshref (2013) for cream samples collected from Alexandria governorate and Beni-Suef city, respectively.

The relatively high prevalence of *S. aureus* in white cheese compared to the other products may be due to the use of unsuitable heat treated milk in the manufacturing of cheese and favorable conditions for bacterial growth during cheese making.

Table 2: Prevalence of *S. aureus* in raw milk and some dairy products collected from Alexandria governorate

Products	No of examined samples	positive samples	
		No	%
Raw milk	100	29	29
Kariesh cheese	40	5	12.5
White cheese	40	22	55.0
Ras cheese	40	4	10
Yoghurt	70	4	5.7
Cream	70	17	24.3
Overall	360	81	22.5

Antibiotic susceptibility of *S. aureus* isolates

The WHO developed a list of critically important antimicrobials to rank antimicrobials according to their importance in human medicine. It was created for use in developing risk management strategies to know which types of antimicrobials used in animals represent potentially higher risks to human populations. The list categorizes antimicrobials into three groups: critically important, highly important, and important (Collignon *et al.*, 2016).

Antibiotic susceptibility test of the *S. aureus* isolates revealed that all the isolates were susceptible to ciprofloxacin, co-trimoxazole, linezolid, rifampin and vancomycin. Thirty four (42%) of the isolates were resistant to cefoxitin and considered to be suspected methicillin resistant *S. aureus*. Resistance to penicillin G, tetracycline, cefoxitin,

chloramphenicol, erythromycin and gentamycin were detected in 92.6 %, 65.4 %, 42%, 13.6%, 12.3 % and 9.9 % of the isolates, respectively. Antibiotic resistance of *S. aureus* isolates are presented in Table (3)

Similar antibiotic resistance phenotypes were also reported in Egypt for *S. aureus* isolates from raw milk samples but with different percentages. Sadek *et al.* (2014) reported that *S. aureus* isolates from raw milk samples collected from Assiut governorate were sensitive to erythromycin, co-trimoxazole and ciprofloxacin and 7.6% and 17.4% of the isolates were resistance to tetracycline and gentamycin, respectively. Elmonier *et al.* (2018) found that 90.1%, 77.3%, 72.7 %, 22.7% and 9.1% of *S. aureus* isolates from raw milk samples collected from Kafr El-Sheikh governorate were resistance to tetracycline, erythromycin, chloramphenicol, gentamicin, and ceftiofur, respectively.

Many different results for *S. aureus* resistance phenotypes from raw milk and dairy products have been reported all over the world. In Ethiopia, 92.2 % and 66.7% of *S. aureus* isolates from milk samples were resistance to penicillin G and tetracycline, respectively (Mekuria *et al.*, 2013). In Iran, *S. aureus* isolates from raw cow milk, raw sheep milk, traditional cheese and kashk samples (prolonged boiled yoghurt) were sensitive to co-

trimoxazole and ciprofloxacin whereas, 56.1 %, 11.3 %, 7.9 % of the isolates were resistant to tetracycline, clindamycin and erythromycin, respectively (Jamali *et al.*, 2015). Moreover, *S. aureus* isolates from milk samples collected from small scale dairy farms in Penang, Malaysia were sensitive to chloramphenicol, ciprofloxacin and gentamycin, and 50%, 10% and 8.3% were resistance to tetracycline, clindamycin and erythromycin, respectively. Also, isolates showed lower resistance of 23.3% and 15.0% to penicillin G and ceftiofur, respectively (Shamila-Syuhada *et al.*, 2016). In north-western Greece, 99.3 % and 28.2% of the 212 *S. aureus* isolates from 367 samples of bulk milk tank, dairy products, nasal swabs, and swabs from dairy plant equipment surfaces, collected from two dairy plants and 24 dairy farms, were resistant to penicillin and tetracycline, respectively; and 3.8 % were suspected MRSA according to their antimicrobial resistance to oxacillin. Prevalence of MRSA was reported to be 10%, 2.4%, 6.5% and 7.3 % in bovine milk, farm workers, plant workers, and in plant equipment, respectively; and no MRSA was detected in any dairy product tested (Papadopoulos *et al.*, 2018)

Thirteen different resistance patterns were observed (Table 4). Resistance to penicillin G (P) alone was the highest reported pattern for resistance

Table 3: Antibiotic resistance of *S. aureus* isolates from raw milk and dairy products collected from Alexandria governorate

Antibiotic	Raw milk		Kariesh cheese		white cheese		Ras cheese		Yoghurt		Cream		Total isolates	
	N*= 29		N= 5		N= 22		N= 4		N= 4		N= 17		N= 81	
	No	%	No	%	No	%	No	%	No	%	No	%	No	%
Ceftiofur	12	41.4	1	20.0	9	40.9	3	75.0	2	50.0	7	41.8	34	42.0
Chloramphenicol	4	13.8	0	0	4	18.2	1	25.0	1	25.0	1	5.9	11	13.6
Ciprofloxacin	0	0	0	0	0	0	0	0	0	0	0	0	0	0
Clindamycin	6	20.7	0	0	2	9.1	2	50.0	0	0	0	0	10	12.3
Co-trimoxazole	0	0	0	0	0	0	0	0	0	0	0	0	0	0
Erythromycin	6	20.7	0	0	2	9.1	2	50.0	0	0	0	0	10	12.3
Gentamycin	2	6.9	1	20.0	2	9.1	2	50.0	0	0	1	5.9	8	9.9
Linezolid	0	0	0	0	0	0	0	0	0	0	0	0	0	0
Penicillin G	25	86.2	5	100	21	95.5	4	100	4	100	16	94.1	75	92.6
Rifampin	0	0	0	0	0	0	0	0	0	0	0	0	0	0
Tetracycline	18	62.1	4	80.0	15	68.2	1	25.0	3	75.0	12	70.6	53	65.4
Vancomycin	0	0	0	0	0	0	0	0	0	0	0	0	0	0

* N : number of isolates

to one antibiotic which was observed in 15 isolates. Resistance to penicillin G and tetracycline (P-TE) was the highest reported frequency in 24 isolates. Whereas resistance to penicillin, tetracycline and cefoxitin (P-TE-FOX) was the highest multi drug resistance (MDR) pattern which was reported in 10 *S. aureus* isolates Table (4).

Table (5) shows the distribution of *S. aureus* isolates according to their resistance to one, two or more antibiotics. It can be noted from the Table that 93.8% of the isolates were resistant to at least one antibiotic. Sixteen out of the 81 isolates (19.8%) were resistant to one antibiotic, Twenty eight out of the 81 isolates (34.86%) were resistant to two antibiotics, Thirty two out of the 81 isolates (39.5%) showed MDR. In accordance to these results, 34.8% of *S. aureus* isolates from raw milk

samples sold in some Assiut city markets showed MDR (Sadek *et al.*, 2014). Samaha *et al.* (2012) reported that 95.5% of *S. aureus* isolates from raw milk samples collected from Kafr El-Sheikh governorate showed multidrug resistant.

Antibiotic resistance genes

The 34 phenotypically suspected MRSA isolates (*S. aureus* showed resistance to cefoxitin) were confirmed by detecting the presence of *mecA* gene using PCR. Table (6) showed that 31 out of the 34 cefoxitin resistant *S. aureus* isolates were found to harbor the *mecA* gene, whereas it could not be detected in 3 isolates, one isolate from the raw milk samples and 2 isolates from cream samples. It can be noted from the Table that the highest MRSA prevalence was reported in white cheese

Table 4: Frequency of antibiotic resistance patterns of *S. aureus* isolates

Antibiotic resistance pattern	Raw milk	Kariesh cheese	White cheese	Ras cheese	Yoghurt	Cream	Total
TE	1	-	-	-	-	-	1
P	5	1	6	1	-	2	15
P-TE	8	3	5	-	2	6	24
P-FOX	1	-	-	-	1	2	4
P-TE-FOX	2	-	4	-	-	4	10
P-TE-C	-	-	1	-	-	1	2
P-TE-FOX-C	1	-	2	1	1	-	5
P-TE-FOX-CN	2	1	1	-	-	1	5
P-FOX-E/DA	2	-	-	-	-	-	2
P-FOX-E/DA-CN	-	-	-	2	-	-	2
P-TE-FOX-E/DA	1	-	-	-	-	-	1
P-TE-FOX-E/DA-C	3	-	1	-	-	-	4
P-TE-FOX-E/DA-CN	-	-	1	-	-	-	1

C: Chloramphenicol CN: Gentamicin DA: Clindamycin E: Erythromycin FOX: Cefoxitin P: Penicillin TE: Tetracycline

Table 5: Distribution of *S. aureus* isolates according to their resistance to one, two or more antibiotics

Product	No of isolates	Resistance to one antibiotic		Resistance to two antibiotics		Multidrug Resistance		Overall resistance	
		No	%	No	%	No	%	No	%
Raw milk	29	6	20.7	9	31.0	11	37.9	26	89.7
Kariesh cheese	5	1	20.0	3	60.0	1	20.0	5	100
white cheese	22	6	27.3	5	22.7	10	45.5	21	95.5
Ras cheese	4	1	25.0	0	0.0	3	75.0	4	100
Yoghurt	4	0	0.0	3	75.0	1	25.0	4	100
Cream	17	2	11.8	8	47.1	6	35.3	16	94.1
Total	81	16	19.8	28	34.6	32	39.5	76	93.8

samples of 22.5% followed by raw milk, Ras cheese, cream, yoghurt and Kariesh cheese samples with prevalence of 11%, 7.5% 7.14% 2.9% and 2.5%, respectively, and the overall prevalence of MRSA in the tested raw milk and dairy product samples was 8.6%.

Erythromycin resistant *S. aureus* isolates were tested for the presence of macrolide resistance genes (*ermA* and *ermC*). *ermC* gene was detected in all the 10 *S. aureus* isolates showed macrolides resistance whereas *ermA* could not detected in any of them. The eight *S. aureus* isolates which showed resistance to gentamycin were tested for the presence of *aacA-D* gene and seven of them were found to harbor the target gene.

In accordance to these results, a prevalence of 9.1 % for MRSA isolates identified by resistance to cefoxitin and PCR detection of *mecA* gene was reported for milk samples collected from Kafr El-Sheikh, Egypt (Elmonier *et al.*, 2018). Al-Ashmawy *et al.* (2014) found that all *S. aureus* isolates from milk and dairy products collected from Mansoura city, Egypt, were genetically verified as MRSA, harboring *mecA* gene. MRSA prevalence reported to be 75%, 65%, 40%, 50% and 35% in raw milk, Damietta cheese, Kariesh cheese, ice cream and yoghurt samples, respectively, and 53% of all milk and dairy products tested. Also, Fadel & Ismail (2015) reported prevalence of 33.3%, 20%, 13.3%, 40%, 13.3%, 33.3% and 13.3% for MRSA isolates with positive *mecA* gene for goat milk, sheep milk, buffalo milk, yoghurt, ice cream, Kariesh cheese samples and dairy workers swaps, respectively, collected from Ismailia city, Egypt.

Variable prevalence of MRSA in milk and dairy products in different countries has been reported. In central Italy, MRSA isolates which were positive to both *mecA* and *bla_Z* genes reported to be 1.3% of all the *S. aureus* isolates from different milk and dairy products. MRSA was found in pasta filata cheese and ovine bulk tank milk from two different farms (Carfora *et al.*, 2015). In Balikesir, Turkey, only three samples (2 from cow milk bulk tank and one from Tulum cheese) of 175 milk and dairy products tested were MRSA phenotypically being resistant to both oxacillin and cefoxitin, and only one was confirmed as MRSA carrying the *mecA* gene (Ektik *et al.*, 2018). In Faisalabad, Pakistan the prevalence of confirmed MRSA isolates by molecular methods targeting *mecA* was 34% in overall bovine milk samples (Aqib *et al.*, 2017). In Jordon, the prevalence of MRSA isolates confirmed by detecting *mecA* gene in bulk milk samples from cattle, sheep and goat milk was 20% 39.1% and 11.9%, respectively (Obaidat *et al.*, 2018).

CONCLUSION

A considerable portion of the examined samples was found to be contaminated with *S. aureus* which may constitute public health hazard and may lead to food poisoning. Moreover, the detection of antibiotic resistant *S. aureus* and the associated resistant genes in the examined samples indicates that raw milk, Kariesh cheese, white cheese, Ras cheese, yoghurt and cream may possess a public health hazard as there is strong evidence that human consumption of food contains antibiotic resistant bacteria resulted in acquisition of antibiotic resistant infections.

Table 6: Prevalence of confirmed methicillin-resistant *S. aureus* isolates in raw milk and dairy products collected from Alexandria governorate

Product	No. of examined samples	Suspected MRSA isolates *		Confirmed MRSA isolates **	
		No	%	No	%
Raw milk	100	12	12	11	11
Kariesh cheese	40	1	2.5	1	2.5
white cheese	40	9	22.5	9	22.5
Ras cheese	40	3	7.5	3	7.5
Yoghurt	70	2	2.9	2	2.9
Cream	70	7	10	5	7.14
Total	360	34	9.4	31	8.6

*Suspected MRSA isolates: cefoxitin- resistance isolates

**Confirmed MRSA isolates: *mecA* gene positive isolates

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انتشار المكورات العنقودية الذهبية المعزولة من اللبن الخام ومنتجات الألبان التي تم جمعها من محافظة الإسكندرية - مصر ومقاومتها للمضادات الحيوية

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الهدف من هذه الدراسة هو تحديد مدى انتشار المكورات العنقودية الذهبية في اللبن الخام وبعض منتجات الألبان التي تم جمعها من محافظة الإسكندرية ودراسة مقاومتها للمضادات الحيوية وتحديد الجينات المسؤولة عن مقاومة بعض المضادات الحيوية الهامة باستخدام تقنية تفاعل البلمرة المتسلسل. أظهرت النتائج ان ٢٢,٥٪ من أصل ٣٦٠ عينة من اللبن الخام ومنتجات الألبان التي تم فحصها كانت إيجابية للمكورات العنقودية الذهبية حيث تم عزلها من ٢٩٪ من عينات اللبن الخام، ١٢,٥٪ من عينات الجبن القريش، ٥٥٪ (٤٠/٢٢) من عينات الجبن الأبيض، ١٠٪ من عينات الجبن الرأس، ٥,٧٪ من عينات الزبادي و ٢٤,٣٪ من عينات القشدة. وكشف اختبار الحساسية للمضادات الحيوية أن جميع العزلات كانت حساسة للسيبروفلوكساسين، الكوتريموكسازول، اللينزوليد و الفانكوميسين. بينما تم اكتشاف مقاومة للبنسلين G، التتراسيكلين، السيفوكسيتين، الكلورامفينيكول، الإريثروميسين والجنتاميسين في ٩١,٤٪، ٦٥,٤٪، ٤٢٪، ١٣,٦٪، ١٢,٣٪ و ٩,٩٪ من العزلات، على الترتيب. تم تحديد جين المقاومة mecA في ٩١,٢٪ من عزلات المكورات العنقودية الذهبية المقاومة للسيفوكسيتين، تم الكشف عن aacA-D في عزلات مقاومة للجنتاميسين بنسبة ٨٧,٥٪، واكتشف جين ermC في ١٠٠٪ من العزلات التي أظهرت مقاومة للماكروليدات في حين لم يكن بالإمكان اكتشاف جين ermA في أي منهم. وكان معدل انتشار المكورات العنقودية الذهبية المقاومة للمضاد الحيوي ميثيسلين في عينات اللبن الخام ومنتجات الألبان التي تم دراستها هو ٨,٦٪. حيث كانت أعلى نسبة تواجد للمكورات العنقودية الذهبية المقاومة للمضاد الحيوي ميثيسلين في الجبن الأبيض بنسبة ٢٢,٥٪ يليها عينات اللبن الخام والجبن الرأس و الزبادي والجبن القريش وذلك بنسبه ١١٪، ٧,٥٪، ٧,١٤٪، ٢,٩٪ و ٢,٥٪ على الترتيب.

