

Research Paper

Enterotoxigenicity of *Staphylococcus aureus* isolated from traditional and commercial dairy products marketed in Iran

Ebrahim Rahimi

Department of Food Hygiene, College of Veterinary Medicine, Islamic Azad University, Shahrekord Branch, Shahrekord, Iran.

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Abstract

This study was conducted to determine the prevalence rate, enterotoxigenicity, and antimicrobial resistance of *S. aureus* isolated from dairy products in Iran. From September 2010 to July 2011, a total of 347 samples from various dairy products, traditional and commercial, were collected from randomly selected retail stores. Overall, 20 samples (5.8%) were found to be contaminated with *S. aureus*. The highest prevalence of *S. aureus* was found in traditional cheese (11.1%), followed by traditional ice-cream (5.9%), cream (5.6%), and butter (5.3%). The ability to synthesize classical staphylococcal enterotoxins (SEA-E) was determined in 7 of 20 (35%) isolates by using ELISA. SE type C was the most common enterotoxin found in the isolated *S. aureus* (42.9%), followed by SE type A (28.6%), SEA+SEC and SE type D (14.3%). Of the 20 isolates, 16 (80.0%) were positive for one or more enterotoxin genes and 8 different genotypes were observed. Susceptibilities of the isolates were determined for 14 antimicrobial drugs using the disk diffusion assay. Most of the isolates (95.0%) were resistant to one or more two antimicrobial agent and 45.0% of the isolates were resistant to three or more of drugs. Resistance to ampicillin was the most common finding (55.0%), followed by tetracycline (40.0%) and penicillin G (30.0%). The results of this study showed the wide spread of enterotoxigenic and multidrug-resistant *S. aureus* strains in traditional dairy products in Iran and highlighted their public health hazards.

Key words: antimicrobial resistance, dairy products, SE genes, staphylococcal enterotoxins, *Staphylococcus aureus*.

Introduction

Staphylococci are ubiquitous in nature; although the primary reservoir is on the skin and mucous membranes of mammals and birds. The presence of *Staphylococcus aureus* (*S. aureus*) in products for human consumption is important to the food industry as some strains are the cause of foodborne intoxication (Pelisser *et al.*, 2009). The genus *Staphylococci* includes over 30 species; whit 18 of these species and subspecies are of potential hazard in food poisoning as they produce either coagulase, heat stable nuclease or enterotoxins. The coagulase positive species are *S. aureus* subsp. *aureus* which is the most common enterotoxigenic species (Le Loir *et al.*, 2003, Loncarevic *et al.*, 2005). *S. aureus* is a common pathogen associated with se-

rious community and hospital acquired diseases and has for long been considered as a major problem of public health (Pesavento *et al.*, 2007).

The pathogenicity of *S. aureus* and its ability to cause diseases is attributed to a number of virulence factors such as the heat stable enterotoxins (Sandel and McKillip, 2004). *S. aureus* produces 15 enterotoxins (Atichou *et al.*, 2004). The five classical enterotoxins (SE, type A, B, C, D and E) were known to be responsible for 95% of staphylococcal food poisoning (SFP) cases, the rest of cases were due to the new types of SEs (SEG- SEO) (Jay *et al.*, 2005). However, the role of newly identified enterotoxins in food poisoning is not fully clarified, and the development of methods for the detection of these novel se genes is of critical importance for food poisoning investigations.

SFP is a mild intoxication occurring after the ingestion of food containing from 20 ng to 1 µg of staphylococcal enterotoxin (SE) which is enough to induce symptoms in human beings (Normanno *et al.*, 2007). SFP symptoms appear 1-6 h after ingestion of contaminated food, depending on individual susceptibility and toxic dose ingested. They include nausea, abdominal cramps, diarrhoea and a characteristic projectile vomiting. Clinical signs of SFP generally disappear within 24-48 h. Deaths occur rarely and specifically in the very young or elderly (Jay *et al.*, 2005).

Pasteurization kills *S. aureus* cells, and fermentation or ripening of cheese may prevent growth of *S. aureus* in raw milk cheese. However, once formed the thermostable SEs generally retain their biological activity (Becker *et al.*, 2007, Le Loir *et al.*, 2003).

Currently, there is limited information regarding the prevalence and antimicrobial susceptibility patterns of *S. aureus* in milk and dairy products in Iran. The aims of this study were to determine the prevalence rate, enterotoxigenicity, and antimicrobial resistance of *S. aureus* in commercial and traditional cheese, yoghurt, ice cream, butter, cream, kashk and dough in Iran. Although, governmental regulation of milk pasteurization and sanitation in dairy processing plants has been in existence in Iran for many years, direct sale of unpasteurized milk and dairy products from producers to the consumer is common in many regions including Isfahan, Chaharmahal va Bakhtyari, and Khuzestan provinces.

Materials and Methods

Sample collection

A total of 347 samples of various traditional and commercial dairy products including cheese (n = 90), ice-cream (n = 85), butter (n = 57), cream (n = 36), yoghurt (n = 30), kashk (n = 24), and Iranian dough (n = 25) from different brands were obtained from randomly selected retail stores located in 25 major cities of Isfahan, Chaharmahal va Bakhtyari, and Khuzestan provinces From September 2010 to July 2011. Kashk and doogh are two popular dairy products in Iran that are available both as traditional and commercial products. Kashk is prepared by prolonged boiling yogurt and doogh which is also called yogurt soda is prepared by beating unflavored yogurt until smooth, and then diluting with water to a consistency similar to whole milk. Traditional dairy products in Iran are produced in small productive centers mostly located in urban areas and distributed unpacked. These products may be produced from unpasteurized milk. All samples were immediately transferred to the food microbiology laboratory, Islamic Azad University of Shahrekord Branch, in portable insulated cold-boxes. The samples were analyzed on the day they were collected.

Isolation of *S. aureus*

A 10 g portion of each food product was added to 90 g of sterile phosphate buffered saline (137 mM NaCl, 10 mM phosphate, 2.7 mM KCl, pH is 7.4) and stomached for 30-90 s. Then microbiological processing of the samples for the isolation of *S. aureus* was performed using Baird-Parker agar (Difco) following a standard procedure. Decimal dilutions of sample suspensions were spread plated on the agar plates in duplicate following incubation for 24-48 h at 35 °C and then finally examined for colonies with typical appearance of *S. aureus*. To identify *S. aureus*, Gram stain, catalase, coagulase, and Voges-Proskaver (VP) tests were conducted on suspected colonies (Huong *et al.*, 2010).

Detection of classical staphylococcal enterotoxins (SEs)

To detect SEs, the isolates were cultured overnight aerobically in 10 mL nutrient broth (Merck, Germany) at 37 °C. Bacterial culture supernatants were collected by centrifugation at 4,000 x g for 10 min and used for detection of SEA, SEB, SEC, SED, and SEE using an enzyme linked immunosorbent assay (ELISA) detection kit (RIDASCREEN® SET A, B, C, D, E Art. No: R4101, R-Biopharm AG, Germany). The assay was performed according to the manufacturer's recommendation and as described elsewhere (Rahimi and Ghasemian Safai, 2010). The mean lower detection limit of the assay was 0.1 mg/mL. All experiments were performed in duplicate.

Detection of *sea*, *seb*, *sec*, *sed*, *see*, *seg*, *seh*, *sei*, and *sej*

Purification of DNA was achieved using a Genomic DNA purification kit (Fermentas, GmbH, Germany) according to the manufacturer's instruction and the total DNA was measured at 260 nm optical density according to the method described by Sambrook and Russell (2001). After DNA isolation, amplification of selected enterotoxin genes (*sea*, *seb*, *sec*, *sed*, *see*, *seg*, *seh*, *sei*, and *sej*) was achieved using 9 primer sets in the one reaction mixture. The sequences of the primers used for gene amplification are presented in Table 1. All oligonucleotide primers were obtained from a commercial source (Cinna Gen, Iran). Polymerase chain reaction (PCR) assays for the detection of *S. aureus* enterotoxin genes were performed according to the methods described previously (Rall *et al.*, 2008, Zouharova and Rysanek, 2008). Briefly, amplification reactions were performed in a 25 µL mixture containing 1 U Taq polymerase (Fermentas, GmbH, Germany), 200 µM of each dNTP (Fermentas, GmbH, Germany), 2.5 µL PCR buffer 10x (Fermentas, GmbH, Germany), 1.0 µM MgCl₂ (Fermentas, GmbH, Germany), 10 pmol of each primers, and 3 µL DNA. The final volume was adjusted to 25 µL by adding sterile ultrapure water (Rall *et al.*, 2008). Amplifica-

Table 1 - Primers and temperature used for the detection of *Staphylococcus aureus* SE genes.

Gene	Primer	Sequence	Base pair	Annealing temperature (°C)
<i>sea</i>	SEA-1	ttgaaacggttaaacgaa	120	50
	SEA-2	gaaccttccatcaaaaaca		
<i>seb</i>	SEB-1	tcgcatcaaaactgacaacg	478	50
	SEB-2	gcaggtactctataagtgcc		
<i>sec</i>	SEC-1	gacataaaagctaggaattt	257	50
	SEC-2	aaatcggattaacattatcc		
<i>sed</i>	SED-1	ctagtttgtaaatatctct	317	50
	SED-2	taatgctatatcttatagg		
<i>see</i>	SEE-1	aggtttttcacaggtcatcc	209	50
	SEE-2	cttttttctcgtcaatc		
<i>seg</i>	SEG-1	aagtagacattttggcggtcc	287	55
	SEG-2	agaacctcaaaactcgtatagc		
<i>seh</i>	SHE-1	gtctatatggaggtacaacact	213	46.4
	SHE-2	gaccttactatttcgctgtc		
<i>sei</i>	SIE-1	ggtgatattggttaggtaac	454	50
	SIE-2	atccatattctttgccttaccag		
<i>sej</i>	SEJ-1	catcagaactgtgttccgctag	142	50
	SEJ-2	ctgaattttaccatcaagggtac		

tion reactions were carried out using a DNA thermal cycler (Master Cycle Gradient, Eppendorf, Germany) with the following program: one cycle of 5 min at 94 °C, 30 cycles each consisting of 45 s at 94 °C, 1 min and 30 s at 65 °C, 1 min at 72 °C and a final extension step of 8 min at 72 °C. The PCR products were stained with 1% solution of ethidium bromide and visualized under UV light after gel electrophoresis on 1.5% agarose. *S. aureus* ATCC 19095, ATCC 23235 and ATCC 700699 were used as the positive controls and DNase free water was used as the negative control.

Antimicrobial susceptibility testing

One colonia from each *S. aureus*-positive sample was selected for susceptibility tests. Antimicrobial susceptibility testing was performed by the Kirby-Bauer disc diffusion method using Mueller-Hinton agar (HiMedia Laboratories, Mumbai, India) supplemented with 5% defibrinated sheep blood, according to the Clinical Laboratory Standards Institute (2006). The following antimicrobial impregnated disks (HiMedia Laboratories, Mumbai, India) were used: penicillin G (10 IU), cephalotin (30 µg), chloramphenicol (30 µg), clindamycin (15 µg), ciprofloxacin (30 µg), erythromycin (15 µg), tetracycline (30 µg), oxacillin (15 µg), gentamycin (10 µg), ampicillin (10 µg), enrofloxacin (10 µg), trimethoprim-sulfamethoxazole (25 µg), methicillin (5µg), and vancomycin (30 µg). After incubation at 37 °C for 48 h, the susceptibility of the *S. aureus* isolates to each antimicrobial agent was measured and the results were interpreted in accordance with interpretive criteria provided by CLSI (2006).

Statistical analysis

Data were transferred to Microsoft Excel spreadsheet (Microsoft Corp., Redmond, WA, USA) for analysis. Using SPSS 16.0 statistical software (SPSS Inc., Chicago, IL, USA), chi-square test and fisher's exact two-tailed test analysis was performed and differences were considered significant at values of $p < 0.05$.

Results

In the present study, a total of 347 samples of various commercial and traditional milk and dairy products were tested for enterotoxigenic *S. aureus* (Table 2). Using cultural techniques, 20 of 347 samples (5.8%) were positive for *S. aureus*. No *S. aureus* was isolated from commercial cheese, ice cream, butter and cream samples and all yogurt, doogh, and kashk samples were negative for *S. aureus*. In contrast, 10 cheese (11.1%), 5 ice cream (5.9%), 2 cream (5.6%) and 3 butter (5.3%) samples were contaminated with *S. aureus*. In this study, total *S. aureus* counts determined were between 1×10^2 and 1×10^6 cfu/g in samples. In 3 (6.0%) of traditional cheese, and 1 (3.7%) butter samples, CPS counts were above 10^5 cfu/g (Table 2).

The ability to synthesize classical enterotoxins was found in 7 of 20 (35.0%) isolates by using ELISA technique. Two isolates produced SEA (28.6%), 3 isolates produced SEC (42.9%), 1 isolates produced SEA+SEC (14.3%), and 1 isolates produced SEA + SED (14.3%) (Table 2). No SEB and SEE were identified in dairy product samples. Type A enterotoxin was found in 1 butter and 1 ice-cream samples. Type C and type A+C enterotoxin was

Table 2 - Occurrence and enterotoxigenicity of *Staphylococcus aureus* from traditional dairy products in Fars Iran.

Sources	No. of samples	<i>S. aureus</i> positive samples			No. of SES	SEs			
		Number (%)	Distribution (CFU/G)			A	C	A+C	A+D
			1x10 ² -1x10 ⁴	1x10 ⁴ -1x10 ⁶					
Cheese	90	10 (11.1)	7	3	4	-	3	1	-
Commercial cheese	30	-	-	-	-	-	-	-	-
Traditional cheese ^a	60	10 (16.7)	7	3	4	-	3	1	-
Ice-cream	85	5 (5.9)	5	-	1	1	-	-	-
Commercial ice-cream	30	-	-	-	-	-	-	-	-
Traditional ice-cream	55	5 (9.1)	5	-	1	1	-	-	-
Butter	57	3 (5.3)	2	1	2	1	-	-	1
Commercial butter	20	-	-	-	-	-	-	-	-
Traditional butter	37	3 (8.1)	2	1	2	1	-	-	1
Cream	36	2 (5.6)	2	-	-	-	-	-	-
Commercial cream	10	-	-	-	-	-	-	-	-
Traditional cream	26	2 (7.7)	2	-	-	-	-	-	-
Yoghurt	30	-	-	-	-	-	-	-	-
Commercial yoghurt	11	-	-	-	-	-	-	-	-
Traditional yoghurt	19	-	-	-	-	-	-	-	-
Kashk ^b	24	-	-	-	-	-	-	-	-
Commercial kashk	8	-	-	-	-	-	-	-	-
Traditional kashk	16	-	-	-	-	-	-	-	-
Iranian dough ^c	25	-	-	-	-	-	-	-	-
Commercial dough	10	-	-	-	-	-	-	-	-
Traditional dough	15	-	-	-	-	-	-	-	-
Total	347	20 (5.8)	16 (80)	4 (20)	7 (35)	2	3	1	1

^aMade from raw sheep or cow milk. ^bA dairy product prepared by beating unflavored yogurt until smooth, and then diluting with water to a consistency similar to whole milk; it is also called yogurt soda. ^cA dairy product prepared by prolonged boiling yogurt.

detected in only 4 traditional cheeses. Type A + D enterotoxin was found in 1 butter (Table 2). No SEs was identified in cream samples.

Table 3 shows the results of PCR assays for the detection of genes encoding the SEA-SEJ toxins. Of the 20 strains of *S. aureus* tested, 16 (80.0%) were positive for one or more SE genes and 9 different genotypes were observed.

Five strains possessed only one type of toxin gene and the remaining isolates harbored more than one toxin genes. The most commonly detected genes were *seg* + *sei* (20.0%), *sej* (15.0%), *sec* + *sej* (15.0%), *sei* + *seh* (10.0%), *sea* + *sed* + *sej* (5.0%), *sea* + *sej* + *sei* (5.0%), and *sea* + *sec* + *seh* (5.0%). No *see* or *seb* gene was detected in any of the isolates.

Table 3 - Detection rates of genotypes of *Staphylococcus aureus* isolated from traditional dairy products according to *sea*, *seb*, *sed*, *see*, *seg*, *she*, *sei* and *sej* genes.

Genotypes	Cheese (N = 10)	Ice-cream (N = 5)	Butter (N = 3)	Cream (N = 2)	Total (N = 20)
Sea	-	-	1	-	1
sej	2	1	-	-	3
Sec, sej	3	-	-	-	3
Seg, sei	2	-	1	1	4
Sei, seh	-	1	-	1	2
Sea, seg, sei	-	1	-	-	1
Sea, sed, sej	-	-	1	-	1
Sea, sec seg, seh	1	-	-	-	1

On comparing the data relative to the strains isolated from chesses, ice cream, butter, and cream, 8 of the 10 (80.0%) strains isolated from cheese samples were *se* gene positive in which the *sec* genes were found more frequently. Three of the 5 (60.0%) *S. aureus* isolated from ice-cream and all of the *S. aureus* isolated from butter and cream samples were *se* gene positive.

The resistance pattern of *S. aureus* isolates to 14 antimicrobial agents tested in this study is shown in Table 4. Most of the isolates (95.0%; n = 19) were resistant to one or more antimicrobial agent. Three isolates (15.0%) were resistant to single antibiotic and 7 isolates (35.0%) showed resistance to 2 antimicrobial agents. Multiresistance which was defined as resistance to 3 or more of antimicrobial agents tested was found in 45.0% of *S. aureus* isolates.

Results and Discussion

In the present study, 20 of the 220 traditional (9.1%) dairy product samples were positive for *S. aureus*. No *S. aureus* was isolated from 127 commercial dairy products and the difference was statistically significant ($p < 0.05$). The high occurrence of *S. aureus* in traditional dairy products could be due to environmental contamination with infected animal wastes or unsanitary food production and storage practices. However, this could be also due to the use of unpasteurized milk because the shedding of bacteria from the infected mammary glands of dairy animals is most likely the primary source of *S. aureus* contamination of

milk and dairy products. While commercial products are produced with pasteurized milk under sanitary condition.

S. aureus count was determined about 10^3 cfu/g in 20 samples period. In 3 of sheep cheese, and 1 butter samples, *S. aureus* counts were above 10^5 cfu/g (Table 2). The prevalence of *S. aureus* in traditional cheese, ice cream, cream and butter was 11.1%, 5.9%, 5.6%, and 5.3%, respectively, which is comparable with those reported by Manfreda *et al.* (2005), Ertas *et al.* (2010), Bostan *et al.* (2006), and Pelisser *et al.* (2009). Necidová *et al.* (2009) reported, *S. aureus* count higher than 10^5 cfu/g is unsuitable for the production of cheese. Staphylococcal counts should reach approximately 10^6 cfu/g to produce enterotoxin (Necidová *et al.*, 2009, Pelisser *et al.*, 2009).

In this study, 35.0% of isolated *S. aureus* strains produced classical enterotoxins. This result is in agreement with those reported by other investigators (Morandi *et al.*, 2007, Normanno *et al.*, 2005). Most of the isolated strains produced SEA, SED, and SEC. Our data show that SEs and *se* genes are in close correlation with the *S. aureus* strain origin. For example, a higher ratio of strains isolated from traditional sheep and goat cheeses produced mainly SEC, while the strains isolated from other dairy products made from cow milk produced SEA and SED. These results are in agreement with the data reported from other countries (Akineden *et al.*, 2008, Loncarevic *et al.*, 2005, Morandi *et al.*, 2005, Rall *et al.*, 2008, Scherrer *et al.*, 2004). Of the 20 strains of *S. aureus* tested, 17 (85.0%) were positive for one or more SE genes and 8 different genotypes were observed. Five strains possessed one type of toxin gene, while the remaining 12 harbored more than one toxin genes. The most commonly detected genes were *seg + sei*, *sej*, *sec + sej*, *sei + seh*, *sea + sed + sej*, *sea + sej + sei*, and *sea + sec + seh*. Similar results were presented by Normanno *et al.* (2005), Manfreda *et al.* (2005), and Tkacikova *et al.* (2003). In contrast, the incidence of *S. aureus* and their SEs was reported relatively high by several authors (Fotta *et al.*, 2000, Holeckova *et al.*, 2002) when compared with our results. Variation in the results reported in other studies may be a result of different sampling techniques employed, seasonal effects, and/or laboratory methodologies employed in different studies.

S. aureus strains are known to be frequently resistant to antibiotic therapy due to their capacity to produce an exopolysaccharide barrier and because of their location within microabscesses, which limit the action of drugs (Gündogan *et al.*, 2006). The resistance pattern of *S. aureus* isolates to 12 antimicrobial agents tested in this study is shown in Table 3. Most of the 20 *S. aureus* isolates (95.0%) were resistant to one or more antimicrobial agent. Resistance to ampicillin was the most common finding, followed by resistance to tetracycline, penicillin G, oxacillin, erythromycin, trimethoprim-sulfamethoxazole, enrofloxacin, gentamicin, chloramphenicol, clindamycin, and ciprofloxacin. All isolates tested for antibiotic sensitivity were susceptible

Table 4 - Antimicrobial resistance profiles of *Staphylococcus aureus* isolated from dairy product samples in Iran.

Antimicrobial agent	<i>S. aureus</i> (N = 20)
Ampicillin	11 (55.0%)
Cephalotin	0 (0.0%)
Chloramphenicol	2 (10.0%)
Ciprofloxacin	1 (5.0%)
Clindamycin	1 (5.0%)
Enrofloxacin	3 (15.0%)
Erythromycin	5 (25.0%)
Gentamicin	3 (15.0%)
Methicillin	0 (0.0%)
Oxacillin	5 (25.0%)
Penicillin G	6 (30.0%)
Tetracycline	8 (40.0%)
Trimethoprim-sulfamethoxazole	5 (25.0%)
Vancomycin	0 (0.0%)
Total	
Resistance to 1 antimicrobial	3 (15.0%)
Resistance to 2 antimicrobials	7 (35.0%)
Resistance to > 2 antimicrobials	9 (45.0%)

to methicillin cephalotin and vancomycin. These results are comparable to those reported by other investigators (Gündogan *et al.*, 2006, Normanno *et al.*, 2007, Peles *et al.*, 2007, Pereira *et al.*, 2009, Pesavento *et al.*, 2007). The results of antimicrobial resistance found in this study are correlated to antibiotics that are being used to treat infection in food animals in Iran. A national monitoring program for antibiotic resistance, including both human and food isolates, is needed in Iran.

National legislation of *S. aureus* in milk and dairy products which is 10^2 cfu/g. *S. aureus* in about 9% of traditional dairy products found to be above which shows the importance of some traditional dairy products as potential sources of *S. aureus* infection in people who consume traditional dairy products. We recommend that coordinated actions are needed to reduce or eliminate the risks posed by this organism at a number of stages in the food chain. These include Good Agricultural Practice, Good Manufacturing Practice and Hazard Analysis of Critical Control Points at every stages of the meat supply chain, from the farm, to the retailer, and those involved with the handling and processing of raw milk products in the home environment.

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