Antimicrobial resistance of *Staphylococcus* spp. from small ruminant mastitis in Brazil¹

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Abstract.- França C.A., Peixoto R.M., Cavalcante M.B., Melo N.F., Oliveira C.J.B., Veschi J.L.A., Mota R.A. & Costa M.M. 2012. **Antimicrobial resistance of** *Staphylococcus* **spp. from small ruminant mastitis in Brazil.** *Pesquisa Veterinária Brasileira 32(8):747-753*. Projeto de Irrigação Senador Nilo Coelho s/n, Universidade Federal do Vale do São Francisco, Campus Ciências Agrárias, Rodovia BR 407 Km 12, Lote 543, Petrolina, PE 56300-990, Brazil. E-mail: mateus.costa@univasf.edu.br

The study aimed to determine the antimicrobial resistance patterns and to identify molecular resistance markers in *Staphylococcus* spp. (n=210) isolated from small ruminant mastitis in Brazil. The antimicrobial resistance patterns were evaluated by the disk diffusion test and by detection of the presence of mecA, blaZ, ermA, ermB, ermC and *msr*A genes by PCR. The efflux pump test was performed using ethidium bromide and biofilm production was determined by Congo red agar test along with PCR for detection of the *icaD* gene. The isolates were most resistant to amoxicillin (50.0%), streptomycin (42.8%), tetracycline (40.4%), lincomycin (39.0%) and erythromycin (33.8%). Pan-susceptibility to all tested drugs was observed in 71 (33.8%) isolates and 41 Staphylococcus isolates were positive for the efflux pump. Although phenotypic resistance to oxacillin was observed in 12.8% of the isolates, none harbored the *mecA* gene. However, 45.7% of the isolates harbored *blaZ* indicating that beta-lactamase production was the main mechanism associated with staphylococci resistance to beta-lactams in the present study. The other determinants of resistance to antimicrobial agents *ermA*. *ermB*. *ermC*, and msrA were observed in 1.4%, 10.4%, 16.2%, and 0.9% of the isolates, respectively. In addition, the icaD gen was detected in 32.9% of the isolates. Seventy three isolates (54 from goats and 19 from sheep) were negative for all resistance genes tested and 69 isolates presented two or more resistance genes. Association among blaZ, ermA, ermB, ermC and efflux pump were observed in 17 isolates, 14 of which originated from goats and three from sheep. The data obtained in this study show the resistance of the isolates to beta-lactamics, which may be associated with the use of antimicrobial drugs without veterinary control.

INDEX TERMS: Biofilm, efflux pump, goats, milk, sheep.

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⁵ Centro de Ciências Agrárias, Universidade Federal da Paraíba, Rod. PB 079 Km 12, Zona Rural, Areia, PB 58397-000, Brazil. **RESUMO.-** [**Resistência antimicrobiana de** *Staphylococcus* **spp. de mastite de pequenos ruminantes no Brasil.**] O presente trabalho teve como objetivo determinar os padrões de resistência a agentes antimicrobianos e identificar marcadores moleculares de resistência em *Staphylococcus* spp. (n=210) isolados de mastite de pequenos ruminantes no Brasil. Os padrões de resistência a agentes antimicrobianos foram avaliados pelo teste de difusão em disco e pela detecção da presença dos genes *mecA, blaZ, ermA, ermB, ermC* e *msrA* via PCR. O teste da bomba de efluxo foi realizado utilizando brometo de etídio e a produção de biofime foi determinada pelo teste do vermelho congo em paralelo com o PCR para detecção do gene *ica*D.

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Os isolados foram mais resistentes a amoxicilina (50,0%), estreptomicina (42,8%), tetraciclina (40,4%), lincomicina (39,0%) e eritromicina (33,8%). Setenta e um (33,8%) isolados foram sensíveis a todas as drogas testadas e 41 foram positivos para a bomba de efluxo. Embora a resistência fenotípica a oxacilina tenha sido observada observada em 12,8% dos isolados, nenhum possuiu o gene mecA. Entretanto, 45,7% dos isolados continham a gene blaZ, indicando que a produção de beta-lactamases foi o principal mecanismo associado com a resistência dos Staphylococcus aos beta-lactâmicos. Os outros determinantes de resistência a agentes antimicrobianos ermA, ermb, ermC e msrA foram observados em 1,4%, 10,4%, 16,2% e 0,9% dos isolados respectivamente. Além disso, o gene icaD foi detectado em 32,9% dos isolados. Setenta e três isolados (54 de cabras e 19 de ovelhas) foram negativos para todos os genes de resistência testados e 69 isolados apresentaram dois ou mais genes de resistência. A associação entre blaZ, ermA, ermB, ermC e bomba de efluxo foi observada em 17 isolados dos quais 14 eram oriundos de cabras e três de ovelhas. Os dados obtidos no presente estudo indicam a resistência dos isolados aos beta-lactâmicos, o que pode estar associado ao uso sem controle veterinário destas drogas nos animais.

TERMOS DE INDEXAÇÃO: Biofilme, bomba de efluxo, cabra, leite, ovelha.

INTRODUCTION

Mastitis is a serious concern to both meat and milk producers since the infection can lead to considerable economic losses due to reduction in milk yield, decreased milk quality and treatment costs (Sawant et al. 2009). Staphylococci are considered to be the most important mastitis causing agents in ruminants (Aires-de-Sousa et al. 2007, Taponen & Pyörälä 2009) and antimicrobial therapy is usually required in treatment (Bergonier et al. 2000, Contreras et al. 2007, Pÿorälä & Taponen 2009). However, there is evidence that resistance to antibiotics is increasing in staphylococci infected food animals (Phillips et al. 2004). The emergence of multidrug resistance (MDR) in *Staphylococcus* spp. is a problem for concern in animal production and issues related to public health (Kumar et al. 2009, Sawant et al. 2009, Virdis et al. 2010).

More than 90% of the Brazilian goat and sheep population is concentrated in the Northeast region, which makes it the leading goat milk supplier in the country. However, the production system is predominantly extensive or semi--intensive and mostly comprised of small scale producers with major technical limitations. Although staphylococci species have been identified as the most common mastitiscausing agent in small ruminant species in Brazil (Silva et al. 2004), knowledge about antimicrobial resistance mechanisms in those bacteria is still lacking.

Therefore, the aim of this study was to determine the antimicrobial resistance patterns and to characterize the antimicrobial resistance mechanisms of *Staphylococcus* species isolated from subclinical mastitis in goats and sheep in Northeastern Brazil.

MATERIALS AND METHODS

Staphylococci isolates

The investigation included Staphylococci (n=210) from subclinical mastitis cases in goats (n=171) and sheep (n=39) raised in 25 herds (17 goat herds, 6 sheep herds and 2 mixed herds) from the states of Pernambuco and Bahia in Northeastern Brazil. The isolated *Staphylococcus* spp. were identified by morphological and biochemical tests according to Quinn et al. (1994).

The bacteria were deposited in the bacteria bank of the Animal Microbiology and Immunology Laboratory of the Universidade Federal do Vale do São Francisco (Federal University of São Francisco Valley) (Univasf) and at the Bacteriosis Laboratory of the School of Veterinary Medicine, Universidade Federal Rural de Pernambuco (Federal Rural University of Pernambuco) (UFR-PE) and included 37 *S. epidermidis* (17.6%), 35 *S. intermedius* (16.6%), 35 *S. hyicus* (16.6%), 34 *S. aureus* (16.2%), 23 *S. caprae* (11.0%), five *S. gallinarum* (2.3%), three *S. simulans* (1.4%) and one *S. saprophyticus* (0.5%). Thirty seven isolates (17.6%) were identified as coagulase negative *Staphylococcus* (CNS) since the species could not be identified.

Antimicrobial susceptibility test

Antimicrobial resistance patterns were determined by the Kirby-Bauer disk diffusion test performed according to recommendations (CLSI 2006). The drugs tested and their concentrations were: gentamycin (10µg), streptomycin (10µg), enrofloxacin (05µg), norfloxacin (10µg), ciprofloxacin (05µg), amoxicillin (10µg), oxacillin (01µg), doxycycline (30µg), tetracycline (30µg), erythromycin (15µg), lincomycin (02µg) and rifampicin (05µg). As a control for antimicrobial testing, we used the reference strains *Staphylococcus aureus* ATCC 25923 and *Escherichia coli* ATCC 49247. The multidrug resistance rate (MDR rate) was calculated by dividing the number of antimicrobial resistant groups (Krumperman et al. 1983).

Efflux pump mechanism detection

Efflux pump screening in *Staphylococcus* spp. was carried out as previously described in Bjorland et al. (2005). Thus, isolates were grown on Mueller Hinton Agar (MH) containing ethidium bromide (0.5mg/ml) for 24 hours at 37°C. Isolates were considered positive for efflux mediated resistance when no fluorescence under UV examination was observed.

Screening for biofilm production

Biofilm production was assessed by inoculating isolates onto Congo Red Agar, according to Greco et al. (2008). After incubation at 37°C for 24h, black colonies were considered as biofilm producers.

Polymerase chain reaction (PCR)

Antimicrobial resistance genes (*mecA*, *blaZ*, *ermA*, B, C and *msrA*) and the biofilm producing gene *icaD* were identified by PCR using the primers and cycle conditions described by Greco et al. (2008), Murakami et al. (1991) and Sawant et al. (2009) (Table 1). Shortly, bacterial genomic DNA was obtained by thermal extraction (Greco et al. 2008). DNA template (2µl) was added to a 23µl master mix containing 4pmol of each primer, 0.4mM of dNTPs mix, 1U of *Taq* DNA polymerase (Ludwig- Biotec), buffer (10mM Tris-HCl [ph 8.5] and 50mM KCl) and 2 mM MgCl₂. PCR products were electrophoresed in 1.5% agarose gel stained with ethidium bromide (0.5μ g/mL) and observed under UV illumination. DNA from reference isolates harboring the target genes were used as positive controls.

Table 1. Oligonucleotide primers, amplification conditions (temperature/ denaturation time/annealing/
extension) and amplicon sizes for the polymerase chain reactions

		, i			
Genes	Primers	Oligonucleotide sequences	Amplification conditions	Amplicon size (pb)	Reference
icaD SeicaDFw		AAGCCCAGACAGAGGCAATATCCA	30s - 94°C	249	Greco et al. 2008
			30s – 53.5°C		
	SeicaDRev	AGTACAAACAAACTCATCCATCCGA	30s - 72°C		
mecA	SA-1	CGGTAACATTGATCGCAACGTTCA	30s - 94°C	214	Murakami et al. 199
			68°C - 15 cycles		
	SA-2	CTTTGGAACGATGCCTAATCTCAT	60°C - 20 cycles		
			30s - 72°C		
blaZ	blaZ F	AAGAGATTTGCCTATGCTTC	30s - 94°C	517	Sawant et al. 2009
			30s - 50°C		
	blaZ R	GCTTGACCACTTTTATCAGC	30s - 72°C		
ermA	ermA F	ATCGGATCAGGAAAAGGACA	1 min - 94°C	486	Sawant et al. 2009
			30s - 49°C		
	ermA R	CACGATATTCACGTTTTACCC	30s - 72°C		
ermB	ermB F	AAGGGCATTTAACGACGAAA	1min - 94°C	423	Sawant et al., 2009
			30s – 49.5°C		
	ermB R	CTGTGGTATGGCGGGTAAGT	30s - 72°C		
ermC	ermC F	TGAAATCGGCTCAGGAAAAG	1min - 94°C	272	Sawant et al. 2009
			30s - 52°C		
	ermC R	CAAACCCGTATTCCACGATT	30s - 72°C		
msrA	<i>msr</i> A F	TGGTACTGGCAAAACCACAT	30s - 94°C	1000	Sawant et al. 2009
			30s - 52°C		
	<i>msr</i> A R	AAACGTCACGCATGTCTTCA	30s - 72°C		

 Table 2. Antimicrobial resistance rates (%) of Staphylococcus spp. causing mastitis in goat and sheep in Brazil determined by the disk diffusion test

	Staphylococci species												
	S. aureus	S. caprae	S. epidermi-	S. gallinarum	S. hyicus	S. interme-	S. saprophyti-	S. simulans	CNS	Total			
	(n=34)	(n=23)	<i>dis</i> (n=37)	(n=5)	(n=35)	<i>dius</i> (n=35)	<i>cus</i> (n=1)	(n=3)	(n=37)	(n=210)			
Gentamycin (5µg)	0 (0.0%)	1 (4.3%)	3 (8.11%)	0 (0.0%)	3(8.5%)	4 (11.4%)	0 (0.0%)	0 (0.0%)	5 (13.5%)	16			
Streptomycin (10µg)	10 (29.4%)	10 (43.4%)	18 (48.6%)	1 (20%)	13(37.1%)	13 (37.1%)	1(100%)	0 (0.0 %)	24 (64.8%)	91			
Enrofloxacin (5µg)	6 (17.6%)	1 (4.3%)	4 (10.8%)	0 (0.0%)	3 (8.5%)	2 (5.7%)	0 (0.0%)	0 (0.0%)	1 (2.7%)	17			
Norfloxacin (10µg)	10 (29.4%)	1 (4.3%)	6 (16.2%)	1 (20%)	7 (20%)	3 (8.5%)	1 (100%)	0 (0.0%)	0 (0.0%)	29			
Ciprofloxacin (30µg)	5 (14.7%)	3 (13.0%)	1 (2.7%)	2 (40%)	6 (17.1%)	0 (0.0%)	0 (0.0%)	0 (0.0%)	2 (5.4%)	19			
Amoxicillin (30µg)	12 (35.3%)	17 (74.0%)	24 (64.8%)	0 (0.0%)	12 (34.3%)	18 (51.4%)	1 (100%)	0 (0.0%)	21 (56.7%)	105			
Oxacillin (1µg)	1 (2.9%)	3 (13.0%)	6 (16.2%)	0 (0.0%)	9 (25.7%)	6 (17.1%)	1 (100%)	0 (0.0%)	9 (24.3%)	35			
Doxycycline (30µg)	4 (11.7%)	3 (13.0%)	6 (16.2%)	0 (0.0%)	4 (11.4%)	4 (11.4%)	0 (0.0%)	0 (0.0%)	4 (10.8%)	25			
Tetracycline (30µg)	5(14.7%)	11 (47.8%)	18 (48.6%)	0 (0.0%)	14 (40.0%)	18 (51.4%)	0 (0.0%)	0 (0.0%)	19 (51.3%)	85			
Erythromycin (10µg)	4 (11.7%)	10 (43.4%)	16 (43.2%)	0 (0.0%)	12 (34.3%)	9 (25.7%)	0 (0.0%)	0 (0.0%)	20 (54.0%)	71			
Lincomycin (2µg)	4 (11.7%)	11 (47.8%)	19 (51.3%)	1 (20.%)	11 (31.4%)	15 (42.8%)	0 (0.0%)	0 (0.0%)	21 (56.7%)	82			
Rifampicin (30µg)	13 (38.2%)	1 (4.3%)	5 (13.5%)	0 (0.0%)	6 (17.1%)	4 (11.4%)	1 (100%)	0 (0.0%)	5 (13.5%)	35			

RESULTS

The antimicrobial resistance patterns of the Staphylococcus spp. isolates are shown in Table 2. The isolates were most resistant to amoxicillin (50.0%), followed by streptomycin (43.4%), tetracycline (40.5%), lincomycin (39.0%) erythromycin (33.8%), rifampicin (16.7%), oxacillin (16.7%), norfloxacin (13.8%), doxycicline (12.0%), ciprofloxacin (8.5%), enrofloxacin (8.1%) and gentamycin (7.62%). Considering the resistance patterns of staphylococci according to the host species (Fig.1), higher resistances were observed in the goat isolates as compared to sheep isolates for the majority of the antimicrobials tested, such as amoxicillin (57.3% vs. 18.0%), streptomycin (46.8% vs. 10.3%), tetracycline (45.6% vs. 10.3%), lincomycin (45.6% vs. 7.7%) erythromycin (35.1% vs. 23.1%), rifampicin (25.7% vs. 23.1%), oxacillin (15.8% vs. 12.8%), norfloxacin (10.5% vs. 5.1%), doxycicline (10.5% vs 7.7%) and enrofloxacin (2.9% vs. 0%). Lower resistance rates in goat isolates were only observed for ciprofloxacin (5.9% vs. 10.3%) and

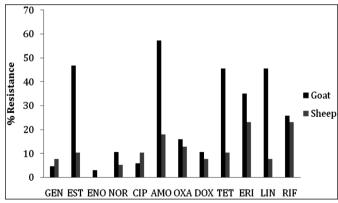


Fig.1. Frequencies of resistance of *Staphylococcus* spp. isolated from mastitis in goats and sheep according to the disk diffusion test. GEN: gentamycin, EST: estreptomycin, ENO: enrofloxacin, NOR: norfloxacin, CIP: ciprofloxacin, AMO: amoxicillin, OXA: oxacillin, DOX: doxycycline, TET: tetracycline, ERI: eritromycin, LIN: lincomycin and RIF: rifampicin.

Table 3. Distribution of *Staphylococcus* spp. isolates from small ruminant mastitis in Brazil according to resistance genes and multidrug resistance rates (MDR)

-		-				-	-				
Resistance gene pattern		Multidrug resistance rate (MDR)									
	0	0.14	0.29	0.43	0.57	0.71	0.86	1			
blaZ	1	6	4	5	9	4	7	-			
ermA	-	-	-	-	-	-	-	1			
ermB	-	-	-	1	-	1	-	-			
ermC	-	-	-	-	-	3	3	1			
icaD	4	6	8	2	-	-	-	-			
blaZ, icaD	5	7	6	3	5	1	3	-			
blaZ, ermB	1	1	2	-	-	-	1	-			
blaZ, ermC	-	-	-	1	-	4	5	-			
blaZ, msrA	-	-	-	-	-	-	1	-			
blaZ, ermA, icaD	-	-	-	-	-	-	1	-			
blaZ, ermB, ermC	-	-	-	-	-	4	-	2			
blaZ, ermB, icaD	-	1	1	-	1	1	-	-			
blaZ, ermC, icaD	-	1	1	-	1	-	-	-			
blaZ, ermA, ermB, icaD	-	-	-	-	-	-	-	1			
ermB, ermC	-	-	-	-	-	1	-	-			
icaD, msrA	1	-	-	-	-	-	-	-			
icaD, ermC	-	-	1	-	-	2	1	-			
icaD, ermB, ermC	-	1	1	1	-	-	-	-			
No pattern	4	15	18	14	8	7	4	3			
TOTAL	16	38	42	27	24	28	26	8			

gentamycin (4.7% vs. 7.7%). No resistance to gentamycin was observed in *Staphylococcus aureus* isolates.

The MDR rate ranged from 0 to 1 (Table 3). Fifteen (7.1%) isolates were susceptible to all tested antimicrobial drugs, ten of which (4.8%) originated from mastitis in goats and five (2.4%) in sheep. Multidrug resistance was more frequently observed in goat isolates (44.7%) than in sheep isolates (23.3%). Seven goat isolates (3.4%) and one sheep isolate (0.5%) were resistant to all tested drugs. A high number of isolates (n=88) exhibiting simultaneous resistance to four or more groups of antimicrobial drugs were observed in this study. Eight (3.81%) isolates were resistant to all tested drugs.

Of the 210 isolates tested, 41 (19.6%) were considered positive for the efflux pump mechanism, 34 isolates originating from goat mastitis and seven isolates from sheep mastitis. Considering the host species, a similar rate for the efflux pump was observed between goat (18%) and sheep (23.3%) isolates. Considering the red Congo agar method for biofilm production, only 16 (7.6%) isolates (15 isolates from goat and one from sheep) tested positive for biofilm.

Table 4 presents the frequencies of antimicrobial resistance markers in the Staphylococci isolates investigated. Seventy three isolates (54 from goats and 19 from sheep) were negative to all resistance genes tested and 69 isolates presented two or more resistance genes simultaneously. All isolates tested were negative for *mecA*, although the *blaZ* gene was detected in 96 (45.8%) isolates. A higher occurrence of *blaZ* was observed among CNS, reaching 73% in *S. epidermidis* and 56.6% in *S. caprae*. On the other hand, the frequency of *blaZ* in *S. aureus* and *S. intermedius* was 20.6% and 25.7%, respectively. Considering the host species, 78 (45.7%) isolates from goats and 18 (46.1%) from sheep harbored the *blaZ* gene.

In regard to other resistance markers, *ermA*, *ermB* and *ermC* markers were observed in three (1.4%), 22 (10.5%) and 34 (16.2%) isolates, respectively. The three *ermA* positive isolates and all 34 staphylococci harboring the *ermC* gen originated from goat mastitis. Moreover, of the 22 isolates harboring *ermB*, only one (4.6%) was isolated from sheep. Association among *blaZ*, *ermA*, *ermB*, *ermC* and the efflux pump were observed in 17 isolates, 14 originating from goats and three from sheep.

From the 41 staphylococci positive for the efflux pump mechanism, only two (1.0%) harbored the *msr*A gene, according to the PCR results. The biofilm-encoding gene *ica*D was detected in 69 (33.0%) isolates (Table 4).

DISCUSSION AND CONCLUSIONS

A high rate of antimicrobial resistance to staphylococci isolated from goat and sheep mastitis in Northeastern Brazil was observed in the present study. This may be associated with the greater use of antimicrobial drugs in this species, because sheep in the region under evaluation have the double use of producing milk and meat and are carried in a more

 Table 4. Distribution of resistance genes in 210 Staphylococcus spp. isolates from small ruminant mastitis in Brazil according to PCR results

Species	Total				Goats				Tota	1			Sheep)		
		icaD	mecA	blaZ	ermA	ermB	ermC	msrA		icaD	mecA	blaZ	ermA	ermB	ermC	msrA
S. aureus (n=34)	13	5 (38.5%)	-	6 (46.1%)	-	2 (15.4%)	-	-	1	-	-	1 (100%)	-	-	-	-
S. caprae (n=23)	27	10 (37.0%)	-	12 (44.5%)	-	1 (3.7%)	3 (11.1%)	1 (3.7%)	1	-	-	1 (100%)	-	-	-	-
S. epidermidis (n=37)	58	14 (24.1%)	-	24 (41.4%)	-	9 (15.5%)	11 (19.0%)	-	5	2 (40%)	-	3 (60%)	-	-	-	-
S. gallinarium (n=5)	10	3 (30%)	-	2 (20%)	-	3 (30%)	2 (20%)	-	0	-	-	-	-	-	-	-
S. hyicus (n=35)	27	11 (40.7%)	-	11 (40.7%)	-	1 (3.7%)	4 (14.9%)	-	6	3 (50%)	-	3 (50%)	-	-	-	-
S. intermedius (n=35)	17	5 (29.4%)	-	4 (23.5%)	2 (11.8%)	2 (11.8%)	4 (23.5%)	-	12	6 (50.0%)	-	5 (41.7%)	-	1 (8.3%)	-	-
S. saprophyticus (n=1)	0	-	-	-	-	-	-	-	1	-	-	1 (100%)	-	-	-	-
S. simulans (n=3)	4	3 (75%)	-	-	-	-	-	1 (25%)	0	-	-	-	-	-	-	-
Staphylococcus spp. (n=37)	38	5 (13.2%)	-	19 (50%)	1 (2.6%)	3 (7.9%)	10 (26.3%)	-	6	2(33.3%)	-	4 (66.7%)	-	-	-	-
TOTAL	194	56	-	78	03	21	34	2	32	13	0	18	0	1	0	0

extensive way. The same fact is reported by Aires-de-Sousa et al. (2007) comparing the sensitivity of *Staphylococcus aureus* isolates from bovine, goats and sheep.

One hundred fifty five (74%) isolates were resistant to two or more antimicrobial drugs simultaneously and pan-susceptibility was observed in only fifteen *Staphylococcus* spp. isolates (7.1%). Although the antimicrobial resistance observed in the present study seems to be higher than those previously reported in Brazil and elsewhere (Aires-de-Sousa et al. 2007, Kumar et al. 2009), direct comparisons must be performed carefully since most investigations focused only on Staphylococcus aureus. Except for enrofloxain and norfloxacin, the present results indicate lower resistance rates to streptomycin, tetracycline and amoxicillin in Staphylococcus aureus as compared to CNS (Table 2). However, the multidrug resistant rate indicated here is higher than that (8%) recently reported in a study on antimicrobial resistance in coagulase positive and negative staphylococci from sub-clinical mastitis in goats (Virdis et al. 2010).

The frequency of resistance to beta-lactams varied considerably and ranged from 12% (doxycycline) to 50% (amoxicillin). This rate of resistance to amoxicillin was higher than that described in *S. aureus* from mastitis in Brazilian dairy goats (Aires-de-Sousa et al. 2007) but very similar to beta-lactam resistance rates reported worldwide for *Staphylococcus aureus* from cattle mastitis (Rabello et al. 2005, Sawant et al. 2009, Virdis et al. 2010).

There is no previous study regarding distribution of the *blaZ* gene in small ruminant species. In the present study, *bla*Z was detected in 45.8% of all *Staphylococcus* spp. isolates, which supports the high resistance level against beta-lactams observed in the disk diffusion test, particularly for the goat species. In Staphylococcus spp. isolates from sheep, the higher *bla*Z gene frequency is unrelated with the disk diffusion test (Fig.1). Discrepancies in genetic and phenotypic tests are reported in previous studies in dairy cattle (Sawant et al. 2009). The frequency of the *bla*Z gene in coagulase positive staphylococci (20.6% in S. aureus and 25.7% in S. intermedius) was lower than that observed in many CNS (73% in S. epidermidis; 56% in S. *caprae*, and 62% in CNS). The frequency of *bla*Z described here was higher than the 18.4% reported by Sawant et al. (2009) but lower than the 100% observed in penicillin resistant S. aureus from Denmark (Vesterholm-Nielsen et al. 1999). The results indicate that beta-lactamase production encoded by *bla*Z might be the main mechanism associated with resistance to beta-lactams in staphylococci causing mastitis in goats and sheep in Brazil, as previously reported in other countries for dairy cattle (Olsen et al. 2006, Taponen & Pyörälä, 2009).

Resistance to methicillin was found in 16.7% of the *Staphylococcus* isolates, but none of them harbored the *mecA* gene. Although a recent study (Onni et al. 2010) reported multi-drug resistance mediated by the *mecA* gene in *S. epidermidis* isolated from dairy sheep herds, methicillin resistant *Staphylococcus* spp. are considered unusual in veterinary medicine (Guérin-Faublée et al. 2003). This type of resistance is associated with resistance to all beta-lactam drugs, despite the apparent susceptibility obtained *in vitro*

tests (CLSI 2006, Sawant et al. 2009). In Brazil, a frequency of 5.6% of the *mecA* gene in CNS is of animal origin, but no detection of CNS in humans was reported (Soares et al. 2008). Transfer of beta-lactam resistance genes may occur among human and animal *Staphylococcus* spp. isolates and is a public health concern (Vesterholm-Nielsen et al. 1999, Katayama et al. 2005, Olsen et al. 2006). Caution should be exercised in regard to raw milk ingestion, which is a common practice in rural areas under development. However, further studies for the purpose of determining the real importance of MRSA from animal-derived foods on public health issues are required, since MRSA isolated from food animals do not necessarily correspond genetically to those clones linked to human infections (Kwon et al. 2005).

Of the 71 erythromycin resistant isolates, only 41 presented efflux pump in MH agar screening and two and 48 isolates were positive to msrA and erm genes by PCR respectively. The resistance to macrolides and tetracycline may occur due to several mechanisms in addition to efflux pump, such as modification enzymes and ribosome protection (Butave et al. 2003, Gatermann et al. 2007, Hassan et al. 2007). The results showed high resistance rates against erythromycin in S. caprae (43%), S. epidermidis (43%), and in other non-identified CNS (56%). This corroborates a previous report showing high resistance rates for erythromycin in *S. epidermidis*, and a high frequency of the mrsA gene in those species (Sawant et al. 2009). Possible differences in phenotypic and genotypic resistance to erythromycin may be explained by other unidentified macrolide resistance mechanisms which are common in CNS isolates (Fiebelkorn et al. 2003). These are ultimately related to geographic location and therapeutic drug usage (Pitkälä et al. 2004, Gatermann et al. 2007). The presence of two or more antimicrobial resistance genes may be associated with multidrug resistant Staphylococcus spp. (Sawant et al. 2009).

The high multidrug resistance rates observed in CNS as compared to CPS are in accordance with previous reports (Kumar et al. 2008, Sawant et al. 2009) and support the hypothesis that CNS might play an important role as a source of genes resistant to S. aureus (Taponen & Pyörälä, 2009). Considering such an assumption, studies about host specificity in Staphylococcus aureus and other pathogenic agents could provide useful information about the epidemiological importance of CNS as reservoirs of genes resistant to pathogenic strains for humans. However, because CNS are the major mastitis causing agents in small ruminants, the high frequency of resistant genes in such species reported here is worth noting. Although goats and sheep are raised on pasture in extensive or semi-intensive production systems in Brazil, the widespread use of some antimicrobials for clinical treatment purposes might be associated with the high resistance rates (Ribeiro et al. 2009, Schmidt et al. 2009), as observed in the present study. Uncontrolled antimicrobial usage was reported by farmers and agricultural technicians in the sampling region. Indeed, high resistance levels against specific antimicrobials may be associated with their frequent and long-term use on the farm (Pitkälä et al. 2004, Kumar et al. 2009).

The drugs showing the lowest resistance rates were gentamycin, enrofloxacin and ciprofloxacin, which are in accordance with other studies on mastitis in several ruminant species (Hussain et al. 2007, Kumar et al. 2009). Virdis et al. (2010) reported high sensitivity to quinolones, but not to aminoglycoside in subclinical mastitis-causing staphylococci.

No association was observed between phenotypic and molecular tests for biofilm production, corroborating previous studies (Baselga et al. 1993, Arciola et al. 2001, Vasuvedan et al. 2003, Oliveira et al. 2006). Only seven isolates were positive for both methods simultaneously. The Congo Red Agar test is a very simple and inexpensive technique, but correct interpretation of the results depends on many factors, such as colony color characteristics (Baselga et al. 1993). PCR detected more biofilm forming isolates than the Congo red agar test, supporting previous findings (Oliveira et al. 2006). However, the present study focused on the icaD gene alone, while other genetic loci (bap, agr and sar) might be associated with biofilm formation as well (Oliveira et al. 2006). The main occurrence of biofilm in goat isolates is associated with antimicrobial resistance and multidrug resistance, as reported in previous studies (Baselga et al. 1993, Oliveira et al., 2006).

The high antimicrobial resistance rates detected in staphylococci causing mastitis in small ruminants in Northeastern Brazil might be associated with several resistant determinant genes. Considering the lack of information about antimicrobial resistance in staphylococci causing mastitis in small ruminants in Brazil, the present findings could be of great importance in developing strategies for mastitis treatment in the region.

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