

OXACILIN-RESISTANT COAGULASE-NEGATIVE STAPHYLOCOCCI (CONS) BACTEREMIA IN A GENERAL HOSPITAL AT SÃO PAULO CITY, BRASIL

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ABSTRACT

In the last decades, coagulase-negative staphylococci (CoNS), especially *Staphylococcus epidermidis* have become an important cause of bloodstream infections. In addition, rates of methicillin-resistance among CoNS have increased substantially, leading to the use of glycopeptides for therapy. The objective of this study was to evaluate eleven consecutive clinically relevant cases of oxacillin-resistant CoNS bacteremia in a general hospital localized in São Paulo city, Brazil. Five different species were identified by different phenotypic methods, including *S. epidermidis* (5), *S. haemolyticus* (3), *S. hominis* (1), *S. warneri* (1) and *S. cohnii* subsp *urealyticus* (1). A variety of Pulsed Field Gel Electrophoresis profiles was observed by macrorestriction DNA analysis in *S. epidermidis* isolates, but two of three *S. haemolyticus* isolates presented the same profile. These data indicated the heterogeneity of the CoNS isolates, suggesting that horizontal dissemination of these microorganisms in the investigated hospital was not frequent. One *S. epidermidis* and one *S. haemolyticus* isolates were resistant to teicoplanin and susceptible to vancomycin. The selective pressure due to the use of teicoplanin in this hospital is relevant.

Key words: Coagulase-negative staphylococci; bacteremia; oxacillin; PFGE

INTRODUCTION

Coagulase-negative staphylococci (CoNS) are major causes of nosocomial bloodstream infection and responsible for high morbidity and mortality rates, mainly in hospitalized patients (9). Members of the genera *Staphylococcus* are catalase-positive, gram-positive cocci, coagulase-negative, aerobes and, when present in human infections, can present multiresistant profiles (8,20). These strains may constitute a dangerous reservoir of resistance genes in a hospital (20).

Staphylococci generally present a benign or symbiotic relationship with their host. However, they may become pathogens when entering the host tissue through break of the cutaneous barrier, inoculation by needles or implantation of medical devices (5). It is increasingly important to accurately identify CoNS isolates to the species level in order to determine

the clinical significance of these bacteria, the proper epidemiological surveillance, and the management of patients infected with CoNS in case of relapse (15).

A substantial increase in the frequency of oxacillin-resistance (methicillin-resistant) in CoNS isolates has occurred over the last decades (4). Between 50% and 80%, depending on the species, are *mec A* positive or oxacillin resistant (1,6).

According to the results of the SENTRY antimicrobial surveillance program, carried out with Brazilian bloodstream isolates over a five-year period from 1997 to 2001, the oxacillin susceptibility in *Staphylococcus aureus* was 68.2% and 19.2% in CoNS (17).

Staphylococcus epidermidis and *Staphylococcus haemolyticus* are the most frequent species in nosocomial infections, and the frequency of oxacillin resistance is higher in CoNS clinical isolates (3). *S. haemolyticus* have been reported

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to show multiple resistance to antimicrobials and quite frequently clinical isolates present with reduced susceptibility or are resistant to teicoplanin (16).

Vancomycin is usually considered the treatment of choice for infections caused by these microorganisms. However, due to the emergence of vancomycin-resistant enterococci (9) and vancomycin-resistant staphylococci (19), reduction in the use of this drug has been recommended (20). A few reports have shown that the mechanism of glycopeptide resistance in *S. epidermidis*, *S. haemolyticus* and *S. hominis* is similar to that described in VISA and hetero-VISA strains (13). The objective of this study was to evaluate eleven consecutive clinically relevant cases of oxacillin-resistant CoNS bacteremia in a general hospital where therapy with the glycopeptide teicoplanin is broadly utilized.

MATERIALS AND METHODS

Bacterial isolates

The study was carried out with eleven consecutive bloodstream CoNS isolates, obtained between June and July 2005 from patients at 9 de Julho Hospital, a 250 beds general hospital localized in the city of São Paulo, Brazil. According to the CDC criteria, these isolates were considered clinically relevant by the National Nosocomial Infections Surveillance Committee of the hospital (12).

Identification

Staphylococci identification was carried out by test for oxidation-fermentation, coagulase (Laborclin, Brazil), catalase, alkaline phosphatase (Sigma-Aldrich, Germany), ornithine (Merck, Germany), urease (Oxoid, UK), PYR (pyrrolidinyll- β -naphthylamide hydrolysis, Probac do Brasil, Brazil), hemolysis in sheep blood agar, acid production from trehalose (Sigma-Aldrich, Germany), mannitol (Nuclear, Brazil), mannose (Vetec, Brazil), sucrose (Reagen, Brazil), maltose (Sigma-Aldrich, Germany), lactose (Difco, USA), cellobiose (Sigma-Aldrich, Germany) and anaerobic growth in thioglycolate (Merck, Germany). Susceptibility to novobiocin (Oxoid, UK), polymyxin B (Oxoid, UK), bacitracin (CECON, Brazil), desferrioxamine (Ciba Geigy, Switzerland) and fosfomicin (Oxoid, UK) was also determined. Isolates were kept frozen at -20°C in Skim Milk (Difco, USA). Bacteria to be tested were suspended in 0,5 ml of saline to a McFarland standard and 50ml were added to each sugar carbohydrate tube. The acid production from carbohydrates was evaluated after 24, 48 and 72hs of incubation at 35-37°C. The final evaluation was at the 7th day. The phenotypic tests were accomplished in parallel with a positive control (*S. epidermidis* ATCC 12228).

Antimicrobial susceptibility

The isolates were tested for susceptibility to oxacillin, vancomycin and teicoplanin by the agar disk diffusion method

with Mueller-Hinton agar plates (Difco, USA) according to Clinical Laboratory and Standards Institute (CLSI 2005) recommendations and confirmed by Etest (AB Biodisk, Sweden). The susceptibility to novobiocin, polymyxin B, bacitracin, desferrioxamine and fosfomicin was determined according to Monsen *et al.* (10).

PFGE typing

Chromosomal DNA CoNS was prepared in agarose blocks and digested with *Sma*I (New England BioLabs, USA), as described elsewhere (14). The isolates were run on a 1% agarose gel (Invitrogen, USA) in a CHEF DRIII system (Bio-Rad, USA) under the following conditions: run time, 23 h; temperature, 13°C; voltage, 200 V; initial forward time, 5 s; final forward time, 60 s. The molecular weight markers (New England BioLabs, USA) were run in the first and in the last lane. The gels were stained with ethidium bromide, washed in water, and photographed under UV light by using the Gel Doc 1000 system (Bio-Rad, USA). The gel patterns were read by visual inspection. The isolates were classified as identical if they shared the same band profile, and isolates differing by more than six bands were considered to represent distinct DNA types (22).

RESULTS AND DISCUSSION

A total of eleven CoNS isolates belonging to five different species were identified including *S. epidermidis* (5 isolates), *S. haemolyticus* (3 isolates), *S. hominis* (1 isolate), *S. warneri* (1 isolate) and *S. cohnii* subsp *urealyticus* (1 isolate) (Table 1). All isolates, except the number 20994, were identified to the species level by the conventional method of Kloos and Banermann (7,8). The isolate 20994 could not be identified by the conventional method, so it was identified by Vitek-2 (bioMérieux, France) as *S. cohnii* subsp *urealyticus*.

The two species most frequently encountered were *S. epidermidis* and *S. haemolyticus*.

The *S. epidermidis* and *S. hominis* isolates were identified by the disk diffusion susceptibility test to desferrioxamine, since other species of CoNS are resistant to desferrioxamine. To differentiate *S. epidermidis* from *S. hominis*, other phenotypic tests were used, as fermentation of trehalose (negative for *S. epidermidis*), alkaline phosphatase (positive for *S. epidermidis*) and growth in thioglycolate (positive for *S. epidermidis*).

The test for production of urease allowed the differentiate *S. haemolyticus* (urease negative), from *S. epidermidis*, *S. hominis* and *S. warneri* (urease positive). The test of positive PYR along with the hemolytic properties in sheep blood agar and absence of fermentation of mannose allowed the differentiation of *S. haemolyticus*, the second most prevalent species.

S. cohnii subsp *urealyticus* (isolate 20994) was the less common specie. Classical tests of resistance to novobiocin,

Table 1. Bloodstream Methicillin resistant Coagulase-negative staphylococci isolated from patients at Hospital 9 de Julho (June/July 2005).

Specie	Clinical isolate	Isolation date	Ward	MIC Vanco (µg/ml)	MIC Teico (µg/ml)	PFGE Profile
<i>S. epidermidis</i>	21170	17/07/05	internal medicine	2.0	16.0	“A”
<i>S. epidermidis</i>	21169	11/07/05	ICU	1.5	4.0	“B”
<i>S. epidermidis</i>	21168	20/07/05	oncology	1.5	4.0	“C”
<i>S. epidermidis</i>	20944	29/06/05	ICU	2.0	4.0	“D”
<i>S. epidermidis</i>	21171	01/07/05	oncology	2.0	4.0	“E”
<i>S. haemolyticus</i>	20995	07/07/05	internal medicine	2.0	12.0	“G”
<i>S. haemolyticus</i>	21172	13/07/05	oncology	1.5	4.0	“F”
<i>S. hominis</i>	20947	20/06/05	ICU	0.75	0.75	NC
<i>S. warneri</i>	20993	08/07/05	internal medicine	1.0	2.0	NC
<i>S. haemolyticus</i>	20946	24/06/05	internal medicine	2.0	3.0	“F”
<i>S. cohnii</i> subsp <i>urealyticus</i>	20994	06/07/05	oncology	1.0	3.0	NC

NC – without classification (only one isolate).

production of urease and absence of sucrose fermentation confirmed the identification of the specie. A discrepancy in the alkaline phosphatase production was noted: the result was negative in the conventional test and positive in the automated system.

All isolates were methicillin-resistant by the disk diffusion test, with MICs ≥ 256 µg/ml by E-Test. Two isolates (*S. epidermidis* and *S. haemolyticus*) presented reduced susceptibility to teicoplanin (Table 1). Strains with this characteristic have been reported by Nunes *et al.* (13) and may be associated with treatment failures or may become precursors of glycopeptide-resistant strains (18).

S. cohnii subsp *urealyticus* is an unusual opportunist species that has been found in hospital environment like pediatric ICUs (24), and may constitute a dangerous reservoir of multiple antimicrobial plasmid mediated resistance genes (21,23).

Among the five clinical isolates of *S. epidermidis* five different patterns of PFGE were observed, indicating absence of clonal dissemination among the patients. The same did not occur with the three clinical isolates of *S. haemolyticus*, where two isolates from patients at different wards, more than one month apart, presented the same profile, suggesting nosocomial transmission (Fig. 1).

In the last years, the importance of CoNS has been increasing due to their pathogenicity and involvement in human diseases. Their identification species in the clinical laboratories is important but not an easy task, because classical phenotypic tests do not differentiate them from other staphylococci require more time in the identification compared to commercial kits. Many clinical laboratories use automated systems for identification of *Staphylococcus* spp., although the reliability of results for

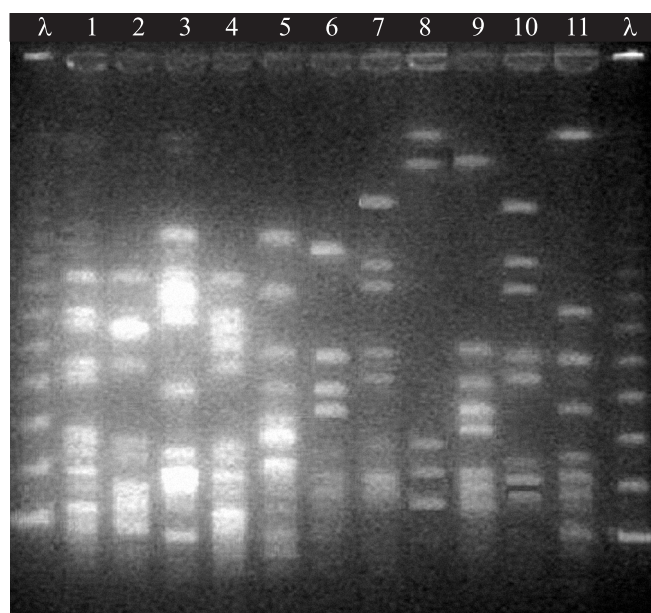


Figure 1. PFGE profile of *Sma*I-digested chromosomal DNA of CoNS isolates, obtained from patients in 9 de Julho Hospital in São Paulo city, Brazil. λ lambda ladder DNA markers; lanes 1-5 *S. epidermidis*; lanes 6,7 and 10: *S. haemolyticus*; lane 8: *S. hominis*; lane 9: *S. warneri*; lane 11: *S. cohnii* spp *urealyticus*.

certain species is not always satisfactory, particularly for species other than *S. epidermidis*.

Two isolates, one *S. epidermidis* (isolate 21170) and one *S. haemolyticus* (isolate 20995) presented high MIC for teicoplanin, but were susceptible to vancomycin (Table 1).

At 9 de Julho hospital, staphylococcal infections, mainly those caused by *S. aureus*, have been successfully treated with teicoplanin instead of vancomycin, for more than a decade, particularly IV-catheter related infections. In counter part, treatment of central nervous infections and endocarditis with teicoplanin has been less effective, probably due to the occurrence of oxacillin-resistant CoNS also resistant to teicoplanin. A surveillance program of glycopeptide resistance and adequate CoNS specie identification have great importance in determination of risk factors and implementation of nosocomial infection control measures.

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RESUMO

Bacteremias por *Staphylococcus* coagulase negativos oxacilina resistentes em um hospital na cidade de São Paulo, Brasil

Staphylococcus coagulase negativos (SCoN), especialmente *Staphylococcus epidermidis* tem se tornado causa importante de infecções da corrente circulatória nas últimas décadas. Além disso, percentuais de resistência a meticilina entre os SCoN têm aumentado significativamente, levando ao uso de glicopeptídeos nestes pacientes. O objetivo deste estudo foi avaliar onze casos consecutivos de bacteremia clinicamente relevantes por SCoN oxacilina resistentes em um hospital localizado na cidade de São Paulo, Brasil. Cinco diferentes espécies foram identificadas por diferentes métodos fenotípicos, incluindo *S. epidermidis* (5), *S. haemolyticus* (3), *S. hominis* (1), *S. warneri* (1) e *S. cohnii* subsp *urealyticus* (1). Diferentes perfis eletroforéticos obtidos pela técnica de “Pulsed Field Gel Electrophoresis” foram observados na análise da macrorestrição do DNA nos isolados de *S. epidermidis*, mas dois dos três isolados de *S. haemolyticus* apresentaram o mesmo perfil. Esses dados indicam uma heterogeneidade nos isolados SCoN, sugerindo que a disseminação horizontal no hospital investigado não é freqüente. Um isolado de *S. epidermidis* e um de *S. haemolyticus* foram resistentes à teicoplanina e sensíveis à vancomicina. Observa-se a relevância da pressão seletiva pelo uso de teicoplanina nos pacientes deste hospital.

Palavras-chave: *Staphylococcus* spp. coagulase negativo; bacteremia; oxacilina; PFGE

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