ARTIGO ORIGINAL

**Staphylococcus** spp. resistentes em hemoculturas e superfícies hospitalares e a segurança do paciente

*Staphylococcus* spp. resistant in blood cultures and hospital surfaces and the patient safety

*Staphylococcus* spp. resistentes em hemocultivos y superfícies hospitalarias y la seguridad del paciente

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**RESUMO**

Justificativa e Objetivos: Este estudo teve como objetivo avaliar a presença e resistência de *Staphylococcus aureus* e estafilococos coagulase-negativos resistentes à oxacilina isolados em superfícies e hemoculturas em uma Unidade de Terapia Intensiva de um hospital de emergência na cidade de Caruaru, Pernambuco, Brasil. **Métodos:** Trata-se de um estudo descritivo, do tipo transversal, sendo as coletas realizadas a partir das superfícies dos leitos de uma Unidade de Terapia Intensiva (UTI). Paralelamente, amostras de hemoculturas dos respectivos pacientes também foram obtidas. As amostras foram coletadas por amostragem de conveniência e a identificação dos microrganismos isolados se deu através de espectrometria de massas. **Resultados:** Observou-se uma prevalência maior de bactérias do grupo dos *Staphylococcus* coagulase negativo-SCN (76,92%) em relação à espécie de *Staphylococcus aureus* (23,07%). Dentre os micrororganismos identificados como resistentes à oxacilina, 12,5% das espécies apresentaram semelhança nas amostras de superfície do leito e hemocultura do mesmo paciente. **Conclusões:** As bactérias resistentes identificadas reforçam a importância de enfatizar as medidas gerais de higienização do ambiente no que se refere a segurança do paciente nos serviços de saúde.

**Descritores:** Infeção Hospitalar. Contaminação de Equipamentos. Unidades de Terapia Intensiva. Infecções Estafilocócicas.

**ABSTRACT**

**Background and Objectives:** This study aimed to evaluate the presence and resistance of *Staphylococcus aureus* and coagulase-negative *Staphylococcus* resistant to oxacillin isolated in surfaces and blood cultures in an Intensive Care Unit of an emergency hospital in the city of Caruaru, Pernambuco, Brazil. **Methods:** This is a descriptive, cross-sectional study, the collections being performed from the bed surfaces of an Intensive Care Unit (ICU). In parallel, blood samples from the respective patients were also obtained. Samples were collected by convenience sampling and the identification of the isolated microorganisms was done by mass spectrometry. **Results:** We observed a higher prevalence of bacteria from the group of coagulase-negative *Staphylococci* - CoNS (76.92%) in relation to the species *Staphylococcus aureus* (23.07%). Among the microorganisms identified as resistant to oxacillin, 12.5% of the species were similar in the bed surface samples and blood cultures from the same patient sam-ple. **Conclusions:** Resistant bacteria identified reinforce the importance of emphasizing the general measures of environmental hygiene in relation to patient safety in health care.

**Keywords:** Cross Infection. Equipment Contamination. Intensive Care Units. Staphylococcal Infections.
RESUMEN

Justificación y Objetivos: Este estudio tuvo como objetivo evaluar la presencia y resistencia de Staphylococcus aureus y estafilococos coagulasa-negativos resistentes a la oxacilina aislados en superficies y hemocultivos en una Unidad de Terapia Intensiva de un hospital de emergencia en la ciudad de Caruaru, Pernambuco, Brasil. Métodos: Se trata de un estudio descriptivo, del tipo transversal, siendo las colectas realizadas a partir de las superficies de los lechos de una Unidad de Terapia Intensiva (UTI). Paralelamente, se obtuvieron muestras de hemocultivos de los respectivos pacientes. Entre los microorganismos identificados como resistentes a la oxacilina, el 12,5% de las especies presentaron semejanza en las muestras de superficie del lecho y hemocultura del mismo paciente. Conclusiones: Las bacterias resistentes identificadas refuerzan la importancia de enfatizar las medidas generales de higienización del ambiente en lo que se refiere a la seguridad del paciente en los servicios de salud.


INTRODUCTION

The Intensive Care Units (ICUs) are considered a critical hospital environment due, among other factors, the significant occurrence of infections by multidrug-resistant bacteria. These infections make the alternative treatment being reduced, extending the period of hospitalization of patients, raising costs for the hospital and generating increased mortality rates. The knowledge of the microbiota of each ICU is important in order to contribute in a more targeted and rational use of antibiotics.1,2

Colonized and infected patients may contaminate the environment, turning inanimate surfaces and equipment potential reservoirs of bacteria, particularly the resistant to antimicrobials.3 The importance of the environment as a reservoir of microorganisms was approached, especially in recent decades, when it was considered a secondary factor in the transmission of nosocomial infections, which are those acquired in the hospital during the delivery of health care or a few days after discharge from patient including bloodstream infections.4,5

The realization of blood cultures represents a significant role in the diagnosis of infections, in order to verify the occurrence of the microorganisms that cause bacteremia.6 The frequency of pathogenic Gram positive isolated organisms, especially Staphylococcus aureus and coagulate-negative Staphylococi (CoNS), have increased significantly in ICU environments.2,7,8 The nasal cavity has been identified as the most prevalent region for the isolation of these microorganisms in hospitalsover, the hands have also been considered an important mean of bacterial transmission in the hospital between patients and professionals, contributing to the increase of cross-infection cases.9

The frequent use of penicillins, such as methicillin and oxacillin in the treatment of staphylococcal infections, favored the emergence of strains resistant to these antibiotics, called ORSA/ORCS, these microorganisms generally exhibit extended resistance to all β-lactam antibiotics and were first recognized as nosocomial pathogens.10 These strains are considered ORSA due to the production of penicillinase enzyme that confers resistance to semi-synthetic penicillins (methicillin/oxacillin) and β-lactam antibiotics making difficult the treatment of staphylococcal infections.11 The resistance to oxacillin presented by Staphylococcus is determined by the presence of a mecA gene located on chromosome.12

The recurring increased rates of ORSA strains, brings great interest on the bodies responsible for the control of nosocomial infections.13 The tracking of these microorganisms by appropriate methods, should be extended not only to laboratory testing of patients, particularly blood cultures, but also to the environment which the patients rests, like inanimate surfaces near them, given that different resistant organisms have been reported as possible causes of cross-infection.13-15

Nosocomial infections can be controlled through effective measures that interfere with the microorganism transmission process, such as frequent hand washing, proper cleaning of equipment and surfaces of beds, the use of personal protective equipment and the adoption of aseptic measures.15

This study aimed to evaluate the presence and resistance profile of Staphylococcus aureus and coagulase-negative Staphylococci resistant to oxacillin isolated in surfaces and blood cultures in an Intensive Care Unit of an emergency hospital in the city of Caruaru, Pernambuco, Brazil.

METHODS

Cross-sectional study, conducted between April 2015 and April 2016, on the ICU of a hospital emergency in the city of Caruaru, Pernambuco, Brazil. The study hospital has 225 beds for hospitalization, 19 of these composing the adult ICU. The unit belongs to the macro-regional health population of Caruaru, which covers 87 municipalities of the micro-regions of Caruaru, Garanhuns, Arcoverde, Afoados da Ingaízea and Serra Talhada, being a reference in traumatology, traumato-orthopedics, surgery general and maxillofacial complex of high complexity.

The blood bottles were supplied by the hospital to the study participants on spontaneous demand, according to the diagnostic routine performed by the Laboratory of Clinical Microbiology by BD BACTEC™ Instrumented Blood Culture Systems (Becton, Dickinson and Company), previously collected as their own methodology of aseptic standards, with inclusion criteria patients admitted to the ICU hospital for a maximum of two
months during the study period. The biological samples obtained from blood culture were being in Blood Agar, MacConkey Agar and Mueller Hinton bacterial isolation purposes and implementation of standardized test of antimicrobial susceptibility.

The collections of inanimate surfaces were performed by convenience sampling from five pre-defined surfaces of the ICU bed (right and left bars of the beds, bed control buttons, bedside table and buttons from the infusion pump). The inclusion criterion was samples from surfaces whose beds were occupied by their respective patients with positive blood culture indication by **BD BACTEC™ Instrumented Blood Culture Systems**. The obtaining of samples was performed by using moistened sterile swabs on Tryptic Soy Broth (TSB), that immediately after the collection were stored in tubes containing the broth and incubated in a greenhouse at a temperature of 37°C for 24 hours for analysis of the bacterial growth. After checking the turbidity in TSB was performed by seeding exhausted in blood agar and MacConkey, incubated in 37°C for 24 hours.

The genus level identification and species of isolated microorganisms was made by mass spectrometry type MALDI-TOF (Matrix Associated Laser Desorption-Ionization - Time of Flight). The mass spectra were obtained in duplicate and compared with the stored software database Biotype MALDI 2.0 (Bruker Daltonics) for the identification of bacterial genus and species.

The identified bacteria were subjected to the antibiogram carried by the diffusion method of Kirby-Bauer and for analyzing of the results regarding the resistance pattern, was used the standardize of the Clinical and Laboratory Standards Institute 2016 (CLSI). Discs containing the following antimicrobials were used: ciprofloxacin, nitrofurantoin, trimethoprim/sulfamethoxazole, chloramphenicol, cefoxitin, tetracycline, amikacin, erythromycin, clindamycin and gentamicin. The tracking of ORSA and ORCNS occurred by the use of cefoxitin antimicrobial, which is a surrogate marker for detection of oxacillin resistance in strains that have the gene meca. For the isolates classified as ORSA/ORCNS by disk diffusion method (Kirby-Bauer method), a test was performed to determine the minimum inhibitory concentration (MIC) for the antibiotic vancomycin using strips containing different concentrations of this antimicrobial agent (**E-test**). This method was used to determine with certainty the phenotypic resistance pattern ORSA strains and ORCNS front vancomycin.

This work was approved by the Research Ethics Committee of the Centro Universitário Tabosa de Almeida (ASCES-UNITA) through the number 1.061.201.

**RESULTS**

Samples from 12 ICU beds were obtained and a sample of blood culture of the corresponding patient bed, totaling 60 samples of environmental surfaces and 12 blood cultures.

The bacterial growth was detected in 52 (86.66%) surfaces samples, 4 of them (7.69%) positive for the growth of more than one microorganism, totaling 57 bacteria species identified (Table 1).

Regarding isolated environmental species, there was a prevalence of *Staphylococcus epidermidis* representing 30.76% of total isolated, followed by *S. aureus* (23.07%) and *S. haemolyticus* (23.07%) and *S. cohnii* (7.69%), *S. hominis* (7.69%) and *S. warnerii* (7.69%) (Figure 1). As surfaces where these species were isolated, they are described in table 1.

Other organisms found included *Acinetobacter baumannii*, *Acinetobacter pittii*, *Bacillus cereus*, *Bacillus flexus*, *Bacillus weihenstephanensis*, *Enterococcus faecalis*, *Klebsiella

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**Table 1. Bacterial species isolated by surfaces on ICU beds. Caruaru, PE, Brazil, 2015 - 2016.**

<table>
<thead>
<tr>
<th>BED</th>
<th>LEFT BAROF THE BED</th>
<th>RIGHT BAR OF THE BED</th>
<th>BED CONTROLS OF THE BED</th>
<th>SHELF SUPPORT</th>
<th>INFUSION PUMP BUTTONS</th>
</tr>
</thead>
<tbody>
<tr>
<td><strong>B1</strong></td>
<td>Staphylococcus epidermidis e Bacillus cereus</td>
<td>Staphylococcus hominis e Bacillus cereus</td>
<td>Enterococcus faecalis</td>
<td>Staphylococcus epidermidis</td>
<td>Enterococcus faecalis</td>
</tr>
<tr>
<td><strong>B2</strong></td>
<td>Acinetobacter baumannii</td>
<td>Acinetobacter baumannii</td>
<td>Bacillus cereus</td>
<td>Staphylococcus epidermidis</td>
<td>Staphylococcus epidermidis</td>
</tr>
<tr>
<td><strong>B3</strong></td>
<td>Acinetobacter baumannii</td>
<td>Acinetobacter baumannii</td>
<td>Staphylococcus haemolyticus</td>
<td>N/G</td>
<td>Staphylococcus cohnii e Bacillus cereus</td>
</tr>
<tr>
<td><strong>B4</strong></td>
<td>Acinetobacter baumannii</td>
<td>Escherichia coli</td>
<td>Staphylococcus aureus</td>
<td>Staphylococcus aureus e Enterococcus faecalis</td>
<td>N/G</td>
</tr>
<tr>
<td><strong>B5</strong></td>
<td>Bacillus cereus e Staphylococcus warneri</td>
<td>Bacillus cereus</td>
<td>Acinetobacter baumannii</td>
<td>Acinetobacter baumannii</td>
<td>Bacillus weihenstephanensis</td>
</tr>
<tr>
<td><strong>B6</strong></td>
<td>Staphylococcus haemolyticus</td>
<td>Bacillus flexus</td>
<td>Staphylococcus aureus</td>
<td>Staphylococcus haemolyticus</td>
<td>Providencia stuartii</td>
</tr>
<tr>
<td><strong>B7</strong></td>
<td>Providencia stuartii</td>
<td>Acinetobacter baumannii</td>
<td>Acinetobacter baumannii</td>
<td>N/G</td>
<td>N/G</td>
</tr>
<tr>
<td><strong>B8</strong></td>
<td>Pseudomonas aeruginosa</td>
<td>Pseudomonas aeruginosa</td>
<td>Proteus mirabilis</td>
<td>Acinetobacter baumannii</td>
<td>Proteus mirabilis</td>
</tr>
<tr>
<td><strong>B9</strong></td>
<td>Klebsiella pneumoniae</td>
<td>Klebsiella pneumoniae</td>
<td>Bacillus cereus</td>
<td>Acinetobacter baumannii</td>
<td>Bacillus cereus</td>
</tr>
<tr>
<td><strong>B10</strong></td>
<td>Acinetobacter baumannii</td>
<td>N/G</td>
<td>N/G</td>
<td>Enterococcus faecalis</td>
<td>N/G</td>
</tr>
<tr>
<td><strong>B11</strong></td>
<td>Acinetobacter baumannii</td>
<td>Klebsiella pneumoniae</td>
<td>Acinetobacter pittii</td>
<td>N/G</td>
<td>Acinetobacter baumannii</td>
</tr>
<tr>
<td><strong>B12</strong></td>
<td>Enterococcus faecalis</td>
<td>Klebsiella pneumoniae</td>
<td>Klebsiella pneumoniae</td>
<td>Providencia stuartii</td>
<td>Enterococcus faecalis</td>
</tr>
</tbody>
</table>

**SUBTITLE:** B = Bed; N/G = No Growth.
The antibiograms analyzed showed profile resistance of 21 strains Gram positive, the group of Staphylococci, 13 (61.9%) isolated on surfaces of samples, and 8 (38.09%) of blood cultures. Of these, 16 (76.19%) were resistant to cefoxitin (surrogate marker for detection of oxacillin resistance), are classified as ORSA/ORCNS, respectively (Table 2).

siella pneumoniae, Providencia stuartii, Proteus mirabilis, Pseudomonas aeruginosa and Escherichia coli (Table 1).

Among the 12 analyzed blood cultures, a total of 10 (83.33%) samples showed growth on agar plates, and S. epidermidis (40% of total isolated), S. haemolyticus (20%), S. hominis and S. caprae (10% isolates each), gram positive isolates key (Graphic 2). Klebsiella pneumoniae presented in 2 (20%) of the analyzed blood cultures (Figure 2).

The antibiograms analyzed showed profile resistance of 21 strains Gram positive, the group of Staphylococci, 13 (61.9%) isolated on surfaces of samples, and 8 (38.09%) of blood cultures. Of these, 16 (76.19%) were resistant to cefoxitin (surrogate marker for detection of oxacillin resistance), are classified as ORSA/ORCNS, respectively (Table 2).

Figure 1. Frequency of the main species of Staphylococcus isolated in the surfaces of the ICU. Caruaru, PE, Brazil, 2015 - 2016.

Figure 2. Frequency of Staphylococcus isolated in the samples positives of blood cultures. Caruaru, PE, Brazil, 2015 - 2016.
Table 2. Resistance profile to antimicrobials and tracking ORSA strains/ORCNS isolated in the ICU beds. Caruaru, PE, Brazil, 2015 - 2016.

<table>
<thead>
<tr>
<th>MICROORGANISM</th>
<th>BED</th>
<th>SAMPLE SOURCE</th>
<th>ANTIMICROBIAL TESTED</th>
</tr>
</thead>
<tbody>
<tr>
<td>Staphylococcus hominis</td>
<td>01</td>
<td>BLOOD CULTURE</td>
<td>*FOX</td>
</tr>
<tr>
<td>Staphylococcus epidermidis</td>
<td>01</td>
<td>SURFACE</td>
<td>R</td>
</tr>
<tr>
<td>Staphylococcus hominis</td>
<td>01</td>
<td>SURFACE</td>
<td>R</td>
</tr>
<tr>
<td>Staphylococcus epidermidis</td>
<td>01</td>
<td>SURFACE</td>
<td>R</td>
</tr>
<tr>
<td>Staphylococcus haemolyticus</td>
<td>02</td>
<td>BLOOD CULTURE</td>
<td>R</td>
</tr>
<tr>
<td>Staphylococcus epidermidis</td>
<td>02</td>
<td>SURFACE</td>
<td>R</td>
</tr>
<tr>
<td>Staphylococcus haemolyticus</td>
<td>03</td>
<td>BLOOD CULTURE</td>
<td>R</td>
</tr>
<tr>
<td>Staphylococcus epidermidis</td>
<td>03</td>
<td>SURFACE</td>
<td>R</td>
</tr>
<tr>
<td>Staphylococcus aures</td>
<td>04</td>
<td>SURFACE</td>
<td>R</td>
</tr>
<tr>
<td>Staphylococcus epidermidis</td>
<td>05</td>
<td>BLOOD CULTURE</td>
<td>R</td>
</tr>
<tr>
<td>Staphylococcus epidermidis</td>
<td>06</td>
<td>BLOOD CULTURE</td>
<td>R</td>
</tr>
<tr>
<td>Staphylococcus aures</td>
<td>06</td>
<td>SURFACE</td>
<td>R</td>
</tr>
<tr>
<td>Staphylococcus haemolyticus</td>
<td>06</td>
<td>SURFACE</td>
<td>R</td>
</tr>
<tr>
<td>Staphylococcus aures</td>
<td>07</td>
<td>SURFACE</td>
<td>R</td>
</tr>
<tr>
<td>Staphylococcus epidermidis</td>
<td>08</td>
<td>SURFACE</td>
<td>R</td>
</tr>
<tr>
<td>Staphylococcus epidermidis</td>
<td>10</td>
<td>SURFACE</td>
<td>R</td>
</tr>
</tbody>
</table>

Regarding MIC determination, all strains tested had MIC ≤ 2 μg/mL for Staphylococcus aureus and ≤ 4 μg/mL Staphylococcus coagulase negative (Table 2), thus being considered sensitive to vancomycin according to CLSI 2016, as shown in figure 3, which presents halos with diameters that remit sensitivity to the tested antibiotic.16

Figure 3. E-test® result for determine the MIC to vancomycin.

DISCUSSION

Frequent nosocomial infection outbreaks cases are related to environmental contamination. Cross infection can be facilitated by the survival of microorganisms on dry surfaces, and the Staphylococci are one of the most important microorganisms associated with Health Care Related Infections (HCRI), especially in Intensive Care Units (ICU).15,17

Contamination lying around the ICU patients was observed in the present study in 95% of the collected samples. For positive Gram, we observed a higher prevalence of bacteria from the group of coagulase-negative Staphylococci - CoNS (76.92%) in relation to the species Staphylococcus aureus (23.07%). According to the results from the susceptibility test antimicrobial, 87.5% of the samples identified as CoNS showed resistance to oxacillin, corroborating with previous studies that have estimated that the CoNS isolated in hospitals in Brazil, 70% to 90%, usually have shown resistance to oxacillin.17-19

During the last decades, the substantial increase in CoNS isolated in hospital settings, as well as their oxacillin resistance rates has increased significantly its clinical importance, becoming recognized as opportunistic agents often associated with nosocomial infections.19,20

As a result, it reinforces the importance of performing increasingly accurate cultures, especially as regards the antibiogram that will contribute to conduct towards the use of more rational way, in a more targeted therapy, particularly in hospital environment.

Since cases of sepsis are the most relevant and aggravating the clinical outcome of the patients was also evaluated in this study samples of blood cultures to check the connection of microorganisms present on nearby surfaces and its relation to the contamination of the patient. An 80% growth of Gram positive bacteria was observed in the analyzed blood cultures, with a prevalence of S. epidermidis (40%).

Similar data were observed in work carried out by Alves et al. (2012) involving 170 samples of blood cultures, which found 45.5% of S. epidermidis. In addition to this, there was a prevalence of 20% of S. haemolyticus, which also plays an important role in opportunistic infections. Infections caused by this species have been considered
as a threat, since the mechanisms involved in biofilm formation between them, are not yet fully elucidated.15,17

The *S. epidermidis* and *S. haemolyticus* are the two species most frequently reported among the isolated CoNS in cases of nosocomial infections, according to several studies.14,20,21

Among the microorganisms identified as resistant to oxacillin, 12.5% of the species were similar in the bed surface samples and blood cultures from the same patient sample. The identification of microorganisms in hospital surfaces has been important in epidemiological investigations and suggest the environment as a possible source of transmission of nosocomial infections.15,17

One of the greatest ways to spread of hospital infections occur through cross-infection, which is caused by the transmission of microorganisms between patients or through the hands of professionals, caregivers and visitors, and health professionals the main contributors to this spread. The acquisition of microorganisms may also occur through direct contact with the patient material or contaminated environment.22,23 The low adherence of hygiene, by some health professionals, contributes to a significant increase of colonized patients, and consequently increasing the degree of environmental contamination. The aspects cleaning measures and disinfection of surfaces with handwashing contribute effectively as control measures in cases of cross-infections by minimizing the rise of environmental reservoirs, which may be considered as potential sources of infection to the patient.22

Because of the problem of global patient safety, the World Health Organization (WHO) established in 2004 the World Alliance for Patient Safety (World Alliance for Patient Safety) which seeks to define priority issues for research in the area of patient safety that are highly relevant.24 Among these, they highlight skills and abilities, often inadequate among health professionals and infections associated with health care.24,25

The similarity of microorganisms found between environmental samples and blood cultures in this study highlights the interference of the hospital in the transmission of pathogens and possible cross-infection. Resistant Gram positive bacteria identified reinforce the importance of emphasizing the general measures of environmental hygiene in relation to patient safety in health care. These include greater attention as handwashing, personal protective equipment for use, application of contact precautions, isolation area of establishment of patients with multidrug-resistant bacteria, as well as the strengthening of the institutional policy of rational use of antimicrobials.

REFERENCES


