Identification and antimicrobial susceptibility of *Staphylococcus* from home-treated peritoneal dialysis patients

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**ABSTRACT:** *Staphylococcus aureus* is the main agent of infections during peritoneal dialysis (PD). The presence of *S. aureus* in the nasal cavity has been extensively studied and suggested as a risk factor of dialysis-related infections, whereas coagulase-negative *Staphylococcus* (CNS) species are frequently considered part of the normal human microbiota. The aim of this study was to identify *Staphylococcus* in the nasal cavity, pericatheter skin and peritoneal effluent from PD patients, as well as to evaluate the antimicrobial activity evolution *in vitro*. Thirty-two chronic PD patients were observed during 12 months and had nasal and pericatheter skin samples collected for culture. When peritonitis was detected, samples were also collected from the peritoneal effluent for culture. The activity of several antimicrobial drugs (penicillin G, oxacillin, cephalothin, ofloxacin, netilmicin and vancomycin) against different *Staphylococcus* species was measured by using the agar drug diffusion assay (Kirby-Bauer method). *Staphylococcus* was separated into *S. aureus*, *S. epidermidis* and other CNS species in order to determine the *in vitro* resistance level. *S. epidermidis* resistance to oxacillin progressively increased during the study period ($p < 0.05$). Resistance to ofloxacin was inexpressive, whereas resistance to netilmicin and vancomycin was not detected. Of the oxacillin-resistant species ($n = 74$), 83% were *S. epidermidis*, 13% other CNS and 4% *S. aureus* ($p < 0.05$). Regarding multi-drug resistant strains ($n = 45$), 82% were *S. epidermidis*, 13% other CNS, and 5% *S. aureus* ($p < 0.05$). This study shows the relevance of resistance to oxacillin and CNS multi-drug resistance, particularly concerning *S. epidermidis*, in PD patients.

**KEY WORDS:** peritoneal dialysis (PD), peritonitis, risk factor, *Staphylococcus*, coagulase-negative *Staphylococcus* (CNS).

**CONFLICTS OF INTEREST:** There is no conflict.

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INTRODUCTION

*Staphylococcus* species represent the most frequent cause of infection in home peritoneal dialysis (PD). In Brazil, during the two first decades of home PD, *S. aureus* was considered the main cause of peritonitis (1-4). However, recent Brazilian data, similarly to those reported in the literature, have demonstrated that coagulase-negative *Staphylococcus* (CNS) species are the main etiological agents of these infections (4-7).

*Staphylococcus* is subdivided into forty species and most of them are coagulase-negative. Nevertheless, the capacity of enzyme synthesis is reserved to *S. aureus*, *S. schleiferi* subspecies *coagulans*, *S. intermedius*, *S. hyicus* and *S. delphini* (8, 9).

Nasal carriers of *S. aureus* have been extensively studied by several authors and the presence of this bacterium in the nasal cavity has been pointed out as a major risk factor of dialysis-related infections, although a reduction in peritonitis has been reported (10-18). CNS is considered part of the normal microbiota in humans; still, it is recognized as a cause of infections of variable severity. Tzamaloukas (19) reported that CNS infections evolved without severe consequences, presenting benign characteristics, whereas other authors have described a different bacterial profile, observing significant levels of recurrent peritonitis (20, 21).

According to Berns (22), the behavior of CNS strains can be similar to that of *S. aureus*, i.e. resistant to methicillin and other β-lactam antimicrobials such as cephalosporins and carbapenems. Therefore, detailed microbiological studies on these species are of high importance for the prognosis and treatment of peritonitis. The present investigation also aimed to identify *Staphylococcus* species and characterize the evolution of its resistance to oxacillin and other drugs in nasal cavity, pericatheter skin and peritoneal effluent samples from patients undergoing PD.

PATIENTS AND METHODS

This study included patients undergoing peritoneal dialysis (manual or automated) and was conducted from January 2003 to July 2004 in the Botucatu Medical School, São Paulo State University (UNESP – Univ Estadual Paulista), Brazil. Patients without episodes of associated peritonitis in the previous three months and not receiving antibiotics in the previous month were included. Patients who did not follow the protocol were excluded from the trial to avoid lack of feedback. The included patients were followed for a minimum period of 12 months and information about
their age, gender, race, diabetes, PD and treatment length was recorded. Sterile swabs were used to collect nasal and pericatheter skin samples. Samplings were done at the beginning of the trial, after two months and between nine and 12 months after the first collection.

Peritonitis was defined as the emergence of unclear peritoneal effluent, followed or not by abdominal pain, painful and abrupt abdominal decompression, nausea, vomiting and fever. The infection was confirmed by a leukocyte count in dialysis effluent greater than 100 cells/mm$^3$, in the presence of polymorphonuclear cells. Cultures were prepared by aseptically collecting 40 mL of peritoneal effluent from the last drainage. After centrifugation, the sediment was inoculated into BACTEC® bottles (BD, USA). For microbial identification, colonies were gram-stained. *Staphylococcus* was differentiated from *Micrococcus* throughout glucose oxidation and fermentation, bacitracin resistance (0.04 U) and sensitivity to furazolidone (100 μg). Coagulase test was employed to differentiate *S. aureus* from CNS (23). The latter was identified by using the simplified method proposed by Cunha *et al.* (24) through analysis of sugar utilization, lactose, anaerobic growth in thioglycollate broth, nitrate reduction, hemolysins, urease and ornithine decarboxylase.

Sensitivity test to antimicrobial drugs was carried out by means of agar drug diffusion assay (Kirby-Bauer method), as recommended by the Clinical Laboratory Standard Institute (CLSI) (25). The employed disks were: penicillin G (10 UI), oxacillin (1 μg), cephalothin (30 μg), netilmicin (30 μg), ofloxacin (5 μg) and vancomycin (30 μg) purchased from CECON (Brazil). Sensitivity, drug resistance and control test based on international reference strains (ATCC 25923) followed the CLSI recommendations (25).

Evaluations were performed by separating the strains from the different sites into three groups: *S. aureus*, *S. epidermidis* and other CNS. We verified the proportion of strains resistant to six drugs (in three evolutional samples), the resistance to oxacillin in all three groups of *Staphylococcus*, and the frequency of multi-drug resistant strains (microbial resistance to more than one tested drug). Considering that six drugs were tested, resistance to oxacillin and cephalothin was interpreted as a single resistance.
Statistical analysis
To compare frequencies, chi-square test was employed. For the tested hypothesis, \( p < 0.05 \) was considered significant.

RESULTS
The characteristics of the 32 patients studied are shown in Table 1. Their mean age was \( 54.7 \pm 17 \) years (between 21 and 85 years); 62.5% patients (\( n = 20 \)) were male; 40.6% (\( n = 13 \)) diabetic; white color prevailed among 84.4% (\( n = 27 \)) and manual dialysis among 75% patients. Dialysis mean length was \( 34.6 \pm 20 \) months.

Ninety-six nasal and pericatheter skin samples were collected from thirty-two PD patients; when peritonitis was diagnosed, peritoneal effluent samples were also obtained. The research resulted in the identification of 288 \textit{Staphylococcus} strains, 155 from the nasal cavity, 122 from skin and 11 from the peritoneal effluent. The frequency of identified species was 21.9% \textit{S. aureus} and 78.1% CNS (Table 2). In the nasal cavity, CNS species were predominant, representing 78.7%, with 50.3% \textit{S. epidermidis}, 13.5% \textit{S. warneri} and 7.1% \textit{S. haemolyticus}. The isolated \textit{S. aureus} strains accounted for 21.3%. In the pericatheter skin, the distribution was similar; 22.1% \textit{S. aureus} and 77.9% CNS were isolated, while the most frequent were: 50.0% \textit{S. epidermidis}, 10.6% \textit{S. haemolyticus} and 9.0% \textit{S. warneri}. In the 11 cases of peritonitis, the distribution percentage of \textit{Staphylococcus} species was similar to that found in the extra-peritoneal sites; 72.7% CNS were isolated, 54.5% were \textit{S. epidermidis} and 18.2% were \textit{S. warneri}. In addition, 27.3% of the peritonitis cases were caused by \textit{S. aureus}. 

Table 1. Characteristics of the 32 patients subjected to peritoneal dialysis at home

<table>
<thead>
<tr>
<th>Characteristics</th>
<th>Number</th>
<th>Percentage (%)</th>
</tr>
</thead>
<tbody>
<tr>
<td>Age Average in years</td>
<td>54.7 ±17</td>
<td></td>
</tr>
<tr>
<td>Gender</td>
<td></td>
<td></td>
</tr>
<tr>
<td>Male</td>
<td>20</td>
<td>62.5</td>
</tr>
<tr>
<td>Female</td>
<td>12</td>
<td>37.5</td>
</tr>
<tr>
<td>Color</td>
<td></td>
<td></td>
</tr>
<tr>
<td>White</td>
<td>27</td>
<td>84.4*</td>
</tr>
<tr>
<td>Nonwhite</td>
<td>5</td>
<td>15.6</td>
</tr>
<tr>
<td>Diabetes mellitus</td>
<td></td>
<td></td>
</tr>
<tr>
<td>Yes</td>
<td>13</td>
<td>40.6</td>
</tr>
<tr>
<td>No</td>
<td>19</td>
<td>59.4</td>
</tr>
<tr>
<td>Dialysis method</td>
<td></td>
<td></td>
</tr>
<tr>
<td>Manual</td>
<td>24</td>
<td>75*</td>
</tr>
<tr>
<td>Automated</td>
<td>8</td>
<td>25</td>
</tr>
</tbody>
</table>

*p < 0.05.

Table 2. Distribution of *Staphylococcus* isolated from the nasal cavity, pericatheter skin and peritoneal effluent

<table>
<thead>
<tr>
<th>Species</th>
<th>Sampling site</th>
<th>nasal cavity % (n)</th>
<th>Pericatheter skin % (n)</th>
<th>Peritoneal effluent % (n)</th>
<th>Total</th>
</tr>
</thead>
<tbody>
<tr>
<td>S. aureus</td>
<td></td>
<td>21.3 (33)</td>
<td>22.1 (27)</td>
<td>27.3 (3)</td>
<td>63</td>
</tr>
<tr>
<td>S. epidermidis</td>
<td></td>
<td>50.3 (78)</td>
<td>50.0 (61)</td>
<td>54.5 (6)</td>
<td>145</td>
</tr>
<tr>
<td>S. warneri</td>
<td></td>
<td>13.5 (21)</td>
<td>9.0 (11)</td>
<td>18.2 (2)</td>
<td>34</td>
</tr>
<tr>
<td>S. haemolyticus</td>
<td></td>
<td>7.1 (11)</td>
<td>10.6 (13)</td>
<td>0</td>
<td>24</td>
</tr>
<tr>
<td>S. saprophyticus</td>
<td></td>
<td>2.6 (4)</td>
<td>1.6 (2)</td>
<td>0</td>
<td>6</td>
</tr>
<tr>
<td>S. capitis</td>
<td></td>
<td>1.9 (3)</td>
<td>1.6 (2)</td>
<td>0</td>
<td>5</td>
</tr>
<tr>
<td>S. simulans</td>
<td></td>
<td>1.9 (3)</td>
<td>1.6 (2)</td>
<td>0</td>
<td>5</td>
</tr>
<tr>
<td>S. lugdunensis</td>
<td></td>
<td>0.6 (1)</td>
<td>1.6 (2)</td>
<td>0</td>
<td>3</td>
</tr>
<tr>
<td>S. hominis</td>
<td></td>
<td>0</td>
<td>1.6 (2)</td>
<td>0</td>
<td>2</td>
</tr>
<tr>
<td>S. schleiferi</td>
<td></td>
<td>0.6 (1)</td>
<td>0</td>
<td>0</td>
<td>1</td>
</tr>
<tr>
<td>Total</td>
<td></td>
<td>155</td>
<td>122</td>
<td>11</td>
<td>288</td>
</tr>
</tbody>
</table>

The in vitro antimicrobial resistance profile of *Staphylococcus* strains to the six studied drugs was done for all samples obtained in the three collections. The percentage of penicillin G-resistant strains did not vary during the collection, including around 80% S. epidermidis, 60% other CNS and 70% S. aureus. As to the in vitro activity of oxacillin (Figure 1), resistant S. epidermidis strains accounted for 17.3% (n
= 9) in the first collection, 26.5% in the second and 44.7% (n = 17) in the third sampling. The increasing rates of oxacillin resistance were statistically significant (p < 0.05); for other Staphylococcus species, resistance results did not differ. The in vitro resistance to cephalothin was inexpressive among Staphylococcus species: 1.9% (n = 1) for S. epidermidis, 10% (n = 2) for S. aureus, and no resistant for other CNS strains. In vitro resistance to ofloxacin during the studied period was 18% for S. epidermidis, 12% for other CNS and 7% for S. aureus. No Staphylococcus sample showed in vitro resistance to netilmicin and vancomycin.

![Graph](Image)

**Figure 1.** In vitro resistance profile of S. aureus, S. epidermidis and other CNS strains to oxacillin in three sample collections from 32 patients subjected to peritoneal dialysis at home.

The distribution of Staphylococcus species resistant to oxacillin (n = 47) or presenting a multi-drug resistance pattern (n = 45) is shown in Figure 2. Concerning strains resistant to oxacillin, 83% (n = 39) were S. epidermidis, 13% (n = 6) other CNS and 4% (n = 2) S. aureus (p < 0.05). Multi-drug resistance was found among 82% S. epidermidis strains, 13% (n = 6) other CNS and 5% (n = 2) S. aureus (p < 0.05).
The pathogens most commonly associated with peritonitis in PD patients are *Staphylococcus* species. The presence of *S. aureus* in nasal cavity and pericatheter skin represents a risk factor for the development of peritonitis (14, 16, 26). However, in CNS-related peritonitis, the relationship between the presence of these microorganisms in the patients’ skin and nasal cavity and the development of subsequent infection is not discussed since the related species are understood to be part of the normal microbiota.

In this investigation, we found a wide variety of *Staphylococcus* species in the nasal cavity, pericatheter skin and peritoneal effluent samples. The most frequent *Staphylococcus* species were *S. epidermidis*, *S. aureus*, *S. warneri* and *S. haemolyticus*. Due to the lack of CNS species identification in routine patients, few studies present data related to CNS in PD patients.

Eisenberg et al. (6) studied the catheter skin and nasal cavity of PD patients and reported the importance of CNS colonization, including the relevance of *S. epidermidis* as a cause of peritonitis. In the present study, similar results were found for CNS species in three collections with serial cultures. The proportional distribution
of species among different anatomic sites suggests that there is bacterial propagation from the natural habitat to the peritoneal effluent. Considering the in vitro antimicrobial activity of oxacillin, S. epidermidis had increasing resistance during the observation period. This bacterium also presented the highest frequency of strains resistant to oxacillin, as well as multi-resistance against two or more drugs, possibly due to its mechanism of acquiring resistance from another agent or due to the in vitro selection of resistant strains (22). Some authors point out that this microorganism is the reservoir of various resistance genes, since it is the main component of skin and mucosal microbiota and is, therefore, subjected to a more selective pressure by commonly used antimicrobials. These results show that, in addition to the epidemiological importance, the identification of Staphylococcus species contributes to information on their correlation to the pathogenic potential and the sensitivity profile of antimicrobials (6).

The in vitro antimicrobial activity of cephalothin resulted in few resistant strains in all three samplings and, according to CLSI (25), Staphylococcus strains resistant to methicillin and oxacillin are also resistant to cephalosporin. Thus, their sensitivity must not be assessed in vitro, since laboratory tests not always represent the interaction between the pathogen and the antibiotic in vivo. Zelenitsky et al. (27) reported limitations of in vitro sensitivity tests for peritonitis treatment in PD, since the tolerance to the antibiotic by the bacteria isolated from the peritoneal effluent differed from that obtained in standard medium, suggesting that growth conditions are important factors in the acquisition of bacterial tolerance.

Nouwen et al. (16) showed that persistent S. aureus nasal carriers did not present resistance to vancomycin, and 2% of the CNS strains isolated from the same site revealed intermediate resistance to that drug.

The results obtained in the present study reinforce the importance of recognizing endogenous sources of different Staphylococcus species and their in vitro susceptibility in the prognosis and treatment of peritonitis (6, 16). Although the isolation of CNS species by means of molecular techniques is complex, it will establish colonization patterns and allow advances in the prevention of infections caused by these species.
REFERENCES

