**Vancomycin resistant Enterococcus spp (VRE): Follow up during 9 years in a Tertiary Teaching Hospital in Southern Brazil**

Ana Maria Sandri¹, Geórgia Lopes da Silva², Silvia Pedroso Tavares Soares², Fabiano Ramos¹², Luciana Ruschel de Alcântara³, Larissa Lutz⁴, Afonso Luís Barth⁵

**ABSTRACT**

**Introduction:** Infection with vancomycin-resistant Enterococcus spp (VRE) has been a worldwide problem since mid 1980’s and, in Brazil, since 1996. This study was conducted to evaluate the experience with VRE in our institution.

**Methods:** A prospective cohort study from 2000 to 2009 was conducted at Hospital Sã Loucas da PUCRS. All hospitalized patients with VRE positive culture were included and followed from their diagnosis until they were negative for VRE or their discharge. Only the first admission for each VRE positive patient was included. Pulsed field gel electrophoresis (PFGE) was performed to determine how VRE had spread.

**Results:** A total of 315 cases of VRE were identified, 224 of which were isolated from rectal swabs. Vancomycin-resistant/ampicillin susceptible Enterococcus faecalis were identified in 312 isolates. PFGE was performed in 47 VRE isolates that presented an indistinguishable migratory profile. The median length of hospital stay and length of stay before VRE isolation were 46 days and 21 days, respectively; 52% of the patients were aged 60 and above. The annual distribution of the new VRE cases showed a clear decrease from 2000 to 2009.

**Discussion:** This study shows a substantial VRE colonization (71%) with a homogenous pattern that emphasizes its transversal spread. Predominance of E. faecalis differs from the literature which largely describes a higher prevalence of vancomycin-resistant Enterococcus faecium. The follow up of VRE during 9 years in our institution highlighted the importance of continuous surveillance to prevent outbreaks in our hospital.

**Keywords:** Vancomycin-resistant Enterococcus, VRE, outbreak, surveillance.

Enterococci are normal inhabitants of the intestinal tract of humans and animals¹. Their capacity to colonize the host for long periods of time, to survive in inanimate objects, and to be resistant to many commonly used antibiotics ensures their success as a nosocomial pathogen¹-⁵. Enterococcus spp can cause serious infections such as bacteremia, urinary tract infection, endocarditis, and intra-abdominal infection, mainly in immunocompromised patients². Among the Enterococcus species, Enterococcus faecalis and E. faecium are the most common, corresponding to 80-90% and 5-10% of clinical isolates, respectively³,⁶. This proportion has shifted somewhat as E. faecium has become increasingly resistant to antimicrobial agents and emerged as a major nosocomial pathogen⁶,⁷.

Infection with vancomycin-resistant Enterococcus (VRE) has been an increasing problem worldwide since the first reports in France and the United Kingdom in 1988⁸,⁹. VRE had spread rapidly throughout the world, particularly in the United States where 30% of the enterococci isolated from intensive...
care units (ICUs) are resistant to vancomycin\textsuperscript{10-12}. In Europe, infections caused by VRE are also a problem and correspond to 7\% of the enterococcal isolates from blood culture; VRE are also important in the community and there is a relatively large reservoir of VRE that had been linked to the use of avoparcin in livestock\textsuperscript{13-15}. In Brazil, the first case of VRE was reported in 1996, in Curitiba, and new cases were reported in the following year in the city of São Paulo and other cities\textsuperscript{16,17}.

There are many risk factors for acquiring VRE: previous use of antibiotics, prolonged hospitalization, long ICU stay, immunosuppression, and abdominal or thoracic surgery\textsuperscript{18,19}. The spread of VRE in hospital settings has been widely reported and are usually due to: 1) patient-to-patient transmission, 2) hand contact of healthcare workers, or 3) contaminated environmental surfaces and medical equipment\textsuperscript{20-24}.

The aim of the present study was to evaluate our local experience with VRE in Hospital São Lucas da Pontifícia Universidade Católica do Rio Grande do Sul (HSL/PUCRS), located in southern Brazil.

METHODS

Patients and Setting

The study was conducted at HSL/PUCRS, a 600-bed tertiary care teaching institution located in the city of Porto Alegre, southern Brazil. The study population consisted of hospitalized patients with clinical or surveillance cultures positive for VRE. The study period was from May 2000, when VRE was isolated for the first time in our General Intensive Care Unit (GICU), through May 2009. Only one hospitalization per patient was considered. Outpatients were not included.

Study Design

We conducted a prospective cohort study including all hospitalized patients colonized and/or infected with VRE, who were followed up from the date of their diagnosis to the date they were negative for VRE or to their discharge from the hospital.

Bacterial Culture and Identification

Enterococci were obtained from clinical specimens or from rectal swabs during surveillance culture. Surveillance culture was performed once a week in patients admitted to the GICU, and at any time in patients who had shared a room with a VRE positive patient for more than 48 hours, according to the recommendations of the Infection Control Service (ICS) of our hospital.

The culture and identification of \textit{Enterococcus} spp was performed according to conventional tests. The susceptibility of enterococci to vancomycin was evaluated using the disk diffusion method and followed the Clinical and Laboratory Standards Institute\textsuperscript{25}. Results for isolates classified as resistant to vancomycin by the disk diffusion method were confirmed by the E-test method.

Molecular Typing

To determine how VRE had been spread, the genotypic profile of VRE isolates obtained from clinical specimens and rectal swabs was determined at the Biomedical Research Unit of the Clinical Pathology Service at Hospital de Clínicas de Porto Alegre (HCPA). Molecular typing was performed by DNA macrorestriction followed by pulsed field electrophoresis (PFGE) as previously described\textsuperscript{26}. Briefly, unshared DNA was extracted from the bacterial culture and digested with \textit{SmaI} for 20 hours in a 30°C water bath. After digestion, the DNA fragments were subjected to PFGE in a CHEF-DR\textregistered II apparatus (Bio-Rad Laboratories, California, USA). In order to evaluate the discriminatory power of PFGE, vancomycin-susceptible \textit{Enterococcus} spp (VSE) obtained from the same patients were also evaluated. The migratory profiles of the isolates were compared visually and analyzed according to the criteria of Tenover et al.\textsuperscript{27}

VRE Surveillance

VRE positive patients were followed up from their diagnosis until they had two negative sequential swabs and clinical specimens with one week apart each other, or until their discharge from the hospital.

In order to prevent further spread of VRE, the ICS of our hospital established specific contact control measures according to the Guideline for Isolation Precautions in Hospitals, 1995, of the United States Centers for Diseases Control\textsuperscript{28}. These precautions have been maintained from May 2000 to May 2009 and included active surveillance with weekly rectal swabs of GICU patients and control of all clinical materials from any unit of the hospital with identified cases of VRE. Positive patients, colonized and/or infected with VRE were subjected to weekly swabs and collection of clinical material; they were taken from the isolation when two subsequent swabs and clinical specimens were negative for VRE.

Once a patient was identified as being infected or colonized with VRE, he/she was put in a private room or in shared rooms with other VRE positive patients and advised not to visit common areas on the ward. Educational sessions were provided to all healthcare workers reinforcing contact precautions including hand hygiene, use of gloves and gowns,
individualized equipments for each room (thermometers, stethoscopes, sphygmomanometers); in addition, if the patient traveled to another part of the hospital, he/she must be conducted in wheel chairs by the health team appropriately worn with gown and glove. Environmental cleaning procedures included daily cleaning of patient rooms and “terminal cleaning” at patient discharge. A sodium hypochlorite-based disinfectant was used for cleaning (Virex). The correct use of vancomycin was emphasized.

Only one hospitalization and one positive culture (the first to appear) per patient were considered.

**Statistical Analysis**

The statistical analyses were performed using SPSS, version 10.0. Results from descriptive statistics were presented in tables and graphics.

**RESULTS**

From May 2000 to April 2009, 315 cases of VRE were identified, of which 224 were isolated from rectal swabs and 91 from other clinical samples, mainly blood, followed by urine (table 1).

Strains of vancomycin resistant and ampicillin susceptible *Enterococcus faecalis* were identified in 312 isolates, of vancomycin and ampicillin resistant *E. faecalis* in 2 isolates, and of vancomycin and ampicillin resistant *Enterococcus faecium* in 1 isolate. Molecular typing was done in 47 VRE isolates, 37 of which were from a rectal swab, 5 from secretion, 4 from blood, and 1 from urine. A well-defined DNA migratory profile could be observed for 46 VRE isolates, as only one did not migrate. The comparison of DNA profiles indicated that 39 isolates were indistinguishable, as there was no visible difference between the bands. This profile was designated profile “A”. Four isolates presented migratory profiles closely related to profile “A”, differing from it in only two bands (genotype A1, three isolates) and in three bands (genotype A2, one isolate). Three other isolates presented migratory profiles that were different from profile “A” by more than seven bands and were designated profiles “B”, “C” and “D”. The discriminatory power of PFGE was evaluated by comparing VRE and VSE isolates obtained from the same patients. VSE isolates presented totally different migratory profiles when compared to clone A.

The annual distribution of new VRE cases showed a clear decrease from 2000 to 2009 in Intensive Care Units (ICUs) and hospital wards (figure 1). The first cases identified at the beginning of 2000 were admitted to the GICU, but the outbreak extended to five different units: clinical and surgical unit, post-cardiac surgical ICU, coronary ICU, pediatric ICU and post-surgical recovery unit (table 2).

Table 1: VRE distribution according to the materials used for bacterial identification.

<table>
<thead>
<tr>
<th>Material</th>
<th>n</th>
<th>%</th>
</tr>
</thead>
<tbody>
<tr>
<td>Rectal swab</td>
<td>224</td>
<td>71.11</td>
</tr>
<tr>
<td>Blood culture</td>
<td>36</td>
<td>11.43</td>
</tr>
<tr>
<td>Urine culture</td>
<td>29</td>
<td>9.21</td>
</tr>
<tr>
<td>Others</td>
<td>26</td>
<td>8.25</td>
</tr>
<tr>
<td>Total</td>
<td>315</td>
<td>100.00</td>
</tr>
</tbody>
</table>

Figure 1: Distribution of the new VRE cases in the ICUs (straight line) and hospital wards (dotted line) from 2000 to 2009.
The median time of hospitalization was 46 days (1 to 393 days) and the median time of hospitalization before VRE was isolated for the first time was 21 days (1 to 366 days). People over 60 years were the most affected (table 3).

DISCUSSION

This study demonstrates substantial VRE colonization (71.1%) in patients admitted to our hospital as assessed by rectal swab; among the clinical materials used for bacterial identification, blood and urine were the most important. The indistinguishable profile observed for the vast majority of the VRE isolates in PFGE indicates a transversal environmental mode of spread. Our results are consistent with earlier studies that indicated clonal dissemination as a major mechanism for spreading of isolates20-22. However, the absolute predominance (99.68%) of \textit{E. faecalis} among VRE cases during the period of the study is not consistent with what has been reported in the literature, which largely describes a higher prevalence of vancomycin resistant \textit{E. faecium}29-32. Studies from Porto Alegre, Southern Brazil, also demonstrated the important occurrence of vancomycin-resistant \textit{E. faecium} during recent years33. The predominance of VRE in older people and in ICUs are also observed in literature. The incidence distribution curve for VRE in our hospital shows that it almost disappeared throughout the 9-year period observed. The implementation of specific and persistent infection control measures probably played a crucial role in these results as we can observe in many publications23,24,34-38.

In conclusion, following the VRE pattern across a 9-year period allowed us to learn how to deal with these resistant bacteria and to understand that its presence can be an excellent indication of the capacity of the unit to maintain the safety measures to avoid the cross transmission of the microorganisms. Continuous surveillance of VRE is mandatory due to its clinical and epidemiologic importance, in order to promptly implement specific control measures to prevent new outbreaks.

Table 2: VRE distribution according to the units of hospitalization.

<table>
<thead>
<tr>
<th>Unit</th>
<th>n</th>
<th>(%)</th>
</tr>
</thead>
<tbody>
<tr>
<td>GICU*</td>
<td>171</td>
<td>54.28</td>
</tr>
<tr>
<td>Clinical and surgical unit</td>
<td>113</td>
<td>35.87</td>
</tr>
<tr>
<td>Post-cardiac surgical ICU**</td>
<td>5</td>
<td>1.59</td>
</tr>
<tr>
<td>Coronary ICU</td>
<td>20</td>
<td>6.35</td>
</tr>
<tr>
<td>Pediatric ICU</td>
<td>3</td>
<td>0.95</td>
</tr>
<tr>
<td>Post-surgical unit</td>
<td>3</td>
<td>0.95</td>
</tr>
<tr>
<td>Total</td>
<td>315</td>
<td>100.00</td>
</tr>
</tbody>
</table>

*GICU= General Intensive Care Unit ; **ICU= Intensive Care Unit.

Table 3: VRE distribution according to patients’ age.

<table>
<thead>
<tr>
<th>Age</th>
<th>n</th>
<th>(%)</th>
</tr>
</thead>
<tbody>
<tr>
<td>0 ┤ 19</td>
<td>4.8</td>
<td></td>
</tr>
<tr>
<td>20 ┤ 39</td>
<td>17.8</td>
<td></td>
</tr>
<tr>
<td>40 ┤ 59</td>
<td>25.4</td>
<td></td>
</tr>
<tr>
<td>60 ┤ 164</td>
<td>52.0</td>
<td></td>
</tr>
<tr>
<td>Total</td>
<td>315</td>
<td>100.0</td>
</tr>
</tbody>
</table>

The median time of hospitalization was 46 days (1 to 393 days) and the median time of hospitalization before VRE was isolated for the first time was 21 days (1 to 366 days). People over 60 years were the most affected (table 3).

REFERENCES


20. Hayden MK, Blom DW, Lye EA, Moore CG, Weinstein RA. Risk of hand or glove contamination after contact with patients colonized with vancomycin-resistant enterococcus or the colonized patients’ environment. *Infect Control Hosp Epidemiol.* 2008;29(2):149-54. http://dx.doi.org/10.1086/524331. PMid:18179370


33. Resende M, Caierão J, Prates JG, Narvaez GA, Dias CAG, d’Azevedo PA. Emergence of vanA vancomycin-


Received: 14/10/14
Accepted: 11/12/14