کارگاه های آموزشی مرکز اطلاعات علمی جهاد دانشگاهی

کارگاه آنلاین کاربرد نرم افزار SPSS در پژوهش

کارگاه آنلاین اصول تنظیم قراردادها

کارگاه آنلاین پروپوزال نویسی
Evaluation of a PCR Assay to Detect *Enterococcus faecalis* in Blood and Determine Glycopeptides Resistance Genes: Van A and Van B

Hamidreza Honarmand¹, Mahsome Falah Ghavidel², Iraj Nikokar³, Morteza Rahbar Taromsari⁴

**Abstract**

**Background:** Bacteremia due to *Enterococcus faecalis* is usually caused by strains resistant to most antibiotics. Effective management of the disease is dependent on rapid detection and characterization of the bacteria, and determination its sensitivity pattern to antimicrobial drugs. The aim of this study was to investigate a more rapid and reliable assay for simultaneous diagnosis of enterococcal bacteremia and its sensitivity pattern to antimicrobial drugs.

**Methods:** Several bacterial suspensions with different content of two standard strains of *Enterococcus faecalis* resistant to vancomycin were used for inoculation to defibrinated sheep blood samples. PCR and routine assay was performed on all blood samples with different bacterial content.

**Results:** Routine assay and PCR for all inoculated blood samples with ≥5 cfu/ml was positive. Mean time for PCR and routine assays was 10 hours and 5 days, respectively.

**Conclusion:** PCR is a more rapid and sensitive assay for simultaneous detection and characterization for *Enterococcus faecalis*, and determination of its sensitivity pattern to vancomycin.


**Keywords**

- *Enterococcus faecalis*
- multiplex-PCR
- Van A
- Van B

**Introduction**

*Enterococcus faecalis* is the cause of 85-90% of enterococcal, and third cause of nosocomial infections, especially bacteremia, sepsis in children, endocarditis, urinary tract infection (UTI), and wound infections.¹² It plays a significant role in treatment of the disease.³⁴ Knowledge of bacterial resistance pattern to antimicrobial agents is important for the successful management of diseases.³ Most hospital isolates are resistant to most usual antibiotics including vancomycin.⁶⁻⁸ There are five resistance genes whose products are responsible for resistance to glycopeptides antibiotics in vancomycin-resistant enterococci strains (VRE). Two of such genes (Van A and Van B) are most common than others, especially in *E. faecalis* and *E. faecium*.⁶⁻⁹ Strains with Van A gene are resistant to vancomycin and tycoplanin, and strains with Van B are resistant to vancomycin but sensitive to tycoplanin.¹⁰⁻¹¹ Resistant enterococcal infections are usually
treated by synergistic action of a glycopeptide and an aminoglycoside.\textsuperscript{1,5} Vancomycin-resistant enterococci strains are usually transferred via the hands of health care workers, who are fecal carrier and are in close contact with patients. Those patients who have long-time hospital stay and long-time antibiotic therapy, as well as children and the elderly with a critical situation, such as those who are hospitalized in intensive care units (ICU), are more prone to take the disease.\textsuperscript{12-14} Culture is the most-used way for detecting enterococci in the blood,\textsuperscript{2} however, for effective treatment of enterococcal bacteremia, characterization of the bacteria and their pattern of resistance to antibiotics is necessary. This requires some diagnostic biological tests as well as determination of its antibiogram pattern and MIC, which usually takes about five days.\textsuperscript{14-17} Some rapid methods such as E test for MIC, API 20 and API 32 for characterization, and selective-differential specific media and chormogenic agars with spectrophotometer. Then, diluted solutions with some single colonies, which were grown on TSA at 37°C for 24 hours, passage to TSA and incubation in 37°C for more 24 hours, identifying by catalase test, PYR test, growth on TSA with 6.5% NaCl, and hydrolysis esculin in the presence of bile on BEA. Differentiation of \textit{E. faecalis} from \textit{E. faecium} was done by three tests including ability to use pyruvate, fermentation of sorbitol, and reduction of tellurite.\textsuperscript{15,24} For screening VRE, we used BEA including 6 µg/ml vancomycin.\textsuperscript{10,15,24}

The extraction of DNA was achieved using the following procedure. Transferring 100 µl of each blood sample to a 2 ml ependorf vial contain 400 µl sterile double distilled H\textsubscript{2}O and incubation in 37°C for 30 minutes, adding 500 µl red cell lysis buffer (N\textsubscript{A}H\textsubscript{CO}_3 10 mM, NH\textsubscript{4}Cl 0.155M, pH=7) and incubation at 37°C for one hour, centrifugation at 10,000 rpm for 15 minutes, discarding the supernatant, adding 200 µl lysis buffer for bacteria (Tris 10 mM, sucrose 0.3 M, MgCl\textsubscript{2} 5 mM) to the pellet with 10 µl lysozyme (0.1 mg/ml, Sinagen, Lot: MR7735) and incubation at 37°C for one hour, adding 4 µl proteinase K (900 u/ml, Fermentaze, Lot: 00022411) and incubation at 65°C for one hour, extraction of DNA by standard phenol-chloroform method and precipitation of DNA by cold isopropanol. PCR mix was prepared as 3 µl 10x PCR buffer, 2 µl of MgCl\textsubscript{2} (25 mM), 0.5 µl of dNTP 10 mM, 100 pM of each primer, 0.2 µl DNA pol (5 u/µl), 2 µl DNA, and double distilled H\textsubscript{2}O to final volume of 25 µl. Special features of primers that were used in this study are shown in table 1.

PCR programs was adjusted as one cycle at 94°C for 7 minutes, 34 cycle at 94°C for 40 seconds, 46°C for 40 seconds, 72°C for 50

**Material and Methods**

We used a standard strain VRE (PTCC 1447, and PTCC 1237) prepared by the division of Bacteria and Fungi Collections, Iranian Institute of Industrial and Scientific Researches, Tehran, Iran). A suspension 10\textsuperscript{8} cfu/ml was made in normal saline by adding some single colonies, which were grown on TSA by adjusting its optical density to half McFarland solution and checking their absorbance in 700 nm with spectrophotometer. Then, diluted solutions with different bacterial contents (10\textsuperscript{4} cfu/ml, 10\textsuperscript{5} cfu/ml and 10\textsuperscript{2} cfu/ml) were made by diluting it in normal saline. They are used for inoculating to blood. By adding certain amount of each bacterial solution to certain amount of defibrinated sheep blood, some blood samples with different bacterial content (10\textsuperscript{4} cfu/ml, 10\textsuperscript{5} cfu/ml, 10\textsuperscript{2} cfu/ml, 10\textsuperscript{1} cfu/ml, 5 cfu/ml and zero as control) were prepared. Ten-milliliter-samples of each dilution were prepared to be used in ten experiments of each of the PCR and routine assays. For routine assay, we used initial enrichment procedure for each specimen by inoculating to TSB and incubation at 37°C for 24 hours, passage to TSA and incubation in 37°C for more 24 hours, identifying by catalase test, PYR test, growth on TSA with 6.5% NaCl, and hydrolysis esculin in the presence of bile on BEA. Differentiation of \textit{E. faecalis} from \textit{E. faecium} was done by three tests including ability to use pyruvate, fermentation of sorbitol, and reduction of tellurite.\textsuperscript{15,24} For screening VRE, we used BEA including 6 µg/ml vancomycin.\textsuperscript{10,15,24}

<table>
<thead>
<tr>
<th>Gene (16S rRNA)</th>
<th>Name</th>
<th>Sequence</th>
<th>Size of PCR product (bp)</th>
<th>Reference\textsuperscript{3}</th>
</tr>
</thead>
<tbody>
<tr>
<td>C1</td>
<td>5-GGATTAGATACCTGGTAGTGTC-3</td>
<td>320</td>
<td>25</td>
<td></td>
</tr>
<tr>
<td>C2</td>
<td>5-TGATGGACGCTGTAACACC-3</td>
<td></td>
<td></td>
<td></td>
</tr>
<tr>
<td>D1</td>
<td>5-ATC AAG TAC AGT TAG TCT TTA G-3</td>
<td>941</td>
<td>19</td>
<td></td>
</tr>
<tr>
<td>D2</td>
<td>5-ACG ATT CAA AGC TAA CTG AAT CAG T-3</td>
<td></td>
<td></td>
<td></td>
</tr>
<tr>
<td>A1</td>
<td>5-GGG AAA ACG ACA ATG GC-3</td>
<td>732</td>
<td>19</td>
<td></td>
</tr>
<tr>
<td>A2</td>
<td>5-GTA CAA TGT GGC CTA TG T-3</td>
<td></td>
<td></td>
<td></td>
</tr>
<tr>
<td>B1</td>
<td>5-ATG GGA AGC CGA TAG TC-3</td>
<td>635</td>
<td>19</td>
<td></td>
</tr>
<tr>
<td>B2</td>
<td>5-GAT TTC GTT CCT CGA CC-3</td>
<td></td>
<td></td>
<td></td>
</tr>
</tbody>
</table>

\textsuperscript{1}The gene of each primer; \textsuperscript{2}Name of primers; \textsuperscript{3}The reference for the primer design
seconds, and final extension at 72°C for 10 minutes. Electrophoresis of PCR product was done on 1.5% agarose gel including Etidium bromide by 100 volts for one hour.

Results

All blood samples with different bacterial content of 5 cfu/ml were positive in the routine assay and PCR (figure 1). So, the sensitivity of PCR was the same as that of routine test.

PCR with species specific primers (D1 and D2) on two blood samples with different bacterial contents of *E. faecalis* (ptcc 1447) is shown in figure 2. The routine assay needed five days, but the PCR assay needed 10 hours. The sensitivity of the test was 95.4% and the specificity was 100%.

Discussion

Rapid diagnosis is very critical in the treatment of bacteremia. Routine assay is time-consuming and expensive, and commercial automatic screening tests and disc diffusion agar are not efficient for highly resistant bacteria that make most of the hospital isolates. Identification by API 20 and API 32 was associated with different sensitivity and specificity. The most-studied selective-differential medias are EVA, CAN-VGA, and BEAA with 60 µg/ml vancomycin. Although is more specific than CAN-VGA, EVA is slower (24 vs 48 hours). Several investigators have used multiplex-PCR with multiple pairs of specific primers. Dutka-Malen used six primer pairs to detect a number of standard strains and clinical isolates. Patel studied 100 clinical isolates (34 *E. faecalis*) using four primer pairs. The multiplex-PCR assay that with 4 primer pairs that Stake used for screening many clinical isolates, had a sensitivity of 85.0% and a specificity of 100% specificity, but the one that with three primers that Jayartne used for screening of 657 isolates had a sensitivity of 95.4% and a specificity of 99.8%. Ke, Ke use primers designed from tuf enterococcal gene to diagnose 14 of 20 enterococcal species. Angelleti, used four pair of primers to detecting 279 isolates, and found that it was more rapid than routine assay. It is in
agreement with our study. Paul,30 used two pairs of specific primers for Van A and Van B to screen clinical isolates, and found it was more sensitive than culture methods. In our study, the sensitivity of PCR was the same as that of routine test. It seems that the quality and quantity of DNA, which is related to DNA extraction method, is critical to the higher sensitivity. Kariyama,31 used Multiplex-PCR with seven pairs of primers to screen many clinical isolates and standard strains, and found that it was simpler and more efficient than the routine method. It is also compatible with our study.

False-positive cases is mainly the result of amplifying DNA of dead bacteria in the sample and amplifying resistance genes like Van A and Van B that are present in some other bacteria. This is critical for fecal samples that contain different bacteria, but not for blood ones.25 Regulating the concentrations of several primers in PCR mix is a technical problem for multiplex-PCR. Kariyama found that inhibition of Van A primers can be neutralized by increasing their concentrations to two-folds.31 Angeletti,29 and Stake,18 performed the multiplex-PCR in two steps, and in one step they only used Van A primers. We used one step and the same concentrations of primers, which seems to be the cause of weak view of bands (figure 3). The other cause of false-positivity is the specificity of primers. Primers designed by Ke,26 from tuf gene could amplify two species of Abiotrophia and four species of Lysteria. However, they are not the usual causes of bacteremia and this problem is important for fecal screening. It has been recommended to use molecular typing methods such as RFLP on PCR product,26 or very specific primers for E. Faecalis,19,27 for characterization. The present study used the latter method (figure 2). It has been recommended to use genus specific or universal primers as the internal control for detecting false-negative cases.27 However, we used species specific primers to diagnose E. Faecalis and as an internal control (figure 2). One of the main technical problems in the diagnosis of bacteremia by PCR is the obtainment of high quality and quantity bacterial DNA from the whole blood. This may not be easily possible because of high contents of PCR inhibitors. Therefore, DNA extraction method is critical.25,32 Zang,33 used Quiagen kit, and detected five cfu/ml bacteria in the blood.33 Newcomb,34 used Boom method and Klausegger,35 used DNA ZOl buffer for lysis bacteria in blood. However, Anthony,36 used double distilled H2O for lysis blood cell, proteinase K for eliminating PCR inhibitors, phenol-chloroform and alcohol precipitation to extract DNA of bacteria in blood, however, the quality of extracted DNAs was not very good. It seems that the poor quality of DNA was the main cause of weak bands of PCR products in our study (figure 1). We recommend a very efficient method or a commercial kit to increase the accuracy of PCR assays in such a case. We also recommend further studies to overcome other technical problems of PCR assay to make it more simple and reliable relative to new and very improved culture based methods such as VRE-BMX chromogenic commercial media.38 Such kits has several advantages such as differentiation at the species level, inhibition of growth of sensitive enterococci to vancomycin, promoting growth of resistant strains, and shorter duration than the traditional methods (24 vs 48 hours).

Conclusion

PCR is a more rapid and a sensitive method for simultaneous detection and characterization of E. Faecalis, and determination of its antimicrobial pattern to vancomycin. It can be considered as an alternative assay, but a more efficient DNA extraction method to increase the sensitivity of the assay is suggested.

Acknowledgment

We thank Mr. Mojtahab Hoseinpour for his kind help.

Conflict of Interest: None declared.

References


کارگاه های آموزشی مرکز اطلاعات علمی جهاد دانشگاهی

کاربرد نرم افزار SPSS در پژوهش
کارگاه آنلاین

اصول تنظیم قراردادها
کارگاه آنلاین

بروپوزال نویسی