

Biological changes of *Enterococcus faecalis* in the viable but nonculturable state

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ABSTRACT. Enterococcus faecalis may enter a viable but nonculturable (VBNC) state under adverse conditions. E. faecalis, the major bacterial species present in failed root canal treatments, is thought to survive after endodontic treatment by entering a VBNC state. In this study, we characterized the VBNC state of E. faecalis. We designed 3 different protocols to successfully induce the VBNC state. Approximately one-third of bacteria entered a VBNC state after 15-30 days, and all remained viable for at least 2 months. The morphology, glycometabolism, and adhesion capabilities of VBNC cells differed from those of E. faecalis during the exponential growth phase. Specifically, VBNC E. faecalis cells could not decompose lactose, D-mannitol, or D-sorbitol, although they were able to metabolize sucrose. Transmission electron microscopy showed that the morphology of the VBNC *E. faecalis* cells changed significantly; the cytoplasmic matrix was unevenly condensed and the overall morphology of the cells became irregular, but the cell membranes remained intact. Although the adhesion ability of the bacteria decreased, VBNC E. faecalis could still adhere to collagen fiber type

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I and tooth dentine. The persistence of this adhesion ability may be important in the virulence of VBNC *E. faecalis*.

Key words: Adhesion ability; *Enterococcus faecalis*; Morphology; Viable but nonculturable

INTRODUCTION

Root canal treatment (RCT), which is the classical treatment for endodontic infections, can seal and separate the root canal system from sources of bacterial infection. However, the failure rate of RCT is as high as 10-15% (Smith et al., 1993). Moreover, the risk of repeat infection in teeth with failed RCT is higher than in teeth with successful RCT (Molander et al., 1998; Sundqvist et al., 1998). Enterococcus faecalis constitutes a small percentage of the microbial species isolated from the root canals of teeth with necrotic dental pulps. However, E. faecalis is the most common species isolated from secondary endodontic infections. Polymerase chain reaction (PCR) detection identified E. faecalis in 82 and 76% of primary and secondary root canal infections, respectively. In contrast, E. faecalis was cultured from only 4% of untreated necrotic pulps (primary infections) and from 42% of root-treated canals (secondary infections) using traditional culture methods (Gomes et al., 2006). These findings demonstrate that E. faecalis is likely involved in the failure of endodontic treatment and that the bacteria in root-treated canals cannot be quantitatively detected using traditional culture methods. E. faecalis is the major species thought to cause endodontic treatment failure and refractory periapical periodontitis, and it is detected in 50-70% of these cases (Peciuliene et al., 2000; Pinheiro et al., 2003). The treated root canal is a unique niche that is low in nutrients and is comparatively well-sealed. E. faecalis can survive in this environment and induce root canal infection. The persistence of E. faecalis is thought to result from its ability to enter a viable but nonculturable (VBNC) state, which was first described for *E. faecalis* by Lleò et al. (1998), but the mechanism is unknown.

The VBNC state is a stress-response mechanism triggered by adverse growth conditions. When bacteria enter the VBNC state, they lose their ability to form colonies on culture media (Lleò et al., 1998), but continue to express antigen components (Smith and Oliver, 2006) and virulence factors (Pruzzo et al., 2002). VBNC bacteria may be able to resume cell division and promote disease transmission and persistence when permissive conditions are restored (Lleò et al., 2001). VBNC *E. faecalis* cells appear to maintain their viability, pathogenicity, and other biological characteristics such as gene expression (Lleò et al., 2000) and metabolic activity (Lleò et al., 1998). However, there are a number of differences in the biology, biochemistry, and genetics of VBNC cells and exponentially dividing cells, including changes in morphology, metabolism, and gene expression (Oliver, 2005) which help VBNC bacteria survive under adverse conditions; however, there are few reports in the current literature.

In this study, we successfully induced a VBNC state in *E. faecalis* cells using 3 different protocols and investigated the characteristics of VBNC cells, including their morphology, metabolic profile, and adherence properties.

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MATERIAL AND METHODS

Bacteriological strains and growth conditions

The *E. faecalis* laboratory strain ATCC33186 was grown in trypticase soy yeast extract broth. Cell growth was monitored using a spectrophotometer at a wavelength of 600 nm (OD_{600}) . Colony-forming unit (CFU) counts of culturable bacteria were performed by seeding samples in duplicate on trypticase soy yeast agar (TSA) plates.

Induction of VBNC state

Three different laboratory growth conditions were used, including an oligotrophic microenvironment that was prepared by adding sterile distilled water to a flask. The other 2 microenvironments consisted of alkaline medium prepared using sterilized sodium buffers at pH 9 and 11. For each condition, 600 mL *E. faecalis* culture in the mid-exponential growth phase (OD₆₀₀ \approx 0.7) (Pfeffer et al., 2006) was centrifuged at 4500 g for 15 min. The cell pellet was washed twice in phosphate-buffered saline. The washed cells were transferred to 3 500-mL flasks at a final density of 10⁷ cells/mL and incubated at 37°C. The cultures were monitored by CFU counts every 3 days from triplicate samples. A population was considered to be in the VBNC state when no colonies were detected after 10 mL cell culture seeded on solid culture medium.

Viability tests

The viability of the inoculated *E. faecalis* cells was tested by reverse transcription-PCR by using a primer pair specific for the *E. faecalis* penicillin-binding protein 5 gene, which encodes a protein involved in peptidoglycan synthesis (Robbi et al., 1996). Two primers within the gene (forward, 5'-CATGCGCAATTAATCGG-3'; reverse, 5'-CATAGCCTGTCGC AAAAC-3') were used. Extracted RNA was also used as template when performing the PCR procedure to rule out DNA contamination. The size of the expected amplification product was 444 bp. In addition, cell viability was measured using a Live/Dead kit (Invitrogen, Carlsbad, CA, USA). This kit utilizes a mixture of the green-fluorescent nucleic acid stain SYTO9 and the red-fluorescent nucleic acid stain, propidium iodide, to evaluate cell membrane integrity (Roth et al., 1997). When used alone, the SYTO9 stain generally labels all bacteria in a population, including those with intact membranes and those with damaged membranes. In contrast, propidium iodide penetrates only bacteria with damaged membranes, reducing the SYTO9 stain fluorescence when both dyes are present. Therefore, in this assay, live cells appeared to be green, while dead cells appeared to be red under fluorescent microscopy. Reverse transcription-PCR and fluorescent staining were performed every 10 days until no culturable cells were observed.

Resuscitation of VBNC E. faecalis

The nonculturable *E. faecalis* resumed division and were monitored on TSA plates after growth in trypticase soy yeast extract broth -0.5% glucose overnight at 37°C as previously described by Lleò et al. (2001).

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Acid production ability and biochemical identification

The pH value was monitored every 3 days after the bacteria were inoculated. Pure cultures and colonies in the mid-exponential phase *E. faecalis* (cultured for approximately 24 h, $OD_{600 \text{ nm}} \approx 0.7$) and VBNC *E. faecalis* cells were selected for biochemical identification using the Rapid ID 32 Strep (BioMérieux, Marcy l'Etoile, France). The ATB Miniapi Software (BioMérieux) was used to automatically read the ID 32 tests. Lactose, maltose, mannitol, sorbitol, and urease were used as biochemical substrates. The Rapid ID 32 Strep is a qualitative experiment rather than a quantitative test.

Transmission electron microscopy (TEM)

Cells in the VBNC state were centrifuged at 2500 g for 15 min and the cell pellets, which were approximately the size of a grain of rice, were fixed with 2% (v/v) glutaraldehyde for at least 2 h. The fixed pellets were washed twice with phosphate-buffered saline for 10 min each and post-fixed with 1% osmium tetraoxide for 1 h. The cells were washed twice with phosphate-buffered saline and dehydrated with successive ethanol washes (30-100%). Each dehydration step was performed for 10 min at 4°C. Two 10-min washes with propylene oxide were carried out, and the preparations were embedded in a mixture (v/v) of propylene oxide and embedded in 618-araldite epoxy resin for 2 h. The mixture was removed and replaced with 100% resin for 6 h. Resin polymerization was performed for 48 h at 60°C. Samples were prepared for TEM using an LKB ultramicrotome. Sections that were 50 nm in thickness were stained using lead citrate and visualized using a Philips CM-120 TEM (Amsterdam, Netherlands). Cells in the exponential growth phase were also analyzed to compare the differences between the 2 phases.

Scanning electron microscopy

The ability of cells to adhere to collagen or dentine was tested by adding bacteria in the VBNC state to dishes containing sterilized collagen fibers or pieces of dentine. The dishes were incubated in an anaerobic culture chamber at 37° C for 48 h. Cells in the exponential growth phase were also cultured as a positive control. We used shaker (180 rpm) during the incubation period to remove the influence of dead cells mixed with the VBNC cells. After incubation, the dishes were washed twice with sterile distilled water to remove nonadherent bacteria. The collagen fibers and dentine pieces were fixed with 2% (v/v) glutaraldehyde for 2 h and washed twice with sterile distilled water. Next, the collagen fibers and dentine pieces were snap-frozen at -80°C for 15 min. The samples were dried in a vacuum freeze dryer (Labconco, Kansas City, MO, USA), gold-coated, and examined by scanning electron microscopy (S-3000N; Hitachi, Tokyo, Japan).

RESULTS

Evaluation of time to enter VBNC state by *E. faecalis* cells in different microenvironments

The 3 different laboratory microenvironments were inoculated in an anaerobic envi-

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ronment at 37°C. Every 3 days, samples were withdrawn aseptically from the microenvironments and culturable cell numbers were evaluated by determining the CFU of diluted samples on TSA plates. When the CFU count was close to zero, 10-mL samples were filtered using 0.22-µm filters (Millipore, Billerica, MA, USA), and the filters were placed face-up on the TSA plates. The bacteria were considered to have reached the VBNC state when the number of culturable cells was less than 0.1 cells/mL.

As shown in Figure 1, the *E. faecalis* in group 1 became nonculturable within 30 days, but the same bacteria grown in the alkaline microenvironments required only 15 days to reach the nonculturable state. Live/dead staining showed that for the 3 tested conditions, 39, 36, and 22% of the bacteria were still alive after 1 month. In addition, reverse transcription-PCR demonstrated that cells cultured in the sterile distilled water expressed penicillin-binding protein 5 gene continuously for 3 months, but the cells cultured in the sodium buffer, pH 9 and 11, expressed penicillin-binding protein 5 gene for only 1.5 months.



Figure 1. Decline of *Enterococcus faecalis* when incubated at 37°C in sterile distilled water (squares), sterilized sodium buffer, pH 9 (triangles), and sodium buffer, pH 11 (lozenges). Each point is the mean of triplicate samples. The error bars were removed for clarity.

Acid production ability and glycometabolism observed in VBNC E. faecalis cells

The pH values of the samples were measured every 3 days from the beginning of the inoculation. The pH of the samples under all 3 conditions decreased from 8.05 to 6.06, 9.68 to 8.30, and 10.74 to 9.82 in the sterile distilled water and from pH 9 and 11 in sodium buffer conditions, respectively (Table 1). The decrease in pH was more evident in the distilled water than in the sodium buffer, indicating that the alkaline condition influenced the ability of the bacteria to produce acids although the alkaline condition was buffered. However, whether this change resulted from the alkaline condition requires further analysis.

Rapid ID 32 Strep tests were placed in a 37°C environment for 4 h, and the results were read automatically using the Mini API Software. The VBNC *E. faecalis* could metabolize sucrose similarly to the rate of exponentially growth *E. faecalis* cells, but the VBNC cells could not decompose lactose, D-mannitol, or D-sorbitol.

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Table 1. pH value of Enterococcus faecalis after incubation for 1 month.									
	0 day ¹	3 days	6 days	9 days	12 days	15 days	18 days	21 days	24 days
Sterilized distilled water	8.05	7.65	7.46	6.64	6.30	6.14	6.09	6.08	6.06
Sterilized sodium buffer group 1 Sterilized sodium buffer group 2	9.68 10.74	9.36 10.58	9.22 10.52	9.25 10.5	8.90 10.45	8.70 10.39	8.80 10.24	8.40 10.03	8.30 9.82

E. faecalis cells incubated in sterilized distilled water, sterilized sodium buffer, pH 9, and sodium buffer, pH 11.

Morphological changes in VBNC E. faecalis cells

TEM of the normal cells showed that the *E. faecalis* were mostly coccoid, with an average diameter of 2.3-2.8 μ m (Figure 2a). The cell membranes were intact, cytoplasmic staining was dark, and a number of extracellular granular secretions were observed.

The VBNC cells exhibited several changes compared with normal cells. The average diameter of the coccoid cells was 3.0-4.5 μ m (Figure 2b, c, and d), which was much larger than that of normal cells. The membranes appeared to be intact, but the cytoplasm was irregular compressed and the staining was lighter. The cell cytoplasm was unevenly condensed, and there were significant gaps between the cytoplasmic membrane and the cell wall. Membrane curling was observed in some of the cells, which was associated with irregular cell morphology. Some cells appeared to have vacuolar degeneration and a loose uneven cytoplasmic matrix.



Figure 2. Morphological changes of the VBNC *Enterococcus faecalis* analyzed using TEM (65,000X). Normal cells (**a**) and VBNC cells (**b**, **c**, and **d**). b, c, and d are corresponding *E. faecalis* cells incubated in sterile distilled water, sterile sodium buffer, pH 9, and sodium buffer, pH 11, respectively.

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Adhesion ability of E. faecalis cells

Scanning electron microscopy showed that both normal and VBNC *E. faecalis* cells had a coccoid shape (Figures 3 and 4). Significantly fewer VBNC cells adhered to the collagen or dentine compared with the normal cells, particularly when the cells were cultured in sodium buffer, pH 11. A subset of VBNC cells appeared to have changes in their shape when adhered to dentine, and they were not able to enter the dentine tubes. In contrast, normal *E. faecalis* cells entered the dentine tubes and retained their coccoid shape.



Figure 3. Scanning electron micrograph of *Enterococcus faecalis* adhered to collagen fiber (5,000X). Exponential phase cells (**a**), VBNC cells in sterile distilled water (**b**), VBNC cells in sodium buffer (**c** and **d**) with collagen fiber incubated at 37°C for 48 h.



Figure 4. Scanning electron micrograph of *Enterococcus faecalis* adhered to dentine pieces (10,000X). Exponential phase cells (**a**), VBNC cells in sterile distilled water (**b**), VBNC cells in sodium buffer (**c** and **d**) with collagen fiber incubated at 37°C for 48 h.

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DISCUSSION

Some species of bacteria enter the VBNC state when they are exposed to adverse environments (Peciuliene et al., 2000; Heim et al., 2002; George et al., 2005). In this state, bacteria lose their ability to form colonies on culture media (Lleò et al., 1998), but remain alive and capable of producing antigen components (Smith and Oliver, 2006) and virulence factors (Pruzzo et al., 2002). When permissive conditions are restored, VBNC cells may resume cell division (Lleò et al., 2001) and participate in disease transmission and persistence (Xu et al., 1982). VBNC *E. faecalis* cells are viable and can maintain their pathogenicity and other biological functions, such as gene expression (Lleò et al., 2000) and metabolic activity (Lleò et al., 1998). However, there are changes in cell physiology, biochemistry, and genetic characteristics (Oliver, 2005), including changes in morphology, metabolism, and gene expression, that help the bacteria adapt to the adverse environments. Most studies examining the VBNC state of *E. faecalis* is frequently associated with cases of failed endodontic treatment (Stuart et al., 2006). Therefore, studies on the VBNC state of *E. faecalis* will lead to a better understanding of treatment failure and suggest possible preventive measures.

Many studies have demonstrated that E. faecalis can withstand harsh environmental conditions, including high alkalinity (Sherman, 1937). E. faecalis can grow at pH 9.6 and can tolerate pH levels as high as 11.9 (Flahaut, 1997). In the RCT process, root canal preparation and clean-up, particularly the typical rinses with NaClO and H₂O₂ and disinfection with Ca(OH),, make the root canal a high-sodium alkaline oligotrophic environment. In this study, we simulated this unique physical microenvironment and developed 3 in vitro protocols to induce the VBNC state. Sterile distilled water represented an oligotrophic condition, while sodium buffer at pH 9 and 11 represented alkaline environments. All studies were conducted at physiological temperature (37°C), which was appropriate for bacterial growth. In this study, E. faecalis entered the VBNC state in both the oligotrophic and alkaline microenvironments, and the VBNC cells were able to remain viable for at least 2 months. Approximately 39 and 36% of *E. faecalis* survived after culture in sterile distilled water and sodium buffer, pH 9, respectively, for 2 months, while 22% of cells cultured in sodium buffer, pH 11, remained viable. These results were similar to those obtained in a study by Pruzzo et al. (2002), which focused on the ability of the VBNC E. faecalis cells to adhere to human cells. In that study, E. faecalis entered the VBNC state 16 days after culture in sterilized lake water and 90% of the bacteria remained alive. Furthermore, approximately one-third of the bacteria were alive after culture for 1 month, which is consistent with our results. In contrast, Heim et al. (2002) reported that 70% of E. faecalis remained 20 days after entry into the VBNC state. This discrepancy may be due to differences in the induction methods used in each study. Changes in the pH of the microenvironments, particularly increases in pH, may alter the physicochemical characteristics of the bacterial cells and trigger the expression of various genes associated with virulence or metabolic regulation (Olson, 1993). As a result, the bacteria may be able to survive and adapt to the adverse alkaline conditions by entering other physical phases.

E. faecalis is the most common species of *Enterococcus*. Most cells are arranged in pairs or short chains and they do not form spores or capsules. *E. faecalis* is catalase-negative and can decompose glucose and maltose into acid. Most strains of *E. faecalis* are capable of mannitol decomposition and bile aescin glycoside hydrolysis, are vancomycin-sensitive, are slightly alkyl aryl-lactamase-sensitive, are resistant to a range of pH conditions, and are resis-

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tant to salt and 40% bile salt. In this study, we used the Rapid ID 32 Strep kit and the ATB Mini API Software to automatically determine how effectively E. faecalis was able to decompose various biochemical substrates, such as lactose, maltose, and mannitol. Compared to exponentially growing cells, VBNC E. faecalis cells lost their ability to decompose lactose, Dmannitol, and D-sorbitol, but retained the ability to decompose most biochemical substrates, including sucrose. Because VBNC E. faecalis were no longer sensitive to certain metabolic substrates, their pathogenicity was decreased, but the carbohydrate metabolism that is retained may still cause harm to host tissues. VBNC E. faecalis produce and release various enzymes and metabolites, which can cause cell toxicity and lead to disintegration of the tissue matrix and spread of the bacteria, or cause indirect tissue damage by inducing an immune response. Enterococcus faecalis exhibits strong resistance to temperature and pH changes in the external environment, particularly to alkaline environments. The maximum pH tolerated by E. faecalis is 11.6, suggesting a relationship between the alkaline resistance of *E. faecalis* and its acid production capability. Our experimental observations showed that E. faecalis cultured in sterile distilled water decreased the pH of the culture medium from 8.05 to 6.06 over 1 month of culture; an equivalent drop in pH was observed in the exponentially growing bacterial culture within 24 h. However, in the sodium buffers, the bacteria reduced the pH values from initial values of 9.68 and 10.74 to 8.30 and 9.82, respectively, when they entered the VBNC state. These decreases were much smaller than those produced by bacteria in sterile distilled water. However, the reduction in pH confirmed the acid production ability of VBNC E. faecalis cells. Overall, the VBNC state of *E. faecalis* was characterized by lower pathogenicity than that of exponentially growing bacteria, as indicated by lower metabolic activity and acid production ability. VBNC cells retained the ability to produce and release several types of metabolites and enzymes and to produce acid, enabling them to maintain their viability and pathogenicity and survive until the external conditions were suitable to support normal growth.

The results of previous studies indicate that when E. faecalis enter the VBNC state, their physiology, biochemistry, and genetic characteristics change (Oliver, 2005), including morphological, metabolism, and genes expression changes. Several studies showed that most Gram-negative bacteria in the VBNC state exhibited a reduced cell volume and a globular morphology. In contrast, VBNC E. faecalis cells increased in size and were slightly elongated (Signoretto et al., 2000). Similarly, in this study, TEM revealed that the average volume of E. faecalis in the VBNC state was 1.5-fold larger than that of normal E. faecalis. Furthermore, no previous studies have reported the ultrastructure of VBNC E. faecalis cells. The bacteria observed in our study were irregular in shape because of cytoplasmic condensation. The cytoplasm appeared to be condensed, was darkly and unevenly stained, and contained many vacuoles that formed large gaps between the cell membrane and the cell wall. The granular extracellular secretions were also significantly reduced. These observations indicated that the VBNC state is associated with substantial changes in the cellular ultrastructure of E. *faecalis*. Bacteria may adapt to harsh environments by concentrating their nuclear material and changing their morphology to resist external pressure. Adhesion to the host is the first step in the infection process (Fux et al., 2005). E. faecalis exerts its pathogenicity by adhering to other bacteria, host tissue, or biomaterials and forming a biofilm. It was demonstrated in a previous study that E. faecalis in the VBNC state could not form a biofilm, but could still synthesize polymeric substances, thus partially maintaining their adhesion ability and pathogenic potential (Lleò et al., 2007). In this study, we used collagen fiber and dentine pieces collected from extracted teeth to evaluate the adhesion ability of VBNC E. faecalis

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cells *in vitro* and compared the results with bacteria in the exponential growth phase. These experiments showed that *E. faecalis* incubated in sterile distilled water adhered to collagen and dentine, but bacteria incubated in alkaline sodium buffer exhibited poor adhesion, and only a few bacteria were detected by scanning electron microscopy. It has been reported that adhesion factors on the cell surface, such as 'aggregation substance' and 'Ace', mediate the adhesion of *E. faecalis* to collagen (Rich et al., 1999; Rozdzinski et al., 2001; Hubble et al., 2003). Significantly, the expression of approximately 200 different proteins was induced in *E. faecalis* upon exposure to various stressful conditions (Flahaut et al., 1997). The function of these proteins is unknown (Rince et al., 2000; Giard et al., 2001), and additional studies are needed to confirm whether these proteins contribute to bacterial adhesion or resistance against environmental stress.

Bacteria only enter the VBNC state under specific environmental conditions. Previous in vitro VBNC induction protocols have focused mainly on changes in physicochemical conditions, such as temperature, nutrient availability, aeration, and illumination. The influence of these parameters on VBNC state entry is highly variable and depends on the bacterial species. The VBNC state of E. faecalis is typically induced by culture in disinfected filtered lake water as an oligotrophic medium for approximately 15 (Signoretto et al., 2000; Pfeffer et al., 2006) to 30 (Lleò et al., 1998) days with direct illumination at 4°C. In this study, we simulated the oligotrophic condition using sterilized distilled water and simulated the alkaline environment using sodium buffer at pH 9 and 11. A Live/Dead kit was used to determine the survival rate of the bacteria. This kit utilizes the green-fluorescent nucleic acid stain SYTO9 and the red-fluorescent nucleic acid stain propidium iodide. When used alone, the SYTO9 stain generally labels all bacteria in a population, including those with intact membranes and those with damaged membranes. However, propidium iodide penetrates only bacteria with damaged membranes, reducing SYTO9 fluorescence when both dyes are present. As a result, bacteria with intact cell membranes appear to be fluorescent green and bacteria with damaged membranes appear to be fluorescent red under fluorescence microscopy. Thus, the survival rate of E. faecalis was calculated based on the observed green fluorescence intensity.

In this study, we found that growth glucose metabolism, acid production, and the morphology of *E. faecalis* changed after entry into the VBNC state. We focused primarily on the VBNC state in *E. faecalis*. However, the mechanism of VBNC entry has not been fully determined, particularly at the level of protein and gene expression. Therefore, additional studies examining the entry and recovery from the VBNC state in *E. faecalis* are necessary. Moreover, this type of basic research may be used to guide clinical practice. Specifically, analyzing the mechanism underlying the changes that accompany the VBNC state in *E. faecalis* would be helpful for guiding clinical practice to improve the outcomes of root canal treatment.

Conflicts of interest

The authors declare no conflict of interest.

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