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### ORIGINAL RESEARCH A Novel Detection of Enterococcus faecalis Using Multiple Cross Displacement Amplification Linked with Gold Nanoparticle Lateral Flow Biosensor

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Background: Enterococcus faecalis, an opportunistic bacterial pathogen, is one of the most frequently isolated bacterial species and cause of serious nosocomial infections in recent decades. A reliable and rapid assay for E. faecalis detection is significant for the diagnosis and follow-up treatment.

Methods: A novel assay method, named multiple cross displacement amplification linked with nanoparticle-based lateral flow biosensor (MCDA-LFB), was applied for detecting E. faecalis strains. A set of special 10 primers was designed according to E. faecalis-specific gene Ef0027. The MCDA amplification conditions, including the target DNA concentration, reaction temperature and time, were optimized. The sensitivity and specificity of MCDA method were tested in the current study, and then, the MCDA-LFB technology was applied to detect the E. faecalis strain from clinical samples.

Results: The E. faecalis specific primers were valid for the establishment of MCDA-LFB technology for the detection of E. faecalis based on the Ef0027 gene. The MCDA amplification condition was optimized at 62°C for 35 min. The MCDA products were directly sensed and displayed with a biosensor. The full process, comprising genomic DNA template preparation (approximately 30 mins), amplification of MCDA (35 mins), and the product identification (approximately 2 mins), could be achieved in 70 mins. The MCDA technique could detect as little as 10 fg per reaction system of pure E. faecalis genomic DNA. The specificity of E. faecalis-MCDA-LFB method is 100%, with no cross-reactions to non-E. faecalis strains.

**Conclusion:** The MCDA-LFB technique established in the present study is a reliable, simple, rapid, sensitive and specific method to assay E. faecalis and can be applied for the detection of clinical samples.

Keywords: Enterococcus faecalis, limit of detection, multiple cross displacement amplification, gold nanoparticle, lateral flow biosensor, MCDA-LFB

#### Introduction

Enterococcus faecalis (E. faecalis), a Gram-positive enterococci, is widely distributed in the gastrointestinal tracts of humans, animals, and insects and natural environment.<sup>1,2</sup> E. faecalis has emerged as a major opportunistic pathogen cause of outbreaks of nosocomial infections affecting various tissues, including human urinary tract infections, intraabdominal infections, infective endocarditis, neonatal sepsis, and bacteremia.<sup>3</sup> Thus, to develop a reliable and rapid assay for *E. faecalis*, detection is significant for the diagnosis and follow-up treatment.

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The traditional and gold standard for the detection of E. faecalis is cultivation on agar plates, while this detection method took more than 48 h of growth.<sup>4,5</sup> Moreover, the sensitivity of cultivation-based techniques was significantly decreased if the clinical samples were collected after antimicrobial therapy.<sup>5</sup> Polymerase chain reaction (PCR) and PCR-based technologies (real-time PCR and multiplex PCR) were applied to detect clinical pathogens because these detection methods are specific, rapid, and sensitive.<sup>6-9</sup> Nevertheless, PCR-based methods need special experimental instruments and skilled personnel that may not be readily available in many resource-poor settings. Herein, advanced assays are urgently required for low-cost, reliable, sensitive, specific, and simple detection of target pathogens to ensure prompt treatment.

Multiple cross displacement amplification (MCDA) technique has been applied to detect a range of pathogens such as *Shigella* spp, *Listeria monocytogenes*, and *Neisseria meningitides*.<sup>10–12</sup> MCDA method can amplify target gene sensitively and efficiently, using a water bath or sample heater at a constant temperature.<sup>13,14</sup> In the MCDA amplification system, six primers for amplification (C1, D1, C2, D2, R1, and R2), two displacement primers (F1 and F2), and two cross primers (CP1 and CP2), designed according to target DNA sequence, were used for the amplification.<sup>15,16</sup> MCDA method is able to yield amplicons from as little as three microbes. These amplification products can be rapidly, reliably, and visually detected with disposable gold nanoparticles-based lateral flow biosensors (LFB).<sup>17</sup>

In the present study, the MCDA-LFB detection was developed for the visual, rapid, highly specific, and sensitive detection of *E. faecalis* carrying the *Ef0027* gene, which appeared to be uniquely present in *E. faecalis* as it showed no homology with other microbial genomes at GenBank by BLAST searches. The detection performance was analyzed with pure cultures and clinical samples.

#### Materials and Methods Ethics Statement

The study was approved by the Human Ethics Committee of the Second Affiliated Hospital of Guizhou University of Traditional Chinese Medicine and complied with the Declaration of Helsinki. All data/isolates were analyzed anonymously.

#### Reagents and Materials

Reagents including DNA extraction kits (Qiagen, Hilden, Germany), visual detection reagent (Malachite Green, MG) and isothermal amplification kits (Beijing HaiTaiZhengYuan technology Co., Ltd., Beijing, China), streptavidin-immobilized 30-nm gold nanoparticles (SA-Gs) (Resenbio Co., Ltd., Xian, China), Antarctic thermal-sensitive uracil-DNA-glycosylase (AUDG), dNTP ((New England Biolabs, Inc, Beijing, China) rabbit anti-fluorescein antibody (anti-FITC Ab), and biotinylated bovine serum albumin (biotin-BSA) (Abcam Co., Ltd., Shanghai, China) were used in this study. Materials including the sample pad, membrane backing card, nitrocellulose membrane (NC), conjugate pad, and absorbent pad were acquired from the Jieyi Biotechnology Co., Ltd. (Shanghai, China).

#### Design of MCDA Assay Primers

Based on the reaction mechanism of MCDA, 5 pairs of special primers according to the *Ef0027* gene (Genbank accession no. 1198935) of *E. faecalis* were devised by PRIMER PREMIER 5.0 and Primer Explorer V4 (Eiken Chemical, Japan).

Hybrids and hairpin structures of the primers were analyzed by the Integrated DNA Technologies design tools. The specificity of *E. faecalis*-MCDA primers was verified by Blast analysis tool. The C1 and D1 primers were labeled by biotin and fluorescein isothiocyanate (FITC) at their 5' end, respectively. All of the primers were synthesized by TsingKe Biotech Co., Ltd (Beijing, China) with HPLC purification grade. The primer sequences are shown in Table 1, and the primer positions of the *E. faecalis* gene are displayed in Figure 1.

### Bacterial Strains and Genomic DNA Template Preparation

In the present study, forty-nine bacterial strains, including standard strain of *E. faecalis* (ATCC 29212), twelve clinical *E. faecalis*-positive isolates, and thirty-six non-*E. faecalis* strains, were used in this study (Table 2). The *E. faecalis* standard strain (ATCC 29212) was applied for the optimization of the target method. The DNA templates were prepared by extracting the genomic DNA from the bacterial isolates using QIAamp DNA Mini Kit in accordance with the instructions from the manufacturer and measured with a Nanodrop ND-2000 (Beijing, China) at A260/280.

Table I	The	Primers	Used	in	the	Present	Study
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Primers Name	Sequences and Modifications	Length	Gene
FI	5'-TTAGGAAAAAAAGCCTATGCA-3'	21 nt	Ef0027
F2	5'-GTCCCTTTGGCAAATAACG-3'	l9 nt	
CPI	5'-ACAAAAATGGCTAAATCCTGTCCACTGATTGGGATGAACAAGC-3'	43 mer	
CP2	5'-GCACGTGTAGCGCAGGAACGACAATTTTTCCAAGGTGCT-3'	39 mer	
СІ	5'-ACAAAAATGGCTAAATCCTGTCCA-3'	24 nt	
CI*	5'-Biotin-ACAAAAATGGCTAAATCCTGTCCA-3'	24 nt	
C2	5'-GCACGTGTAGCGCAGGAAC-3'	19 nt	
DI	5'-TCCAGATTCTTAACTGCTGCC-3'	21 nt	
DI*	5'-FITC-TCCAGATTCTTAACTGCTGCC-3'	21 nt	
D2	5'-GCAAGTGCCGATGATTAGTTTG-3'	22 nt	
RI	5'-AGTTTCTCCTGTATAACTG-3'	19 nt	
R2	5'-AAGGGGATTTTAGCGTAT-3'	18 nt	
RI R2	5'-AGTTTCTCCTGTATAACTG-3' 5'-AAGGGGATTTTAGCGTAT-3'	19 nt 18 nt	

Notes: C1\*, 5'-labeled with biotin when used in MCDA-LFB assay; D1\*, 5'-labeled with FITC when used in MCDA-LFB assay. Abbreviations: FITC, fluorescein isothiocyanate; mer, monomeric unit; nt, nucleotide.

### Gold Nanoparticle-Based Lateral Flow Biosensor Preparation

The LFB (4mm×60 mm) was constructed according to the previous reports.<sup>18,19</sup> In brief, a plastic adhesive backing card was laminated with NC membrane, sample pad, absorbent pad, and conjugate pad, followed by spraying with biotin-BSA (2.5 mg/mL) and anti-FITC Ab (0.15 mg/mL) conjugates onto the NC membrane to produce the line for control sample (CL) and test sample (TL). Each line was separated with 5 mm. And the conjugate pad of the strip was deposited with SA-G in 0.01M PBS (PH 7.4). The 4-mm-wide strips were prepared by cutting the assembled cards. The prepared biosensors were preserved in a plastic box with a desiccant gel at room temperature.

**Figure I** Location and sequence of the *Enterococcus faecalis* specific gene *Ef0027* applied for the MCDA primer design. The nucleotide sequence of the sense strand of the *E faecalis* gene *Ef0027* is shown in the diagram. Right arrows and left arrows indicate sense and complementary sequences which were used in the current study, respectively. **Abbreviation:** MCDA, multiple cross displacement amplification.

#### MCDA-LFB Detection

MCDA was performed in a 25  $\mu$ L reaction system as described by previously reported.<sup>20</sup> In brief, 0.4  $\mu$ M of d F1 and F2 (displacement primers), 1.2  $\mu$ M of R1, R2, D1\* and D2, 0.8  $\mu$ M of C1\* and C2, 1.2  $\mu$ M of CP1 and CP2 (cross primers), 1.25  $\mu$ L of *Bst* DNA polymerase (10 U), and 12.5  $\mu$ L of 2×reaction mix, and 1  $\mu$ L prepared DNA.

Colorimetric indicator (Malachite Green, MG) and lateral flow biosensor (LFB) methods were applied for the determination and verification of the *E. faecalis*-MCDA products, respectively. The color of effectively amplified products could change from colorless to light green by using MG method. However, the color of negative and blank controls kept colorless. For the LFB assay, two visible red lines (the CL and TL) could display for the positive results but only the line (CL) appeared for the negative and blank control samples.

## Temperature Optimization of the *E. faecalis*-MCDA-LFB Assay

The *E. faecalis*-MCDA reaction temperature used for optimization was from 60°C to 67°C for 60 mins with 10 pg *Enterococcus faecium* DNA template. In addition, DNA of *Staphylococcus aureus* (ATCC 25923) and double-distilled water (DW) were used as negative control and a blank control, respectively. The MCDA amplifications were monitored through the turbidity of products. The curves of DNA concentrations of each amplified products were exhibited in the graph. Turbidity >0.1 was considered as positive.

No.	Bacteria	Strain No. (Source of Strains)	No. of Strains	MCDA-LFB Result
I	E. faecalis	ATCC 29212	1	Р
2	E. faecalis	Isolated strains (2nd GZUTCM)	12	Ρ
3	Enterococcus faecium	lsolated strains (2nd GZUTCM)	9	Ν
4	Enterococcus avium	lsolated strains (2nd GZUTCM)	I	Ν
5	Enterococcus raffinosus	lsolated strains (2nd GZUTCM)	I	Ν
6	Staphylococcus aureus	ATCC 25923	1	Z
7	Pseudomonas aeruginosa	ATCC27853	I	Ν
8	Legionellae bacillus	Isolated strains (2nd GZUTCM)	I	Ν
9	Hemophililus parainfluenza	lsolated strains (2th GZUTCM)	I	Ν
10	Bordetella parapertussis	lsolated strains (2nd GZUTCM)	I	Ν
11	Klebsiella pneumoniae	lsolated strains (2nd GZUTCM)	I	Ν
12	Shigella boydii	lsolated strains (2nd GZUTCM)	I	Ν
13	Mycoplasma pneumoniae	lsolated strains (2nd GZUTCM)	I	Ν
14	Enteropathogenic Escherichia coli	Isolated strains (GZCDC)	I	Ν
15	Streptococcus pneumoniae	lsolated strains (2nd GZUTCM)	I	Ν
16	Staphylococcus saprophyticus	Isolated strains (GZCDC)	I	Ν
17	Enterotoxigenic Escherichia coli	Isolated strains (2nd GZUTCM)	I	Ν
18	Invasive Escherichia coli	Isolated strains (2nd GZUTCM)	1	N
19	Enterohemorrhagic Escherichia coli	lsolated strains (2nd GZUTCM)	1	N
20	Enteroaggregative Escherichia coli	Isolated strains (2nd GZUTCM)	1	N
21	Streptococcus suis	lsolated strains (GZCDC)	1	Ν

Table 2 Bacterial Strains Used in the Current Study

(Continued)

No.	Bacteria	Strain No. (Source of Strains)	No. of Strains	MCDA-LFB Result
22	Vibrio cholerae	lsolated strains (GZCDC)	1	Ν
23	Acinetobacter baumannii	Isolated strains (2nd GZUTCM)	1	Ν
24	Bacillus cereus	Isolated strains (GZCDC)	1	Ν
25	Vibrio parahemolyticu	Isolated strains (GZCDC)	I	Ν
26	Shigella flexneri	Isolated strains (2nd GZUTCM)	1	Ν
27	Listeria monocytogenes	Isolated strains (2nd GZUTCM)	I	Ν
28	Mycobacterium tuberculosis	Isolated strains (GZCDC)	I	Ν
29	Leptospira interrogans	Isolated strains (GZCDC)	1	Ν
30	Brucella suis	Isolated strains (GZCDC)	1	Ν

Table 2 (Continued).

Notes: 2nd GZUTCM, the Second Affiliated Hospital, Guizhou University of Traditional Chinese Medicine.

Abbreviations: ATCC, American type culture collection; GZCDC, Guizhou Provincial Center for Disease Control and Prevention; P, positive; N, negative.

#### Analytical Sensitivity of *E. faecalis*-MCDA-LFB Assay

The sensitivity of *E. faecalis*-MCDA-LFB method was determined using 1 $\mu$ L of the serial dilution (10 ng, 10 pg, 1 pg, 100 fg, 10 fg, 1fg and 100 atto gram per microliter) of the genomic DNA extracted from *E. faecalis*. The amplification was carried out at the optimized temperature to measure the limit of detection (LoD) of *E. faecalis*-MCDA-LFB. Three replicates of each dilution were tested.

### Optimization of the Amplification Time for *E. faecalis*-MCDA-LFB Assay

The *E. faecalis*-MCDA reaction system was amplified at the optimized temperature for diverse amplification times (from 15 to 45 mins, 10 mins of the interval). And then, the products were determined using LFB, and each amplification time was tested at least three times.

# Specificity Analysis of *E. faecalis*-MCDA-LFB Detection

Genomic DNA templates from 49 different bacterial strains (Table 2) were used to verify the specificity of *E. faecalis*-

MCDA-LFB method. The 49 bacterial strains included the standard strain of *E. faecalis* (ATCC 29212), 12 *E. faecalis* isolates, and 36 non-*E. faecalis* strains. DW was used as a blank control. All of the MCDA results were tested with LFB and confirmed at least three times.

### Application of *E. faecalis*-MCDA-LFB Method for the Detection of Clinical Samples

Fifty-three urine samples of patients, 30 hospital surface samples, and 35 urine samples of healthy volunteers were collected from the Second Affiliated Hospital of Guizhou University of Traditional Chinese Medicine. All of the samples were detected for *E. faecalis* using traditional culture, PCR, and DCDA-LFB techniques, respectively. Traditional culture methods including colony morphology, Gram stain, and biochemical identification were used for identification of the *E. faecalis* isolates. PCR detections of all the genomic DNA templates were conducted by using *E. faecalis* specific primers.<sup>8</sup> The MCDA-LFB detection results were compared with that of traditional culture and PCR assay.

#### **Results** Verification of *E. faecalis*-MCDA Productions

In order to confirm the validity of primers (Table 1) for the *E. faecalis*-MCDA assay, genomic DNA from pure *E. faecalis* cultures were amplified at 64°C for 60 mins. The products were identified using MG and LFB methods. The results showed that amplification appeared with genomic DNA from *E. faecalis* (ATCC 29212), but not with *E. faecium, S. aureus*, and the blank control (Figure 2A and B). Therefore, the *E. faecalis*-MCDA primers used in the current study were valid for the development of *E. faecalis*-MCDA-LFB.

# Optimal Temperature for *E. faecalis*-MCDA-LFB Detection

To certify the optimum amplification temperature, 10 pg of genomic DNA of *E. faecalis* strain was utilized as the target template in each reaction mixture. The results showed that  $62^{\circ}$ C was the faster amplification as indicated by the kinetics graphs of real-time turbidity (Figure 3).



Figure 2 Confirmation and verification of *E. faecalis*-MCDA products. (A) The *E. faecalis*-MCDA amplification products were detected by the MG method through visual observation of the color change. (B) LFB was applied for the visual detection of *E. faecalis*-MCDA products. Tube 1/Biosensor 1: positive amplification of *E. faecalis* strain 29212; Tube 2/Biosensor 2: negative amplification of *Staphylococcus aureus* (ATCC25923); Tube 4/Biosensor 4: blank control (DVV).

Therefore, the most suitable temperature for *E. faecalis*-MCDA-LFB was determined as  $62^{\circ}$ C, which was applied to the rest of the assays in the current study.

# Sensitivity of MCDA-LFB Method for *E. faecalis* Assay

The sensitivity of *E. faecalis*-MCDA-LFB was tested by serially diluted DNA sample from *E. faecalis* strain (ATCC29212) and determined by visual inspection of reaction productions with MG reagents and lateral flow biosensors, respectively. Both of the two methods demonstrated that the LoD of *E. faecalis*-MCDA-LFB was 10 fg DNA template (Figure 4A and B).

## Optimized Amplification Time for *E. faecalis*-MCDA-LFB Detection

In order to confirm the optimum amplification time for the *E. faecalis*-MCDA-LFB detection during amplification phases, 15-, 25-, 35-, and 45-min amplification times were selected and compared at  $62^{\circ}$ C, respectively. The detection limit level of DNA (10 fg of *E. faecalis* genomic templates per reaction) showed both TLand CL when the reaction proceeded for 35 and 45 mins (Figure 5). Therefore, 35 mins was deemed as an appropriate amplification time for *E. faecalis*-MCDA-LFB detection.

# Specificity of MCDA-LFB Method for *E. faecalis* Assay

The genomic DNA from all of the bacterial strains (Table 1) were used for *E. faecalis*-MCDA-LFB specificity determination. The results showed that only the genomic DNA from *E. faecalis* strains showed positive results, while other non-*E. faecalis* isolates and blank control showed negative results (Figure 6).

### Detection Results of *E. faecalis*-MCDA-LFB for Clinical Samples

In order to demonstrate that the MCDA-LFB method is a reliable and rapid tool for *E. faecalis* detection in clinical samples, 53 urine samples of patients, 30 hospital surface samples, and 35 urine samples of healthy volunteers collected from the Second Affiliated Hospital of Guizhou University of Traditional Chinese Medicine were used for the application of *E. faecalis*-MCDA-LFB established in the current study. The results showed that 23 of 118 samples had been verified as *E. faecalis*-positive results through traditional culture technique, and the other 95



**Figure 3** (A–H) Optimization of reaction temperature for *E. faecalis*-MCDA primers. The MCDA amplifications for the detection of *E. faecalis* were monitored through real-time turbidity, and the corresponding curves of DNA concentrations were displayed in the graph. The threshold value was 0.1 and a turbidity>0.1 was considered as positive. Mixtures with 10-pg genomic templates of *Enterococcus faecium* (isolated strain) and *Staphylococcus aureus* (ATCC 25923) were used as negative controls (NCs), and 1  $\mu$ L of double-distilled water (DW) was used as a blank control (BC). Eight kinetic graphs were obtained at different temperatures (60–67°C, 1°C intervals) with 10-pg target genomic DNA per reaction. The graphs from C to H showed robust amplification.

samples showed *E. faecalis*-negative outcome. The MCDA-LFB assay results were consistent with traditional cultivation detection results. However, only 15 urine samples showed positive results when using conventional *Ef0027* gene-based PCR, and all of the 15 positive urine samples were covered by the *E. faecalis* positive samples detected by traditional culture and MCDA-LFB methods (Table 3).

#### Discussion

*E. faecalis* is becoming increasingly important for causing bacteremia, endocarditis, and other nosocomial infections in recent decades.<sup>21,22</sup> Moreover, more and more *E.* 



**Figure 4** Sensitivity analysis of the MCDA-LFB detection by serial dilutions of genomic DNA extracted from *E. faecalis* strain. Two detection methods, colorimetric indicator (MG; **A**) and lateral flow biosensor (**B**), were used to analyze the amplification productions. The serial dilutions of *E. faecalis* genomic DNA (10 ng, 10 pg, 1 pg, 100 fg, 10 fg, and 100 atto gram per microliter) were subjected to standard MCDA amplifications. Tubes (**A**)/Biosensors (**B**) 1–8 represent the DNA levels of 10 ng, 10 pg, 1 pg, 100 fg, 10 fg, 1 fg, and 100 atto gram per reaction of target templates and blank control (DW), respectively. The target gene levels of 10 ng, 10 fg, 10 fg, 10 fg per reaction showed positive amplifications.

*faecalis* isolates build up antibiotic resistance.<sup>2,23,24</sup> Thus, developing a reliable, rapid, specific, and sensitive detection method for *E. faecalis* is essential for the disease diagnosis and its therapy. Our study showed that the newly established *E. faecalis*-MCDA-LFB is a rapid, sensitive, and specific method to detect *E. faecalis*.

The specificity of MCDA was powerfully confirmed with genomic DNA from E. faecalis pure cultures. All of the E. faecalis isolates showed positive results, but all the non-E. faecalis strains displayed a negative outcome (Figure 6). Thus, the E. faecalis-MCDA-LFB method has a high level of specificity for the identification of E. faecalis species. The high degree of specificity of the MCDA-LFB established in the present study seems due to using the Ef0027 gene, which encodes a putative phosphosugar-binding transcriptional regulator. This gene appeared to be uniquely present in E. faecalis as it showed no homology with other microbial genomes at GenBank by BLAST searches, and the Ef0027 gene also applied to develop some advanced methods for detecting *E. faecalis* species.<sup>8,25</sup> Another reason to explain the high degree of specificity of E. faecalis-MCDA-LFB is because we designed ten primers targeting the different regions of the Ef0027 gene sequence.

Compared with conventional culture and PCR-based methods, the MCDA-LFB developed in this study is more time-saving and economical.<sup>10,12,26</sup> The *E. faecalis*-MCDA-LFB reaction just requires a simple incubation at 62°C for 35 mins. There are various portable user-friendly instruments for MCDA reaction. For example, the dry block heater (Labnet, USA) is one of the suitable instruments, which is a portable, battery-powered equipment supporting 96-channel MCDA reactions per detection.



Figure 5 Optimization of amplification time for *E. faecalis*-MCDA-LFB detection method. Different amplification times (A, 15mins; B, 25 mins; C, 35 mins; D, 45 mins) were tested at 62°C, respectively. Biosensors 1, 2, 3, 4, 5, 6, 7, and 8 represent genomic DNA levels of 10 ng, 10 pg, 1 pg, 100 fg, 10 fg, 1 fg, and 100 atto gram per reaction of target templates and blank control (DW). The best sensitivity was observed when the amplification lasted for 35 mins; C).

Moreover, the universal isothermal amplification kits such as NEB Warmstart and Eiken Loopamp kits can be commercially obtained and applied for MCDA amplification, and the cost of MCDA reaction is estimated to be \$3.5 USD, the LFB detection merely costs approximately \$2 USD, which is cheaper than the common PCR-based methods.<sup>15,16</sup> Additionally, this method can decrease labor costs because performing the MCDA-LFB assay does not need skilled technical personnel. The entire detection process, including template preparation (approximately 30 mins), isothermal amplification (35 mins), and LFB reading (approximately 2 mins), could be accomplished within 70 mins. Thus, this method can save precious time and cost of detection compared with the traditional methods.

In addition, the advanced *E. faecalis*-MCDA-LFB technique was more sensitive than normal PCR and loopmediated isothermal amplification (LAMP) methods. The newly developed *E. faecalis*-MCDA-LFB method was capable of assaying 10 fg of *E. faecalis* genomic DNA (Figure 4), which was more sensitive than *E. faecalis*-LAMP-LFB detection (250 fg per reaction) reported by the previous study.<sup>25</sup> Although the amplification products could be detected equally with the MG method used in the present study, the LFB was deemed as the preferred method as observing the amplification results is more objective and does not need any apparatus.<sup>19,27</sup> In conclusion, the newly *E. faecalis*-MCDA-LFB assay in this study is a valuable method for the convenient, rapid, sensitive, specific, visual, reliable, and low-cost detection of *E. faecalis* in clinical and environmental samples, especially in resource-constrained settings. However, the *E. faecalis*-MCDA-LFB method has some limitations when compared with some methods which can be applied for *E. faecalis* antimicrobial resistance detection. Later, we will study on the rapid and reliable assay of drug resistance genes of *E. faecalis* based on the *E. faecalis* MCDA-LFB method created in the current study.

#### Conclusion

In this study, a reliable and rapid *E. faecalis*-MCDA-LFB method was developed for detecting *E. faecalis*, causing nosocomial infection, including urinary tract infection, bloodstream infection, and surgical site infection in human. This method could rapidly, sensitively, specifically, and reliably detect the *E. faecalis* agent in the urine sample and did not rely on expensive instruments and reagents. Using LFB detection could help to obtain a rapid, objective, and easily interpretable result. Hence, the *E. faecalis*-MCDA-LFB assay could be considered as a useful method for reliable and rapid detection of *E. faecalis* in clinical and environmental samples, particularly in resource-limited regions of the world.



Figure 6 Specificity analysis of the *E. faecalis*-MCDA-LFB detection for different strains. The MCDA amplifications were carried out using different strain genomic DNA as templates and determined by means of visual LFB method. Biosensor I, *E. faecalis* strain (ATCC 29212); biosensor 2–13, twelve isolated strain of *E. faecalis* from the Second Affiliated Hospital, Guizhou University of Traditional Chinese Medicine; biosensor 14–22, nine isolated *Enterococcus faecium* strains, biosensor 23–49, *Enterococcus avium*, *Enterococcus raffinosus*, Staphylococcus aureus, Pseudomonas aeruginosa, Legionellae bacillus, Hemophililus parainfluenza, Bordetella parapertussis, Klebsiella pneumonia, Shigella boydii, Mycoplasma pneumonia, Enteropathogenic Escherichia coli, Streptococcus pneumonia, Staphylococcus saprophyticus, Enterotoxigenic *Escherichia coli*, Invasive Escherichia coli, Enterohemorrhagic Escherichia coli, Enterose, Vibrio parahemolyticu, Shigella flexneri, Listeria monocytogenes, Mycobacterium tuberculosis, Leptospira interrogans, Brucella suis; biosensor 50, blank control (DW).

Detection Methods	Clinical Samples (n=53)		Hospital Surfaces Samples (n=30)		Healthy Volunteers Samples (n=35)	
	Positive	Negative	Positive	Negative	Positive	Negative
Culture	18	35	2	28	3	32
PCR	14	39	0	30	1	34
MCDA-LFB	18	35	2	28	3	32

Table 3 Comparison of Conventional Culture, PCR. and MCDA-LFB Methods for the Assay of E. Faecalis in Clinical Samples

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#### Disclosure

All of the authors declared that there are no competing interests in this article.

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