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## ORIGINAL RESEARCH Rapid Detection Method for Pathogenic Candida Captured by Magnetic Nanoparticles and Identified Using SERS via AgNPs<sup>+</sup>

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Purpose: Candidemia infection is common in the clinic and has a high mortality rate. Candida albicans, Candida tropicalis, and Candida krusei are very important and common pathogenic species. Candida is difficult to isolate from clinical samples and culture, and immunological detection cannot distinguish these related strains. Furthermore, Candida has a complex cell wall, which causes difficulties in the extraction of DNA for nucleic acid detection. The purpose of this study was to establish a protocol for the direct identification of Candida from serum.

**Materials and Methods:** We synthesized  $Fe_3O_4(a)PEI$  (where PEI stands for polyethylenimine) magnetic nanoparticles to capture Candida and prepared positively charged silver nanoparticles (AgNPs<sup>+</sup>) as the substrate for surface-enhanced Raman scattering (SERS). Candida was directly identified from serum by SERS detection.

Results: Orthogonal partial least squares discriminant analysis (OPLS-DA) was used as the multivariate analysis tool. Principal component analysis confirmed that this method can clearly distinguish common Candida. After 10-fold cross-validation, the accuracy of training data in this model was 100% and the accuracy of test data was 99.8%, indicating that the model has good classification ability.

**Conclusion:** The detection could be completed within 40 minutes using Fe<sub>3</sub>O<sub>4</sub>@PEI and AgNPs<sup>+</sup> prepared in advance. This is the first time that Fe<sub>3</sub>O<sub>4</sub>@PEI was used in the detection of Candida by SERS. We report the first rapid method to identify fungi directly from serum without breaking the cell wall to extract DNA from the fungi.

Keywords: capture, surface-enhanced Raman scattering, positively charged silver nanoparticles, orthogonal partial least squares discriminant analysis, 10-fold cross-validation

#### Introduction

Broad-spectrum antibiotics and immune inhibitors are widely used, especially in cancer patients after radiotherapy and chemotherapy. The rate of Candida infection has recently increased; it now has the highest mortality of the invasive fungal infections.<sup>1</sup> Candidemia is the most common type of invasive *Candida* infection. It mainly refers to a bloodstream infection caused by Candida. Candida albicans, *Candida tropicalis*, and *Candida krusei* have high incidence rates in the clinic.<sup>2</sup> The incidence rate of C. albicans is the highest among Candida species; however, a 5-year study by Barchiesi et al in Italy showed that the mortality rate of non-C. albicans and that of C. albicans are similar. It even tends to exceed the mortality of C. albicans.<sup>3</sup> Candidemia has a high fatality rate, with an all-cause in-hospital

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941

mortality rate of up to 30%. As such, it has a worse therapeutic response than bacteremia.<sup>4</sup>

Early diagnosis of invasive fungi disease and timely drug treatment are critical to reducing candidemia mortality. However, candidemia has no specific clinical manifestations, and it is difficult to make an early diagnosis based on clinical symptoms; thus, candidemia must be confirmed by clinical laboratory results.<sup>4</sup> *Candida* has been cultured in vitro for a long time and is difficult to grow.

Biochemical and matrix-assisted laser desorption ionization time-of-flight mass spectrometry (MALDI-TOF MS) methods for the identification of *Candida* in the clinic are still based on culture methods.<sup>5,6</sup> There are some rapid detection methods without culturing in the clinic such as 1-3- $\beta$ -D glucan (G test), mannose lgG, polymerase chain reaction (PCR), and metagenomics next-generation sequencing (mNGS). The G test and mannose lgG cannot determine the *Candida* species.<sup>7</sup> For PCR and mNGS, it is required to break the cell wall to extract DNA; however, the fungal cell wall is complex and difficult to break.<sup>8,9</sup> Thus, it is difficult to extract DNA that can be identified under non-culture conditions.<sup>10</sup> Therefore, a rapid detection method based on impure culture without breaking the cell wall is urgently needed.

It has been reported that it is possible to detect pathogens using the fingerprints of whole organisms via surfaceenhanced Raman scattering (SERS).<sup>11</sup> In recent years, SERS has been applied to the detection of bacteria, but there are still few reports on the detection of fungi. SERS has a strong specificity, requires no contact with samples, has low temperature and pressure requirements, involves fast spectral imaging, and has a high resolution. This is especially true for samples that can be tested in a nondestructive manner. Moreover, the SERS spectrometer is stable with low maintenance costs, and the operation is simple.<sup>12,13</sup>

In this study, we tested the zeta potential of *Candida* and found that the surfaces of *C. albicans, C. tropicalis,* and *C. krusei* were all negatively charged. Thus, we selected AgNPs<sup>+</sup> as the SERS-enhanced substrate.<sup>14,15</sup> We identified and distinguished three common *Candida* species through the following steps: We detected the SERS spectrum of three common *Candida* species with AgNPs<sup>+</sup> as standard spectrum.<sup>14,16</sup> Then magnetic microspheres (Fe<sub>3</sub>O<sub>4</sub>) were modified by polyethylenimine (PEI) to form Fe<sub>3</sub>O<sub>4</sub>@PEI. Fe<sub>3</sub>O<sub>4</sub>@PEI was mixed with the candidemia samples to form the Fe<sub>3</sub>O<sub>4</sub>@PEI@*Candida* mixture. We have used Fe<sub>3</sub>O<sub>4</sub>@PEI for the first time in the detection of

*Candida* by SERS. The supernatant of the mixture was discarded and mixed with SERS-enhanced materials, and the sample was evaluated on silicon wafers.<sup>17</sup> When the mixture dried, the SERS spectrum of the mixture was measured and compared with the standard spectrum. The resulting spectrum was consistent with the standard spectrum. Finally, orthogonal partial least squares discriminant analysis (OPLS-DA) was used to analyze the spectral data of three *Candida* species.<sup>18</sup>

## Materials and Methods Chemicals

Ferric oxide hexahydrate (FeCl<sub>3</sub>·6H<sub>2</sub>O), ethylene glycol (CH<sub>2</sub>OH)<sub>2</sub>), and sodium acetate (NaOAc) were purchased from the Beijing Sinopharm group. PEI, cetyl trimethyl ammonium bromide (CTAB), and sodium borohydride (NaBH<sub>4</sub>) were purchased from Sigma-Aldrich (Shanghai Trading Co., Ltd.), the 99% purity silver nitrate (AgNO<sub>3</sub>) came from Beijing Modern Oriental Co., and ammonium hydroxide (NH<sub>4</sub>OH) was supplied by the Xilong Chemical Group Co. Monocrystalline silicon sheets were purchased from Zhejiang Lijing Group Co., Ltd. The culture medium for Candida was provided by Shanghai Comaga Biotechnology Co., Ltd., and pure water was purchased from Hangzhou Wahaha Co., Ltd. The above reagent materials were all produced in China and are standardized commercial reagents that can be directly used in experiments without further purification. Clinical samples of Candida used in this experiment were provided by the Affiliated Hospital of Xuzhou Medical University.

#### Instrumentation

An Ht-7700 transmission electron microscope (TEM) was purchased from Hitachi (Japan) and a Shimadzu 2600 UVvisible spectrophotometer was purchased from Shimadzu. The Raman spectrometer was an i-Raman Pro Raman spectrometer (B&W Tek, USA). The zncl-bs230 \*230 heating electromagnetic stirrer was purchased from Beijing Century Huake Co., Ltd. (China). The constant temperature fungal incubator was from Beijing Fuyi Electric Co., Ltd. (China).

## Strain Collection and Identification

*Candida* was collected from samples with positive culture results. After 24 hours of constant temperature culture, the strains were identified by MALDI-TOF MS (Bruker Daltonics, Bremen, Germany). Fifteen samples of patient

serum and cultured strains of each of the three *Candida* species required in this study were collected and stored at  $-80^{\circ}$ C for later use. The serum was collected from participants in the experiment.

# Preparation of Fe<sub>3</sub>O<sub>4</sub>@PEI Magnetic Nanoparticles

First, 5.4 g FeCl<sub>3</sub>·6H<sub>2</sub>O was added to 160 mL (CH<sub>2</sub>OH)<sub>2</sub> and stirred for 30 min. Next, 4 g PEG6000 and 10.8 g NaAc was added to the solution and completely dissolved. The solution was then poured into a highpressure reactor and heated for 16 h at a temperature of 210°C. When the reaction finished, the product was washed with ethanol and pure water. We then used a magnet to adsorb the supernatant. The residual product was dried at 60°C for 6 hours in a vacuum to obtain magnetic Fe<sub>3</sub>O<sub>4</sub> (300-500 nm). We next dispersed 500 mg of magnetic Fe<sub>3</sub>O<sub>4</sub> in PEI solution (0.5 g/ 100 mL) in an ultrasonic environment, and the PEI gradually bound to the magnetic Fe<sub>3</sub>O<sub>4</sub>. This process requires one hour to form  $Fe_3O_4$  (@PEI.<sup>19,20</sup> Finally, the  $Fe_3O_4$ @PEI was washed with ultra-pure water five times to remove excess PEI.

#### Preparation of SERS Substrates

We prepared positive AgNPs<sup>+</sup> as an enhanced substrate with CTAB as the stabilizer.<sup>18</sup> We then prepared two solutions. Solution 1 was 10.96 mg AgNO3 with a purity of 99%; 3.65 mg of CTAB and 0.5 mL of NH<sub>4</sub> OH were added at a concentration of 25% to 19.5 mL ultra-pure water. Solution 2 was prepared by adding 3.65 mg CTAB and 6.05 mg NaBH<sub>4</sub> to 20 mL ultrapure water. Solution 1 and solution 2 were placed in an ice bath for 10 min. After 10 min, solution 1 was poured into a conical flask. The conical flask was then heated on an electromagnetic stirrer and ice bath with intense stirring. During the stirring process, we used a sterile syringe to add solution 2 dropwise into solution 1. The solution gradually turned from colorless to dark yellow, indicating the formation of a colloidal dispersion system. This process lasted for four hours.

Silver can easily combine with oxygen and  $H_2S$  in the air, and the bottle mouth should be covered during stirring. After 4 hours, stirring was stopped, and the sample was heated for 10 minutes to remove the residual  $NH_3$  and  $NaBH_4$  in the solution. We began stirring again at the end of the heating. When the sample cooled to room

temperature, we removed it from the heated electromagnetic stirrer, added ultra-pure water, and diluted it to 40 mL. The sample was sealed and stored at 4°C.

## Establishment of Standard Raman Spectrum

*Candida* were grown on Sabouraud Agar medium at 25°C for 72 h; three strains of each *Candida* species were selected to collect standard Raman spectra. We prepared a *Candida* solution at 10 CFU mL<sup>-1</sup> and mixed 10  $\mu$ L of *Candida* solution with 10  $\mu$ L of the nanoparticles. Two microliters of the mixture was dropped onto a monocrystalline silicon wafer and left until the mixture dried. The Raman spectrometer (i-Raman Pro) analyzed the sample at 785 nm with a maximum laser power of 275 mW. The light spot diameter was 105  $\mu$ m, the integration time was 30 s, and the excitation power was 10% of the maximum power. Five sites were averaged from each species of *Candida* to collect one Raman spectrum.

## Fe<sub>3</sub>O<sub>4</sub>@PEI Capture Candida Test

The Fe<sub>3</sub>O<sub>4</sub>@PEI@*Candida* complex was formed after adding the volume of Fe<sub>3</sub>O<sub>4</sub>@PEI with the highest capture efficiency to serum containing *Candida* in a 1 mL Eppendorf tube. The complex was concentrated in the bottom of the tube with a magnet. The supernatant was discarded, and the AgNPs<sup>+</sup> were resuspended and rotated for 15 minutes.<sup>21</sup> We then concentrated the particles with a magnet in a small volume. The Raman spectrum of the precipitate was detected similarly as above and compared to the standard spectrum. If the spectrum could be matched, then the spectrum of the detected precipitate was *Candida*.

## Statistical Analysis

SIMCA 14.1 (Umetrics, Umea, Sweden) software is the most commonly used software for the analysis of spectral data. We used this software for OPLS-DA analysis. OPLS-DA is a supervised discriminant analysis statistical method and was used to establish the relationship model between metabolite expression and *Candida* species to predict the different species of *Candida*. The parameters  $R^2X$  and  $R^2Y$  in the model represent the interpretation rate of the established model to the X and Y matrices, respectively.  $Q^2$  indicates the prediction ability of the model. As the values of  $R^2$  and  $Q^2$  approach 1, the model theoretically becomes better with worse fitting accuracy. In general,  $R^2$  and  $Q^2$  values higher than 0.5 are good, higher than 0.4 is acceptable, and the difference between the two should not be too large. A 10-fold cross-validation can be used to evaluate the OPLS-DA model's accuracy for the classification of samples. For example, the data set is divided into 10 parts, of which 9 parts are the training data and 1 part is the test data. Each test gets a correct rate (or error rate). Generally, a 10-fold cross-validation is required, and the accuracy of the algorithm is then estimated via the average accuracy (or error rate).<sup>22</sup>

#### Results

## A Technical Route for Rapid Identification of *Candida*

Figure 1A shows the production process of  $Fe_3O_4$ @PEI. We improved the method of Liu et al.<sup>19,21</sup> and synthesized  $Fe_3O_4$  microspheres sized 300–500 nm. In an ultrasonic bath, PEI is mixed with  $Fe_3O_4$ . The cationic PEI will be adsorbed on the surface of  $Fe_3O_4$ , resulting in  $Fe_3O_4$ @PEI.

Figure 1B shows the rapid identification of *Candida* by  $Fe_3O_4@PEI$ . First,  $Fe_3O_4@PEI$  is adsorbed on the surface of *Candida* to form  $Fe_3O_4@PEI@Candida$  after placing magnets outside the Eppendorf tube to adsorb the contents of the Eppendorf tube. The  $Fe_3O_4@PEI$  can enrich *Candida* from the solution in seconds. The AgNP<sup>+</sup> substrate is also adsorbed on the surface of the complex to form  $Fe_3O_4@PEI@Candida@AgNPs^+$ . We identified *Candida* species via a SERS fingerprint spectrum.

#### Characterization of Nanomaterials

Figure 2A shows a TEM image of spherical AgNPs<sup>+</sup>. Figure 2B shows that the zeta potential of the AgNPs<sup>+</sup> is positive. Figure 2C shows that the particle size of the AgNPs<sup>+</sup> is more than 100 nm. The UV absorption spectrum of the AgNPs<sup>+</sup> is shown in Figure 2D, and the ultraviolet absorption peak is 404 nm.

Figure 3A is a TEM image of Fe<sub>3</sub>O<sub>4</sub>@PEI, which shows PEI attached to the surface of the Fe<sub>3</sub>O<sub>4</sub> forming a film. Monodisperse Fe<sub>3</sub>O<sub>4</sub>@PEI is uniform in size, spherical in shape, and 300–500 nm in diameter. Figure 3B shows Fe<sub>3</sub>O<sub>4</sub> particles with a zeta potential of -47.8 mV. The Fe<sub>3</sub>O<sub>4</sub>@PEI has a zeta potential of +38.1 mV. *C. albicans, C. tropicalis,* and *C. krusei* have zeta potentials of -12.2 mV, -19.9 mV, and -13.7 mV, respectively. Figure 3C shows that Fe<sub>3</sub>O<sub>4</sub> @PEI and AgNPs<sup>+</sup> are adsorbed on the surface of the *Candida* to form Fe<sub>3</sub>O<sub>4</sub>@PEI@*Candida*@AgNPs<sup>+.17,18</sup>

## Capture Efficiency of Fe3O4@PEI for Candida

We investigated the ability of Fe<sub>3</sub>O<sub>4</sub>@PEI to capture three *Candida* species from 1 mL of serum. The *Candida* titer in the serum samples was  $1 \times 10^{6}$  cells mL<sup>-1</sup>. After treatment, the residual *Candida* in the supernatant was counted to calculate the capture rate.<sup>20</sup> The volume of Fe<sub>3</sub>O<sub>4</sub>@PEI (1, 2, 3, 4, and 5 µL) was used as a variable to study the optimal capture efficiency. According to Figure 4, when the volume of Fe<sub>3</sub>O<sub>4</sub>@PEI  $\geq 4 \mu$ L, the capture efficiency of Fe<sub>3</sub>O<sub>4</sub>@PEI in the serum samples of different *Candida* concentrations reached equilibrium. The capture efficiencies of *C. albicans, C. tropicalis*, and *C. krusei* were 95.9%, 98.0%, and 79.6%, respectively.

#### SERS Spectrum

The SERS spectrum is the *Candida* species' average spectrum; the *x*-coordinate is the Raman shift, and the *y*-coordinate is the Raman intensity. Figure 5 compares the average SERS spectrum of a standard spectrum and







Figure 2 Characterization of AgNPs<sup>+</sup>. (A) TEM image of spherical AgNPs<sup>+</sup>. (B) The zeta potential of AgNPs<sup>+</sup> is +40.3 mV. (C) The particle size of AgNPs<sup>+</sup> is 122.4 nm. (D) The ultraviolet absorption peak of AgNPs<sup>+</sup> is found at 404 nm.



Figure 3 Characterization of  $Fe_3O_4$ @PEI. (A) TEM image of a  $Fe_3O_4$ @PEI that is uniform, spherical, and 300–500 nm in diameter. (B) The zeta potential of  $Fe_3O_4$  is -47.8 mV, and the zeta potential of  $Fe_3O_4$ @PEI is +38.1 mV. The zeta potentials of *C. albicans, C. tropicalis,* and *C. krusei* are -12.2 mV, -19.9 mV, and -13.7 mV, respectively. (C)  $Fe_3O_4$ @PEI and *Candida* adsorb AgNPs<sup>+</sup> on the surface.

the capture spectrum. The black line is the average standard SERS spectrum of the three species of *Candida*, and the red line is the average captured SERS spectrum of the three species of *Candida*. The captured spectrum is consistent with the standard spectrum.

Major spectral bands observed in *C. albicans, C. tropicalis*, and *C. krusei* are shown in Table 1. For *C. albicans*, the Raman shifts at 658, 715, 792, 1000, 1027, 1177, 1251, and 1444 cm<sup>-1</sup> were significantly enhanced. The Raman data for *C. albicans* and *C. tropicalis* were similar, but they have different Raman

945



Figure 4 The capture efficiency of *C. albicans, C. tropicalis*, and *C. krusei* (incubation time was 15 min).

shifts. The Raman shifts of *C. tropicalis* are at 656, 789, 1027, 1178, 1251, and 1443 cm<sup>-1</sup>. The Raman data for *C. krusei* are different from those of *C. albicans* and *C. tropicalis*. The Raman shifts of *C. krusei* are at 680, 964, 1044, 1092, 1163, 1260, and 1471 cm<sup>-1</sup>. Furthermore, the peaks of *C. albicans, C. tropicalis*, and *C. krusei* are all located at 640–675, 1022–1060, 1150–1186, 1215–1295, and 1440–1475 cm<sup>-1.23</sup>

#### Multivariate Statistical Analysis

OPLS-DA was used to compare the SERS spectrum of *Candida* captured from serum by  $Fe_3O_4$ @PEI containing *C. albicans, C. tropical,* and *C. krusei* (15 serum samples each). One point represents one sample (Figure 6). At the 95% confidence interval, the sample analysis results in an ellipse. The three *Candida* species can be nicely separated.

 $R^2X$  (cum) = 0.997,  $R^2Y$  (cum) = 0.93,  $Q^2$  (cum) = 0.924, so  $R^2X$ ,  $R^2Y$ , and  $Q^2$  are all above 0.7, indicating that the model is well established and has good predictive ability. We also conducted a 10-fold cross-validation to evaluate the accuracy of the model classification. The average accuracy of the training data and the data set were 100% and 99.8%, respectively, indicating that the classification ability of the model is good. Table 2 shows specific information on the 10-fold cross-validation of the OPLS-DA model.

#### Discussion

This study first established a method that can identify and distinguish *Candida* in impure cultures in a nondestructive and rapid manner. This method combined the SERS spectrum and OPLS-DA multivariate statistical analysis. The detection could be completed within 40 minutes using  $Fe_3O_4$ @PEI and AgNPs<sup>+</sup> prepared in advance.

Blood culture is currently the golden standard for the diagnosis of candidemia, but it has a long reporting time that can delay treatment.<sup>6</sup> The G test is common for early diagnosis of candidemia, but it cannot determine the species and has a high false positive rate.<sup>7</sup> Mannose IgG is easily cleared in the blood and requires continuous monitoring; thus, the clinical applications of this project require further research.<sup>7</sup> Some new techniques have also been applied to the clinical detection of fungi. MALDI-TOF MS is extremely accurate in identifying yeast, but it must be done with pure samples; the mass spectrometry cost is very high.<sup>5</sup> PCR can quickly detect a variety of clinical specimens with fungal infection at a high specificity. However, PCR requires extraction of DNA, and *Candida* has cell walls that are difficult to lyse.<sup>8</sup> mNGS



Figure 5 Average standard and captured SERS spectrum of *C. albicans*, *C. tropicalis*, and *C. krusei*. (A) Average standard and captured SERS spectrum of *C. albicans*. The Raman shifts at 658, 715, 792, 1000, 1027, 1177, 1251, and 1444 cm<sup>-1</sup> were significantly enhanced. (B) Average standard and captured SERS spectrum of *C. tropicalis*. The Raman shifts at 656, 789, 1027, 1178, 1251, and 1443 cm<sup>-1</sup> were significantly enhanced. (C) Average standard and captured SERS spectrum of *C. krusei*. The Raman shifts at 680, 964, 1044, 1092, 1163, 1260, and 1471 cm<sup>-1</sup> were significantly enhanced.

Assignment	Range (cm <sup>-1</sup> )	C. albicans	C. tropicalis	C. krusei
Guanine, tyrosine	640–675	++	++	++
Adenine, glycoside	713–740	+	-	-
Cytosine, uracil	745–790	-	+	-
C=C deformation, C–N stretching	930–990	-	-	+
Phenylalanine, galactomannan, C–C aromatic ring stretching	1000-1010	+	-	-
C-C stretching (phospholipids carbohydrates, eg chitin), C-N stretching	1022-1060	+	+	+
O-P-O (DNA), C-C or C-O-C stretching (carbohydrates, eg 1,3-b-glucan)	1078-1110	-	-	+
=C–O–C= (unsaturated fatty acids in lipids), galactomannan	1129-1145	-	-	-
C–O ring, aromatic amino acids in proteins	1150-1186	+	+	+
Chitin	1195-1200	-	-	-
Amide III (random), thymine	1215-1295	+	+	+
Amide III (protein), C–H deformation	1315-1325	-	-	-
Adenine, guanine, CH deformation	1330-1345	-	-	-
Galactomannan	~1410	-	-	-
CH <sub>2</sub> deformation	1440-1475	++	++	++
C=C stretching	~1490	-	-	-

Table I Major Spectrum Bands Observed in C. albicans, C. tropicalis, and C. krusei

Notes: ++, strong; +, present; -, absent.

also requires DNA extraction. In addition, the cost of detection is relatively high and needs further improvement for clinical applications.<sup>9</sup> SERS can detect fungi non-destructively, which is a major advantage over PCR and mNGS.<sup>24,25</sup>

The material capture on Fe<sub>3</sub>O<sub>4</sub>@PEI in serum samples containing *Candida* was analyzed by SERS. After enrichment, the Fe<sub>3</sub>O<sub>4</sub>@PEI@*Candida* was combined with AgNPs<sup>+</sup> for SERS detection and compared with a standard spectrum. The three *Candida* species were identified and distinguished by combining the resulting *Candida* spectrum with the stoichiometry of OPLS-DA. The process did not require the destruction of *Candida* cell walls and was fast. Fe<sub>3</sub>O<sub>4</sub>@PEI nanoparticles could capture *Candida* in a broad spectrum because its surface has a positive charge, and the *Candida* surface carries a negative charge. The negative charge on the PEI led to electrostatic attraction and *Candida* interactions, thus obtaining good capture efficiency (95.9% for *C. albicans*, 98.0% for *C. tropicalis*, and 79.6% for *C. krusei*). Moreover, the surface of AgNPs<sup>+</sup> carries a positive charge, and it does not combine with Fe<sub>3</sub>O<sub>4</sub> @PEI, but only with *Candida*. Therefore, AgNPs<sup>+</sup> are a good choice as the SERS-enhanced substrate in this study.

OPLS-DA can easily distinguish the differences between groups and improve the validity and analytical capability of the model. The OPLS-DA score graph of the



Figure 6 OPLS-DA of C. albicans, C. tropicalis, and C. krusei via SERS where one point represents one sample. The data points are nicely separated.

Time	Accuracy of Training Data	Accuracy of Test Data	
I	100%	99.3%	
2	100%	98.7%	
3	100%	100%	
4	100%	100%	
5	100%	100%	
6	100%	100%	
7	100%	100%	
8	100%	100%	
9	100%	100%	
10	100%	100%	
Mean	100%	99.8%	

**Table 2** Results of the 10-Fold Cross-Validation of C. albicans,C. tropicalis, and C. krusei Based on Orthogonal Partial LeastSquares Discriminant Analysis

model established in this study showed a clear separation trend (Figure 6).  $R^2X$  (cum) = 0.997,  $R^2Y$  (cum) = 0.93, and  $Q^2$  (cum) = 0.924, indicating that the model was built with good quality parameters.<sup>26,27</sup> We also estimated the classification efficiency of the model by performing a 10-fold cross-validation. The accuracy of the training data and test data of this model was 100% and 99.8%, respectively; the model has good classification ability.<sup>28</sup>

Our study shows SERS can be used for rapid and nondestructive detection of Candida under non-culture conditions. The equipment cost is relatively low. Candida is difficult to culture and has a cell wall unlike other pathogens, which makes it difficult to detect fungi quickly and nondestructively.<sup>29,30</sup> This highlights the advantages of SERS in detecting pathogenic fungi. Superior to others' former selection of materials,<sup>17,21</sup> we selected and prepared materials of Fe<sub>3</sub>O<sub>4</sub>@PEI (300–500 nm) and AgNPs<sup>+</sup> based on the characteristics of large diameter and negative charge of fungi. Compared with the latest method of detecting fungi SERS, our method does not require pure cultures<sup>18</sup> and does not require lysis of the fungal cell wall.<sup>31</sup> After trapping pathogenic fungi with magnetic beads, they were directly detected instead of cultured, which greatly improved the detection speed of pathogenic fungi and provided a beginning for solving the problem of long detection time of pathogenic fungi in the clinic. OPLS-DA data indicated that SERS could clearly distinguish these common Candida, and the method is suitable of further exploration and clinical application.

## Conclusion

We report a new method for the direct detection of *Candida* under impure culture conditions. Fe<sub>3</sub>O<sub>4</sub>@PEI

can capture *Candida*, and AgNPs<sup>+</sup> can be used as a substrate to enhance the SERS spectrum intensity. The detection could be completed within 1 hour, and it has been reported that the detection speed of fungi is the fastest among the non-culture detection methods that do not require wall lysis, thus indicating that the diagnostic tool based on SERS might be a powerful technique for the early diagnosis of candidemia.

### **Ethical Statement**

This study has been approved by the ethics review committee of the Affiliated Hospital of Xuzhou Medical University, China. Comply with the World Medical Association Declaration of Helsinki, the relevant Chinese laws and regulations, and other ethical principles to protect the rights and interests of the subjects, as well as the regulations of the ethics committee of Jiangsu (accession number: XXYFY2019-KL141-01). The informed consent of the patient was exempted.

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## **Author Contributions**

Y.Y. and B.G. conceived and designed the study. S. H. and F.G. did the experiments. H.K. provided clinical samples. C.W., S.C., W.G., and L.W. did the analysis of surface-enhanced Raman scattering spectrum data. S.H., H.K., and F.G. drafted the manuscript. All authors played a significant role in data collection and analysis. All authors made substantial contributions to conception and design, acquisition of data, or analysis and interpretation of data; took part in drafting the article or revising it critically for important intellectual content; agreed to submit to the current journal; gave final approval of the version to be published; and agree to be accountable for all aspects of the work.

#### Disclosure

The authors report no conflicts of interest in this work.

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