Pathogenic Microorganism Detection in Invasive Pulmonary Aspergillosis (IPA) Patients Using Bronchoalveolar Lavage Fluid

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ABSTRACT

Background: To evaluate the diagnostic values of different forms of pathogenic microorganism detection from the bronchoalveolar lavage fluid (BALF) for Invasive Pulmonary Aspergillosis (IPA) diagnosis.

Methods: Bronchoalveolar lavage fluid (BALF) was collected from 100 patients with suspected clinical pulmonary Aspergillus infections by means of bronchoscopy. The smear microscopy, fungal culture and PCR were used to detect the Aspergillus from the BALF.

Results: 15 cases were diagnosed by pathological data (the proven group), while 12 confirmed cases were supported by clinical examinations including radiology and etiology (the probable group). The 73 cases left were not IPA, among which 21 cases of common pneumonia were taken as the control group. For the practise of smear microscopy method, the sensitivity, specificity, and accuracy of the diagnoses were 40.7%, 100.0%, and 66.7% respectively. As for the fungal culture, the sensitivity, specificity, and accuracy were 37.0%, 100.0%, and 64.6% respectively. For PCR, the sensitivity, specificity, and accuracy were 55.6%, 100.0%, and 77.1% respectively. With the combination of the three methods mentioned, the final results were 92.6%, 100.0% and 95.8%.

Conclusion: The pathogenic detection of bronchoalveolar lavage fluid (BALF) is an ideal diagnostic method for Aspergillus infections. The smear microscopy method, the fungal culture and PCR has their own limitations in diagnosis. The detection of combining the three methods is proved to improve the efficiency of IPA diagnosis significantly.

Introduction

The insufficiency of detective methods during early stage of Invasive Pulmonary Aspergillosis (IPA) has lead to severe misdiagnosis [1]. Surveys and researches had been conducted to proved that the uncertainty and improper treatment could result in relatively high mortality rate [2,3]. Bronchoalveolar lavage (BAL) is the method which can collect secretory fluid in the respiratory tract and alveoli. The fluid can be separated for further analysis so that the content of the components can be quantified. It is an common method in locating the lesion site in clinical practice. Compared with other methods, for example, BALF has fewer chance of contamination rather than collecting sputum specimen. Therefore, it can improve the specificity of etiological diagnosis. At present, this technology is mainly applied in the clinical diagnosis of pulmonary bacterial infection, but there have been few reports on
the evaluation of its pathogenic detection for the diagnosis of fungal infection. In this study, BALF was collected from 100 suspected patients suffering from invasive pulmonary Aspergillus infection. Three pathogenic detection methods, smear microscopy, fungal culture and PCR were used to quantify the Aspergillus examples. Then the results of these three methods would be valued under the clinical criterion. To provide assistance for the clinical diagnosis of pulmonary aspergillosis.

**Method**

**Collection and Storage of the BALF Specimens**

The ethical approval for this study was obtained by First Affiliated Hospital of Sun Yat-sen University. Before bronchoalveolar lavage fluid were collected, the patients involved in this research received detailed explanation and understood the benefits and potential danger, and had signed the consent letters for bronchoscopy procedures. The subject inclusion and exclusion criteria used in this study refer to the consensus of Chinese experts on the detections of pathogens in the bronchoalveolar lavage of pulmonary infectious diseases, which was proposed by respiratory disease branch of the Chinese Medical Association in 2017 [4]. The procedure of collecting bronchoalveolar lavage was as follows:

a) 1-2mL of 2% lidocaine was injected through the biopsy hole for local anesthesia, after the bronchoscope reached the lung segment of the lavage site.

b) The top of the bronchoscope was incarcerated in the opening of the target bronchial segment or sub segment. A dose of 20-50mL sterile normal saline at 37 ℃ or room temperature was injected through the biopsy hole of the bronchoscope. After each injection, the appropriate negative pressure (less than 100 cmH₂O) was used to recover the liquid, and the total recovery rate was secured to be more than 40%. The intensity of negative pressure during lavage was under severe control to avoid the collapse of distal airway caused by excessive negative pressure.

c) 8-10mL of BALF specimens were collected with a sterilized plastic collector. And the specimens were placed in an incubator with ice cubes and immediately sent to -80℃ refrigerator in the laboratory facilities. The bronchoscopy biopsies had also been performed, and pathological diagnoses had been made by the pathology departments of each hospital.

**Detection of the Specimens**

After being collected, The BALF examples were thawed and divided into three parts for the application of smear microscopy, culture, and PCR product analyses. Smear microscopy and culture methods were conducted in the microbiology laboratory facilities of the First Affiliated Hospital of Sun Yat-Sen University. The smear microscopy was performed using fluorescent staining techniques. In the experimental processes, 5 mL of BALF were digested with digestive fluid (dithiothreitil). Then the examples were selected and centrifuged at 3,000 r for 15 to 20 minutes. The supernatant was discarded and the sediment remained. The sediment was placed on a glass slide and one drop of fluorescent staining solution would be added. The slide was observed under microscopy by technicians. After the residual sediment was treated again with digestive fluid, it was inoculated in Sabouraud culture medium and cultured at 30℃ for one week. After the culture was completed, the colony characteristics were observed and identified according to the characteristics of colony and conidia.

**BALF PCR**

**Primers:** After the BALF had been sent for microscopy and culturing, the remaining specimens were sent to Sagene Biotech Company Ltd for PCR detection. Then, the internal transcribed spacer (ITS) primers, which serve in Aspergillus identification, were redesigned based on the methods and sequences provided by the relevant literature [6, 7] (ITS-F:GCCTGTCCGAGCGTCATTG;ITS-R:TTAAGTTCAGCGGGTATCCCT; size of amplified fragments: 224 bp). The primers and the experimental reaction system were verified by the pre-experimental processes. The primers were found to have good specificity, and thus could be used for the identification of Aspergillus.

**DNA Extraction:** A DNA extraction kit (Sagene, China) was used to extract the DNA from each specimen. Following the extraction, gel electrophoresis was performed. 3 μL of DNA was taken and added to 1μL of loading buffer. Electrophoresis was conducted at 120 V for 25 minutes for the purpose of analyzing the quality of the extracted DNA.

**Establishment of the PCR Reaction System**

a) **Reagents:** The reagent types and volumes are shown in the Table 1.

b) **PCR Reaction Conditions:** The following were the PCR reaction conditions in this study: 98℃ for 1 minute; 98℃ for 10 seconds; 60℃ for 5 seconds; 72℃ for 5 seconds; and 72℃ for 2 minutes, for a total of 40 cycles.
c. **Preliminary Analysis**: For the preliminary analysis, 3 μL PCR product was selected to perform the 1.2% agarose gel electrophoresis (Figure 1) and preliminarily analyze the size of the amplified fragments, in order to determine whether or not the targeted fragments had been successfully amplified.

d. **Sequencing (Abi 3500) of the PCR products**: Sequencing of the PCR products was conducted for the specimens which have been successfully amplified to the target fragments. Then, an NCBI BLAST comparison was performed on the sequencing results (Picture 1).

Figure 1: ITS amplification electrophoresis map of aspergillus.

Picture 1: Example of NCBI BLAST comparison results of amplified fragment.

**Table 1**: The reagent types and volumes during the PCR reaction system.

| Reagent             | Volume  |
|---------------------|---------|
| 5xPCR Buffer        | 10μl    |
| dNTP(2.5mM each)    | 4μl     |
| Forward primer (10μM) | 1μl   |
| Reverse primer (10μM) | 1μl   |
| polymerase(2.5U/μl) | 0.5μl  |
| DNA template        | 10-20ng |
| ddH₂O               | up to 50μl |
Statistical Data Method

SPSS (IBM 22.0) software was used for the statistical data in this study.

a. A Chi Square Test Method was used to compare the distributions of the three detection methods among the proven and clinical diagnosis groups. The differences were considered to be statistically significant when $p<0.05$.

b. A Kappa Consistency Test Method was used for the pairwise comparisons of the results of the three detection methods as a whole and among the groups, respectively.

c. The sensitivity (Sn), specificity (Sp), and positive likelihood (PRL) ratios of the three detection methods for IPA diagnoses were calculated.

Results

Table 2: Clinical baseline data of patients.

| Features                      | Proven group N=15 | Clinical diagnosis group N=12 | Common pneumonia group N=21 | P value |
|-------------------------------|-------------------|------------------------------|-----------------------------|---------|
| Age (year, mean, interval)    | 47.93 (28-67)     | 54.92 (35-71)                | 59.14 (23-79)               | 0.286   |
| Gender (male, percentage)     | 10(66.7%)         | 8(66.7%)                     | 10(47.6%)                   | 0.414   |
| Underlying diseases           |                   |                              |                             |         |
| Bronchiectasis                | a9(60%)           | a7(58.3%)                    | b3(14.3%)                   | 0.007   |
| Obsolete pulmonary tuberculosis| a14(93.3%)        | a9(75%)                      | b3(14.3%)                   | 0       |
| Chronic obstructive pulmonary disease | 0(0.0%) | 0(0.0%)                      | 4(19.0%)                    | 0.112   |

In the present study, 100 patients with suspected clinical pulmonary Aspergillus infections were registered. Through clinical examinations including pathology, radiology, etiology. Patients’ data were collected (Table 2). According to the diagnosis standards of IDSA(2016) [5], 15 patients were diagnosed by pathological data(proven group), and 12 cases were diagnosed by radiology, etiology, and other clinical examinations. The other 73 patients were diagnosed as having tuberculosis, common bacterial pneumonia, vasculitis etc. Among those patients, 21 patients with common bacterial pneumonia were used as the control group. The positive number with smear microscopy, culture, and PCR in BALF was 6 (40.0%), 7 (46.7%), and 6 (40.0%) in the proven group respectively; and 5 (41.7%), 3 (25.0%), and 9 (75.0%) in the clinical diagnosis group, respectively. In addition, the total number of patients with positive results as diagnosed using the above-mentioned three methods was 0 in the common pneumonia group, as illustrated in Table 3.

Table 3: Number of positive cases in the diagnostic tests.

| Group                        | Smear microscopy (%) | Culture (%) | PCR(%) |
|------------------------------|----------------------|-------------|--------|
| Proven group (N = 15)        | 6 (40)               | 7 (46.7)    | 6 (40.0) |
| Clinical diagnosis group (N = 12) | 5 (41.7)            | 3 (25.0)    | 9 (75.0) |
| Common pneumonia group (N = 21) | 0(0)                | 0(0)        | 0(0)   |

- $\chi^2 = 2.187$ and $p = 0.335$ could be obtained, there were no significant differences in the distributions of the three detection methods among the two groups ($p > 0.05$). It was determined that the calculated $\chi^2$ and $p$ values were 0.182 and 0.913 (proven group) and 6.241 and 0.044 (clinical diagnosis group). The statistical results showed that in the clinical diagnosis group, the BALF culture and PCR results were significantly different ($p<0.05$). Furthermore, in the proven group, there were no significant differences observed between the three detection methods ($p > 0.05$).

- A Kappa Consistency Test Method was adopted in this study for the pairwise comparison of the results of the three detection methods as a whole, and also within the different groups. The results are shown in Table 4. As shown in the table, there was a weak consistency observed between the microscopy and culture methods. However, the results of both were found to be inconsistent with the PCR method.

- In accordance with the above-mentioned results of the Internal Kappa Consistency Test, the consistency between the smear microscopy and culture methods among the three test methods was weak. However, the results of the PCR were observed to be very inconsistent with those of the other two methods. However, it was speculated that if the microscopy, culture, and PCR methods were
combined for diagnosis procedures, the diagnosis efficiency could be significantly improved. The calculation results are shown in the last row of Table 5.

Table 4: The kappa consistency test for the three diagnostic tests.

| Group                | kappa |
|----------------------|-------|
| Smear vs. Culture    | 0.054 |
| Smear vs. PCR        | -0.111|
| Culture vs. PCR      | -0.486|

Table 5: Diagnostic value indicators of each test method.

| Test method   | Sensitivity (Sn) (%) | Specificity (Sp) (%) | Accuracy (%) |
|---------------|----------------------|----------------------|--------------|
| Smear microscopy | 40.7                 | 100.0                | 66.7         |
| Culture       | 37.0                 | 100.0                | 64.6         |
| PCR           | 55.6                 | 100.0                | 77.1         |
| Combined method | 92.6                 | 100.0                | 95.8         |

Discussion

The molecular biological diagnosis method (represented by PCR) is found to be more sensitive and rapid than the traditional culture method. It could directly detect the nucleic acid molecules of the pathogens without the steps of isolation and culturing, which proved to be helpful for the early diagnoses of the disease [8]. It is known that PCR can be theoretically detected regardless of whether the pathogen are active or inactive. The positive results of the PCR were significantly greater in number than those of the culture method. However, there were no significant differences observed when compared with the smear microscopy (p > 0.05). The reason was that the IPA patients had received antifungal therapy; therefore, the positive rates of the culture method were lower than PCR. This showed fungal culture method is less the diagnosis efficiency than PCR when antifungal therapy were conducted. However, we found that some patients were positive with BALF smear microscopy or fungal culture, but negative with PCR method. A possible cause of this is the fungal composition. The outer sides of Aspergillus (which belong to eukaryote) are wrapped by a layer of compact and hard cell wall components. The complete cell wall will inevitably reduce the extraction effects of nucleic acid substances [9].

Therefore, the interpretations of PCR results should be made in combination with other factors, such as clinical manifestations, other examination results. In our study, it was found that when smear microscopy techniques were performed for the lower sediment of BALF, the concentrations of pathogen components in those specimens were relatively high. The positive rates of smear microscopy were 40.0%, while that of common sputum was only 4.5% [10,11]. This showed the lower sediment of BALF was very high efficiency for microscopic examinations. In summary, the three examined diagnostic methods were found to have high specificity, but insufficient sensitivity. In the present study, considering the low degree of overlap in the results of the three detection methods, it was speculated that if a combination of microscopy, culture, and PCR methods was applied in diagnosis processes.

Conclusion

The pathogenic detections of bronchoalveolar lavage fluid (BALF) have been determined to be an ideal diagnostic method for Aspergillus infections. Because of the limitations of the different detection methods, The combination of BALF smear microscopy, culture, and PCR could significantly improve IPA diagnosis efficiency.

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Conflicts of Interest Statement

No.

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