Expression of surfactant protein-A in exhaled breath condensate of patients with chronic obstructive pulmonary disease

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Abstract. Pulmonary surfactant protein A (SP-A) has been associated with host defense in the lung, and contributes to the pathogenesis of chronic obstructive pulmonary disease (COPD). The present study aimed to determine a non-invasive method of measurement of SP-A, and further examine the expression levels of SP-A in patients with COPD. SP-A was detected in the exhaled breath condensate (EBC) obtained from patients with COPD and from non-COPD subjects. The individuals recruited for the present study comprised 60 subjects with and without COPD, who underwent lobectomy for a solitary peripheral lung nodule. EBC was collected using a condenser, and an enzyme-linked immunosorbent assay (ELISA) was used to measure the levels of SP-A. Tissue samples were obtained during lobectomy through resection of the adjacent lung tissues, located >5 cm from the nodule. Western blot analysis and immunohistochemistry were used to measure SP-A and SP-A-positive type II pneumocytes. The results demonstrated that SP-A was detectable in the EBC of all subjects. The results of the ELISA and western blotting demonstrated that the expression levels of SP-A were significantly decreased in patients with COPD, compared with the non-COPD subjects. The reduction of SP-A-positive type II pneumocytes was associated with the expression levels of SP-A. Decreased expression levels of SP-A in EBC were associated with a higher degree of airway limitation. These results suggested that the measurement of SP-A levels in the EBC may serve as a method for monitoring airway obstruction in patients with COPD. Further investigations are required in order to examine these observations further and to elucidate the underlying mechanisms.

Introduction

Chronic obstructive pulmonary disease (COPD) is one of the leading causes of morbidity and mortality worldwide, characterized by poorly reversible airflow limitation (1,2). The pathophysiology of COPD remains to be fully elucidated, which limits the development of more effective therapies and novel drugs for COPD. Various methods have been used for diagnosing, monitoring and evaluating COPD, predominantly comprising indirect methods, including lung function tests (3) and imaging techniques (high resolution computed tomography) (4), and more direct evaluations using invasive measurements (bronchoscopy and induced sputum) (5,6). Indirect and direct evaluation methods have adverse effects through exposure to noxious agents and the invasiveness of the sampling procedures, respectively (7). Therefore, the identification of biomarkers for COPD has received increasing attention, aiming to reflect the degree of disease severity with non-invasive measurements, and to assist in monitoring pharmacological therapy (8). Exhaled air and blood analysis may be used to develop biomarkers of lung diseases.

Exhaled breath condensate (EBC), obtained through cooling exhaled air during spontaneous breathing, offers promising real-time measurement of pulmonary pathobiology. The advantages of EBC include easy and non-invasive collection from patients using a portable device, and its suitability for sequential and longitudinal sampling of the lower respiratory tract (9,10).

Pulmonary surfactant proteins, defined as SP-A, SP-B, SP-C and SP-D, are synthesized by type II pneumocytes, and SP-A and SP-D have been associated with the host defense of the lung (11-13). SP-A and SP-D lectins have been reported to contribute to the pathogenesis of COPD (14,15), and abnormal inflammatory responses and the involvement of inflammatory cells and cytokines may affect the composition and function of surfactant proteins (16). The present study focused on SP-A, which functions in protecting the lungs from oxidants and inflammatory and infectious stress (17,18). The immunomodulatory protein, SP-A, in bronchoalveolar lavage (BAL) fluid has also been reported to be associated with impaired
functions in chronic smokers (19,20). The preoperative detection methods of SP-A are limited, and predominantly include blood analysis and BAL fluid analysis (18-21). However, blood analysis only indirectly reflects the state of the lung, and BAL fluid analysis cannot be used as a routine procedure due to the invasiveness of the sampling procedure (22). Therefore, a novel non-invasive method, which is easy to perform, is required to measure the expression levels of SP-A and to routinely monitor the severity of COPD.

The present study aimed to measure levels of SP-A in the EBC of patients with COPD, to evaluate whether the expression levels of SP-A in EBC are associated with those in the lung tissue, and to assess whether SP-A in EBC is associated with lung function.

Materials and methods

Subjects. The population recruited for the present study comprised 60 subjects who underwent lobectomy for a solitary peripheral lung nodule between September 2012 to January 2013. All patients were recruited from the Tianjin Chest Hospital (Tianjin, China). Patients who had lung infection or acute COPD exacerbation within the month prior to commencement of the present study were excluded. Patients suffering from severe diseases of the heart, brain, liver or kidney were also excluded.

The diagnosis of patients with COPD was performed, based on the following clinical symptoms: History of exposure to risk factors and laboratory examination. The severity of COPD was assessed according to the American Thoracic Society and European Respiratory Society Consensus Statement (23), where post-bronchodilator forced expiratory volume (FEV) in 1 sec/forced vital capacity ≤70%, and an FEV of 1-80% confirmed the presence of airflow limitation that was not fully reversible. The 60 subjects were divided into two groups, according to pirometric evaluation: A COPD group (28 cases) and a non-COPD group (32 cases) (Table I).

Written informed consent was obtained from all subjects, using a Surfactant Associated Protein A kit (Cloud-Clone Corp, Houston, TX, USA). Briefly, dilutions (100 µl each) of standard, blank or EBC samples were added to the appropriate well and incubated for 2 h at 37°C. Subsequently, liquid from each well was removed, detection reagent A (100 µl) was added to the wells of a micro-titer plate and incubated at 37°C in the dark for 60 min. Following washing with 350 µl of 1X wash solution, detection reagent B (100 µl) was added to the wells, and the plate was incubated for a further 30 min at 37°C in the dark. Subsequently, 90 µl substrate solution was added to the wells and incubated for 15-25 min at 37°C in the dark. The reaction was terminated following the addition of 50 µl stop solution, and the absorbance was measured at 450 nm using a microplate reader 550 (Bio-Rad Laboratories, Inc., Hercules, CA, USA).

Western blot analysis. The expression levels of SP-A were detected using western blot analysis, following standard procedures. Briefly, fresh lung tissues were homogenized in radioimmunoprecipitation assay buffer (Beyotime Institute of Biotechnology, Shanghai, China) and protein concentrations were determined using the Bradford assay (Bio-Rad Laboratories, Inc.). Equal quantities of protein (40 µg) were separated by 12% SDS-PAGE (SDS-PAGE Gel Preparation kit; Beyotime Institute of Biotechnology) and transferred to nitrocellulose blotting membranes (Schleicher & Schuell Bioscience GmbH, Dassel, Germany). The membranes were then blocked by incubating with 10% nonfat dried milk overnight at 4°C. Subsequently, the membranes were incubated with mouse anti-human SP-A monoclonal antibody (1:500; cat. no. PE-10; Fuzhou Maixin Biotech Co., Ltd.) or mouse anti-actin antibody (1:1,000; cat. no. MAB1501; EMD Millipore, Billerica, MA, USA) at room temperature for 1 h, followed by incubation with horseradish peroxidase-conjugated rabbit anti-mouse secondary antibody (1:1,000; cat. no. sc-358914; Santa Cruz Biotechnology, Inc., Dallas, TX, USA). All experiments were repeated in triplicate. The protein expression levels were quantified using Photoshop Image Analysis software (Image-Pro Plus 6.0; Media Cybernetics Inc., Rockville, MD, USA).

Immunohistochemistry. For immunohistochemical analysis, sections were deparaffinized and rehydrated, and antigen retrieval was achieved by boiling (92-98°C) in citric acid buffer (pH 6.0) for 20 min. Immunostaining of SP-A and thyroid transcription factor-1 (TTF-1) were performed using mouse anti-human SP-A monoclonal antibody (1:100; cat. no. PE-10; Fuzhou Maixin Biotech Co., Ltd.) and anti-TTF-1 monoclonal

Table I. Patient characteristics.

<table>
<thead>
<tr>
<th>Characteristic</th>
<th>COPD</th>
<th>Non-COPD</th>
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<tbody>
<tr>
<td>Total cases (n)</td>
<td>28</td>
<td>32</td>
</tr>
<tr>
<td>Male (n)</td>
<td>15</td>
<td>18</td>
</tr>
<tr>
<td>Age (mean, years)</td>
<td>58.71±11.22</td>
<td>61.81±8.33</td>
</tr>
</tbody>
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COPD, chronic obstructive pulmonary disease.
antibody (1:100; cat. no. MX011; Fuzhou Maixin Biotech Co., Ltd.), according to manufacturer's instructions. The sections were counterstained with Mayer's hematoxylin (Fuzhou Maixin Biotech Co., Ltd.). TTF-1 was used to count the total type II pneumocytes including the SP-A-positive type II cells on the adjacent serial sections.

Evaluation of the SP-A-positive type II pneumocytes was performed using a light microscope (BX50; Olympus Corporation, Tokyo, Japan) at x400 magnification (manual counting). A total of 20 microscopic fields on each slide were evaluated for all subjects, with the results expressed in cells/ml².

Statistical analysis. The results are expressed as the mean ± standard deviation. Differences between the groups of subjects were determined using Student’s t-test. Pearson’s correlation coefficients were used to evaluate the associations between variables. The statistical software package, SPSS v.17 (SPSS, Inc. Chicago, IL, USA), was used for analysis. P<0.05 was considered to indicate a statistically significant difference.

Results

SP-A in EBC and lung tissue samples. SP-A was detected in the EBC of all subjects. The results of the ELISA revealed that the expression levels of SP-A in the EBC were significantly decreased in the patients with COPD (4.165±2.950 pg/ml), compared with the non-COPD subjects (7.706±3.923 pg/ml; Fig. 1A). The protein expression levels of SP-A were subsequently examined in the lung tissue samples using western blot analysis. The protein expression levels of SP-A were significantly decreased in the patients with COPD (1,480.7±256.7), compared with the non-COPD subjects (1,954.1±312.2; Fig. 1B).

SP-A-positive type II pneumocytes. The alveolar macrophages and certain type II pneumocytes were SP-A-positive. The expression of TTF-1 was only detected in the type II pneumocytes, with the alveolar macrophages being TTF-1-negative. As type II pneumocytes and alveolar macrophages have morphological differences, immunostaining with TTF-1 and SP-A monoclonal antibodies was used to determine the total
The number of type II pneumocytes and SP-A-positive type II pneumocytes. Images of the SP-A immunostained tissues from a patient with COPD and a patient without COPD are shown in Fig. 2A and B, respectively. The images in Fig. 3 show...
representative TTF-1-immunostained images of lung tissue samples from a patient with COPD (Fig. 3A) and a patient without COPD (Fig. 3B). Although no significant difference was detected, the results demonstrated that the total number of type II pneumocytes was lower in the COPD group, compared with the non-COPD group, and the ratio of SP-A-positive type II pneumocytes (SP-A-positive type II pneumocytes/total type II pneumocytes, expressed as a percentage) was decreased in the COPD group, compared with the non-COPD group (Figs. 2-4).

Correlation between the expression levels of SP-A and EBC. As shown in Fig. 5A, the expression levels of SP-A in the EBC were correlated with the ratio of SP-A-positive type II pneumocytes/total type II pneumocytes \( r=0.278; P=0.026 \). Decreased expression levels of SP-A were associated with a decreased ratio of SP-A-positive type II pneumocytes. In addition, the ratio of SP-A-positive type II pneumocytes was correlated with the predicted FEV 1% (data not shown), suggesting that the increased degree of airway obstruction was associated with an increase in the SP-A ratio. The expression levels of SP-A in the EBC were also correlated with the predicted FEV 1% \( r=0.272; P=0.036 \), suggesting that the decreased expression levels of SP-A in EBC were associated with an increase in the degree of airway obstruction (Fig. 5B).

Discussion

The non-invasive measurement of airway obstruction in patients with COPD may prove useful for monitoring the disease. EBC analysis is a non-invasive method for detecting airway lining fluid composition, and the technique has numerous advantages, as it is easy to perform and less expensive in terms of equipment and personnel costs, compared with sampling BAL and induced sputum (25,26).

Controversial results have been reported regarding the expression levels of SP-A in the BAL or serum of patients with COPD. Decreased expression levels of SP-A have been reported in smokers who are otherwise healthy, elderly people and patients with COPD (19,20,27). Ohlmeier et al (28) reported increased expression levels of SP-A in lung tissue samples and the induced sputum of patients with COPD, compared with normal or fibrotic lung tissue samples. Vlachaki et al (21) also reported that decreased expression levels of SP-A in lung tissue samples of patients with COPD are associated with the pathogenesis of COPD.

In the present study, investigation to identify a non-invasive SP-A measurement technique and subsequent investigation of the expression levels of SP-A in patients with COPD were performed. The expression levels of SP-A were determined in EBC collected, which was collected from patients with COPD and non-COPD subjects. The results of the present study demonstrated that the expression levels of SP-A were significantly decreased in the patients with COPD, compared with the non-COPD subjects, and the expression of SP-A in the EBC was correlated with that in the lung tissue. Immunostaining of TTF-1 revealed that the total number of type II pneumocytes in patients with COPD was increased, compared with the number in the non-COPD subjects. The decrease in the total number of type II pneumocytes may be due to the extensive tissue damage in COPD (12), and has been reported in previous studies following lung injury (29,30). Type II pneumocytes proliferate during lung injury and chronic inflammatory states, including COPD, in order to produce type I pneumocytes and specific products, including SP-A, which are involved in lung defense (31).

The results of the present study demonstrated that the ratio of SP-A-positive type II pneumocytes to the overall type II pneumocytes was decreased in patients with COPD, compared with the non-COPD subjects. One possible explanation is that disease-induced epithelial destruction depletes the type II pneumocytes, which are responsible for the production of SP-A. Furthermore, the decreased expression levels of SP-A in EBC were associated with a higher degree of airway obstruction, suggesting that the measurement of SP-A in EBC may be a potential method for monitoring airway obstruction in patients with COPD.

In conclusion, the present study performed non-invasive SP-A measurement by detecting the expression levels of SP-A in EBC collected from patients with COPD. Decreased expression levels of SP-A in the EBC were observed in the patients with COPD, compared with the non-COPD subjects, and these decreased expression levels were also observed in the lung tissues. In addition, the total number of type II pneumocytes in the patients with COPD were decreased and the ratios of SP-A-positive type II pneumocytes were reduced, compared with those in the non-COPD subjects. The reduction in SP-A-positive type II pneumocytes was associated with the expression levels of SP-A. Finally, the results of the present study demonstrated that decreased expression levels of SP-A in the EBC were associated with a higher degree of airway obstruction. These results suggested that the measurement of SP-A in EBC may be a potential method for monitoring airway obstruction in patients with COPD. Further investigations are required in order to further examine these observations and elucidate the underlying mechanisms.

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