Elevated circulating PAI-1 levels are related to lung function decline, systemic inflammation, and small airway obstruction in chronic obstructive pulmonary disease

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Introduction
Chronic obstructive pulmonary disease (COPD) is a worldwide health care problem characterized by progressive airflow limitation, which is rarely reversible.¹ The World
Health Organization predicted that COPD would become the third leading cause of death and might move up to the fifth leading cause of disability-adjusted life years by 2030.\(^2,3\) Aside from airway inflammation and oxidative stress, recent studies suggested that small airway obstruction (SAO) caused by small airway remolding (SAR) along with aberrant extracellular matrix (ECM) deposition also contributed to the pathogenesis of COPD.\(^4\) On the other hand, more and more evidences showed that airway inflammation might also “spread” into the circulatory system and cause systemic inflammatory injuries and organ damage.\(^5\) However, arguments about how they work in COPD and what factors are involved have never stopped.

As we know, plasminogen activator inhibitor-1 (PAI-1) was first identified as an important PAI expressed by cultured bovine endothelial cells\(^6\) and many other different types of cells in various tissues.\(^7\) During the past decades, numerous clinical and laboratory studies have evaluated the various functions of this unique serpin, including regulating fibrinolysis in the process of thrombus formation, promoting ECM, contributing to atherosclerosis,\(^8\) bone remodeling,\(^9\) renal fibrosis,\(^9\) myocardial infarction, and diabetic vascular damage.\(^10\) Mostly, PAI-1 is found to act in the aforementioned roles by interacting with the urokinase-type activator (uPA) system, which is known for its contribution to the fibrinolysis in blood.\(^11\) Meanwhile, recent studies have found that uPA system, including uPA and urokinase-type activator receptor (uPAR), is involved in inflammation and tissue remolding.\(^12\) Interestingly, sputum levels of PAI-1 and soluble uPAR (suPAR) were found to be increased in COPD patients.\(^13,14\) These findings raised the problem of whether PAI-1 and uPA system is involved in the inflammatory response and tissue remolding in COPD? Unfortunately, there is no definite answer until now.

Therefore, we measured the serum levels of PAI-1 and suPAR in COPD patients and healthy subjects and analyzed if these two circulatory biomarkers were related to lung function decline, systemic inflammation, and SAO so as to explore the potential roles of PAI-1 and suPAR in COPD.

**Methods**

**Study protocol**

The study protocol conformed to the principles of the Declaration of Helsinki and was approved by the Ethic Committee of West China Hospital, Sichuan University, People’s Republic of China. All subjects gave written informed consents before the study. Between October 2013 and March 2014, naïve COPD patients and healthy controls without active pulmonary disorders were recruited from the Outpatient Department of West China Hospital. Every subject received a standard lung function test according to the American Thoracic Society and European Respiratory Society guidelines;\(^15\) briefly, all the subjects underwent three-time eligible prebronchodilation spirometry tests and postbronchodilation spirometry tests. Then, COPD was diagnosed prospectively for this study on the basis of the Global Initiative for Chronic Obstructive Lung Disease (GOLD) criteria.\(^1\) To be included in the study, patients had to show a ratio of forced expiratory volume in the first second to forced vital capacity (FEV\(_1/FVC\)) below 70% after bronchodilation and an increase in FEV\(_1\) below 12% after inhalation of β\(_2\)-agonist (200 mg salbutamol). Meanwhile, during the final 3 months, they had to be clinically stable. All the enrolled patients and controls had never received standard COPD-related therapies such as inhaled corticosteroids, bronchodilator, etc. Chest X-ray and electrocardiogram were performed to exclude the comorbidities such as pneumonia, pulmonary tuberculosis, lung cancer, cardiomegaly, myocardial ischemia, arrhythmia, and other cardiopulmonary disorders. Patients were also excluded if they had conditions known to affect serum levels of PAI-1, suPAR, or C-reactive protein (CRP), including hemorrhagic and clotting disorders, diabetes, stroke, acute infectious disorders, and so on.

**Collection of the basic information**

The basic information of enrolled subjects, including age, sex, height, weight, cigarette status, passive cigarette exposure, main complaints, and medical history, was collected by two individual investigators. Body mass index (BMI) was calculated using the formula weight/square of height. To avoid the contrived bias, the third investigator assessed this information and had it confirmed by the subjects, and all the investigators were blind to the spirometry reports of the subjects.

**Measurement of PAI-1, suPAR, CRP, TIMP-1, and MMP-9**

The COPD patients and healthy controls were requested to fast overnight from 22:00 the night before; the next morning venous blood samples were collected from the median cubital vein, and 1 hour later, serum was separated by centrifugation at 3,000 revolutions per minute for 10 minutes and stored at \(-80^\circ\text{C}\) until analysis. Serum levels of PAI-1, suPAR, CRP, tissue inhibitor of metalloproteinase-1 (TIMP-1), and Matrix metalloproteinase-9 (MMP-9) were detected using a Magnetic Luminex Screening Assay (R&D Systems, Minneapolis, MN, USA) on the Bio-Plex 200 (Bio-Rad, CA, USA) at the Department of Laboratory Medicine of West China Hospital. The detection sensitivities were 0.67 pg/mL for PAI-1, 43.2 pg/mL for suPAR, 116 pg/mL for CRP,
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3.42 pg/mL for TIMP-1, and 13.6 pg/mL for MMP-9. All the measurements were carried out strictly according to the manufacturer’s instructions, and technicians who performed tests were blinded to the clinical details of the subjects.

Statistical analysis
All the data are presented as mean ± standard deviation (SD). Differences between groups were statistically analyzed using one-way analysis of variance or chi-square test when appropriate. Then, correlations between different serum biomarkers and lung functions were tested using Pearson’s partial correlation test adjusting for age, sex, BMI, pack-years, and passive cigarette exposure status. Finally, multivariable linear analysis was conducted to confirm the aforementioned relationships. The threshold of significance was set at two-side 5%. Data were analyzed, and figures were created with GraphPad Prism 6.01 for Windows (GraphPad Software Inc, La Jolla, CA, USA).

Results
Clinical characteristics of subjects
We enrolled 43 COPD smokers, 41 COPD nonsmokers, 18 healthy smokers, and 33 healthy nonsmokers. According to the GOLD standard, among these 84 COPD patients, there were 44 GOLD 1, 35 GOLD 2, 4 GOLD 3, and 1 GOLD 4 patients. Table 1 lists the demographic and clinical characteristics including age, sex, BMI, smoking status, pack-years, passive smoke exposure, lung function data, and serum levels of PAI-1, suPAR, and CRP. No significant difference in age, BMI, and passive smoke exposure was observed among these four groups. However, irrespective of COPD group or the healthy group, almost all the smokers were males. Lung function data illustrated that COPD patients were suffering from more serious SAO as their maximum midexpiratory flow (MMEF)/Pre and FEV1/FVC were largely decreased when compared with healthy controls.

Circulatory PAI-1 but not suPAR was upregulated in COPD patients
First, we compared PAI-1 and suPAR levels between COPD patients and healthy subjects. As listed in Table 1, serum PAI-1 levels in COPD patients was significantly increased in comparison with healthy subjects (125.5±51.74 ng/mL versus 102.98±36.62 ng/mL, P<0.007); more specifically, COPD smokers had higher levels of PAI-1 compared with healthy nonsmokers, healthy smokers, and COPD nonsmokers (136.16±52.61 ng/mL versus 99.29±37.14 ng/mL, 109.74±35.69 ng/mL, and 114.43±49.00 ng/mL, P<0.05, respectively, Figure 1A). Similarly, serum levels of CRP in COPD smokers were the highest among these four groups.

Table 1 Characteristics of healthy volunteers and COPD patients

<table>
<thead>
<tr>
<th>Characteristics</th>
<th>Healthy controls (N=51)</th>
<th>COPD (N=84)</th>
<th>P-value</th>
</tr>
</thead>
<tbody>
<tr>
<td></td>
<td>Nonsmokers (HN, 33)</td>
<td>Smokers (HS, 18)</td>
<td>Smokers (CN, 41)</td>
</tr>
<tr>
<td>Age, years</td>
<td>59±21±10.64</td>
<td>56.94±12.52</td>
<td>60.27±9.32</td>
</tr>
<tr>
<td>Sex, male/female</td>
<td>10/23</td>
<td>17/1</td>
<td>12/29</td>
</tr>
<tr>
<td>BMI, kg/m²</td>
<td>23.04±5.40</td>
<td>20.11±7.75</td>
<td>23.50±2.79</td>
</tr>
<tr>
<td>Current smokers (%)</td>
<td>NA</td>
<td>14 (77.8)</td>
<td>NA</td>
</tr>
<tr>
<td>Former smokers (%)</td>
<td>4 (22.22)</td>
<td>NA</td>
<td>NA</td>
</tr>
<tr>
<td>Pack-years for ever-smokers</td>
<td>NA</td>
<td>23.01±13.91</td>
<td>NA</td>
</tr>
<tr>
<td>Passive smoke exposure (%)</td>
<td>11 (33.3)</td>
<td>5 (27.78)</td>
<td>10 (24.39)</td>
</tr>
<tr>
<td>FEV1, L</td>
<td>2.28±0.84</td>
<td>3.11±0.56</td>
<td>1.68±0.50</td>
</tr>
<tr>
<td>FVC, L</td>
<td>2.99±0.95</td>
<td>4.04±0.62</td>
<td>2.87±0.62</td>
</tr>
<tr>
<td>FEV1/FVC, %</td>
<td>81.38±5.68</td>
<td>77.39±5.51</td>
<td>58.03±10.88</td>
</tr>
<tr>
<td>FEV1/Pre, %</td>
<td>110.0±19.05</td>
<td>111.4±15.05</td>
<td>79.37±21.31</td>
</tr>
<tr>
<td>FEV1/FVC, %</td>
<td>95.00±3.68</td>
<td>92.17±3.91</td>
<td>79.49±9.92</td>
</tr>
<tr>
<td>MMEF25–75, L/s</td>
<td>2.17±0.95</td>
<td>2.70±1.04</td>
<td>0.73±0.35</td>
</tr>
<tr>
<td>MMEF25–75/Pre, %</td>
<td>70.57±23.84</td>
<td>77.69±22.74</td>
<td>24.66±11.59</td>
</tr>
<tr>
<td>PAI-1, ng/mL</td>
<td>99.29±37.14</td>
<td>109.74±35.69</td>
<td>114.43±49.00</td>
</tr>
<tr>
<td>suPAR, pg/mL</td>
<td>697.26±366.34</td>
<td>698.12±339.84</td>
<td>797.06±412.79</td>
</tr>
<tr>
<td>CRP (μg/mL)</td>
<td>1.08±1.12</td>
<td>1.06±1.19</td>
<td>1.91±2.50</td>
</tr>
</tbody>
</table>

Notes: Data are presented as mean ± standard deviation; χ² test was used to test the significance of the difference between two groups; *these characteristics are significantly different between groups; **represents significance for CS versus HN; *for CS versus HS; †for CS versus CN. P<0.05 was considered to be statistically significant.

Abbreviations: COPD, chronic obstructive pulmonary disease; BMI, body mass index; kg/m², kilogram per square meter; FEV₁, forced expiratory volume in the first second; FVC, forced vital capacity; MMEF, maximum midexpiratory flow; PAI-1, plasminogen activator inhibitor-1; suPAR, soluble urokinase-type activator receptor; CRP, C-reactive protein; NA, not applicable.
PAI-1 (justified) 0.001 – 0.002 0.002 A
suPAR (justified) 0.817 0.032

Besides, a positive relationship was observed between PAI-1 levels and MMP-9 levels (justified r = 0.602, p = 0.132). However, serum suPAR levels showed no difference among these different patients and healthy subjects (p = 0.829).

PAI-1 levels were correlated with lung function and systemic inflammation

Next, we studied the relationship between PAI-1 and COPD-related parameters (Table 2). As can be seen from Table 2, serum PAI-1 levels were inversely related to FEV₁/FVC (justified r = -0.308, p < 0.001, Figure 2A) and FEV₁/Pre (justified r = -0.295, p = 0.001, Figure 2B) after adjusting for age, sex, BMI, pack-years, and passive cigarette exposure status. Meanwhile, PAI-1 levels showed a positive relationship with CRP levels (justified r = -0.351, p < 0.001, Figure 3A).

Serum PAI-1 levels reflected SAO and SAR

Also, from the data given in Table 2, we observe inverse correlations between PAI-1 levels and spirometry index of SAO, including the ratio of FEV₁/FVC (justified r = -0.289, p = 0.001) and the ratio of MMEF₂₅–₇₅/Pre (justified r = -0.273, p = 0.002). Furthermore, PAI-1 levels were positively related to TIMP-1 levels (justified r = 0.498, p < 0.001, Figure 3B) and MMP-9 levels (justified r = 0.267, p = 0.002). Besides, a positive relationship was observed between PAI-1 levels and suPAR levels (justified r = 0.210, p = 0.016).

Table 2 Correlations between serum PAI-1 levels, suPAR levels, and COPD-related characteristics in all subjects

<table>
<thead>
<tr>
<th>Parameter</th>
<th>PAI-1 (justified)</th>
<th>suPAR (justified)</th>
</tr>
</thead>
<tbody>
<tr>
<td>FEV₁/FVC</td>
<td>r = -0.308</td>
<td>p = 0.014</td>
</tr>
<tr>
<td></td>
<td>p &lt; 0.001</td>
<td>0.873</td>
</tr>
<tr>
<td>FEV₁/Pre</td>
<td>r = -0.295</td>
<td>p = 0.001</td>
</tr>
<tr>
<td></td>
<td>p = 0.132</td>
<td>0.817</td>
</tr>
<tr>
<td>FEV₁/FVC</td>
<td>r = -0.289</td>
<td>p = 0.021</td>
</tr>
<tr>
<td></td>
<td>p = 0.817</td>
<td></td>
</tr>
<tr>
<td>MMEF₂₅–₇₅/Pre</td>
<td>r = -0.273</td>
<td>p = 0.002</td>
</tr>
<tr>
<td></td>
<td>p = 0.602</td>
<td></td>
</tr>
<tr>
<td>CRP</td>
<td>r = 0.351</td>
<td>p = 0.188</td>
</tr>
<tr>
<td></td>
<td>p &lt; 0.001</td>
<td>0.032</td>
</tr>
<tr>
<td>TIMP-1</td>
<td>r = 0.498</td>
<td>p = 0.456</td>
</tr>
<tr>
<td></td>
<td>p &lt; 0.001</td>
<td>&lt; 0.001</td>
</tr>
<tr>
<td>MMP-9</td>
<td>r = 0.267</td>
<td>p = 0.133</td>
</tr>
<tr>
<td></td>
<td>p = 0.002</td>
<td>0.132</td>
</tr>
<tr>
<td>suPAR</td>
<td>r = 0.210</td>
<td>p = 0.016</td>
</tr>
<tr>
<td></td>
<td></td>
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</tr>
</tbody>
</table>

Notes: The value of r represents the Pearson’s partial correlation coefficient adjusted for age, sex, BMI, cigarette status, and passive smoke exposure (in the column titled PAI-1 (justified)). P < 0.05 was considered to be statistically significant. Data in bold indicates statistical significance.

Abbreviations: PAI-1, plasminogen activator inhibitor-1; COPD, chronic obstructive pulmonary disease; FEV₁, forced expiratory volume in the first second; FVC, forced vital capacity; FEV₁/FVC, ratio of FEV₁ to FVC; FEV₁/Pre, FEV₁ percentile predicted; CRP, C-reactive protein; TIMP-1, tissue inhibitor of metalloproteinase-1; MMP-9, Matrix metalloproteinase-9; BMI, body mass index.
Serum suPAR were related to CRP and TIMP-1
On the other hand, serum suPAR levels showed positive correlations with CRP levels \( (\text{adjusted } r=0.188, P=0.032) \) and TIMP-1 levels \( (\text{adjusted } r=0.456, P<0.001) \). However, it showed no relationship with lung function parameters and MMP-9 levels.

Multivariable linear analysis
Finally, we performed a multivariable linear analysis for circulating PAI-1 levels and found that age, FEV\(_1\)/FVC, CRP, and TIMP-1 were the independent parameters associated with PAI-1, as listed in Table 3.

Discussion
Our main goal was to explore the role of PAI-1 and suPAR in COPD. First, we measured and compared the serum PAI-1 and suPAR levels in healthy nonsmokers, healthy smokers, COPD nonsmokers, and COPD smokers. Then, we analyzed the correlations between serum PAI-1, suPAR levels, and indicators of lung function decline, systemic inflammation, and SAO. Finally, we found that serum PAI-1 levels were significantly increased in COPD patients, especially COPD smokers, and serum PAI-1 levels were related to lung function parameters such as FEV\(_1\)/predicted and FEV\(_1\)/FVC, systemic inflammation indicators represented by serum CRP, and SAO parameters including FEV\(_3\)/FVC, MMEF\(_25-75\)/predicted, serum...
TIMP, and serum MMP-9. On the other hand, however, serum suPAR levels only showed association with CRP and TIMP-1 levels. These findings implied that PAI-1 might be involved in the inflammation responses and airway remodeling process in COPD.

First, we found that serum PAI-1 level, but not suPAR, was increased significantly in COPD patients, especially COPD smokers compared with other subjects, and negatively related to FEV\textsubscript{1}/Pre and FEV\textsubscript{1}/FVC. As a serpin with multiple functions, PAI-1 was first identified as an inhibitor of plasminogen activation system.\textsuperscript{16} By interacting with uPAR, PAI-1 showed profibrogenic properties and acted as a meaningful serum biomarker in various pulmonary diseases such as pulmonary fibrosis\textsuperscript{17} and asthma.\textsuperscript{18} Previous studies illustrated that cigarette smoke exposure could apparently increase the expression of PAI-1 in mice,\textsuperscript{19} and sputum PAI-1 levels are significantly increased in COPD patients.\textsuperscript{11} Thus, our finding confirmed that PAI-1 might be related to cigarette smoke exposure in COPD. Besides, we found that serum PAI-1 was positively related to suPAR, which was consistent with the result of the previous study, which indicated that plasminogen activation system (especially uPA system) was invoked in COPD.\textsuperscript{11}

Then, our data showed a positive relationship between serum PAI-1 and serum CRP. Recently, it has been widely accepted that COPD-related inflammation was not only limited to the local airway, but could also “spread” to the whole body, which was termed systemic inflammation.\textsuperscript{20} As the most widely investigated biomarker of systemic inflammation, CRP was found to relate to airflow obstruction\textsuperscript{21} and disease severity of COPD.\textsuperscript{22} This was confirmed in our study by demonstrating that serum CRP was increased in COPD patients. Meanwhile, basic research indicated that PAI-1 regulated lipopolysaccharide-induced inflammation via targeting on the TLR4/NF-κB signaling pathway in alveolar macrophages,\textsuperscript{23} and elevated PAI-1 expression promoted alveolar epithelial cell apoptosis and exacerbated lung inflammation induced by influenza A virus.\textsuperscript{19} On the other hand, Chen et al\textsuperscript{24} found that CRP could increase the expression of PAI-1 via mediating oxidative stress and MAPK signal pathway in human coronary endothelial cells. Therefore, our findings suggested for the first time that PAI-1 may be associated with systemic inflammatory responses in COPD.

In addition, we observed that serum PAI-1 was related to SAO parameters, including MMEF\textsubscript{25–75}/Pre and FEV\textsubscript{1}/FVC, and SAR indicators such as TIMP-1 and MMP-9. As we know, SAR affecting bronchioles, especially respiratory bronchioles that were the transition zones between the airway and alveolar spaces, was the leading cause of SAO, which induces the persistent airflow limitation in COPD.\textsuperscript{25} More specifically, as the main mechanism of SAR, epithelial–mesenchymal transition (EMT) mainly involving MMP and TIMP families played vital roles in the pathogenesis of COPD.\textsuperscript{26} Previous studies have identified MMPs as a family of enzymes that were capable of degrading all components of the ECM involved in the EMT process\textsuperscript{27} and found that members of the MMP family were selectively inhibited by specific individuals of TIMPs. Particularly, TIMP-1 has been shown to bind to both the active and latent forms of MMP-9 and inhibit its enzymatic activity. Overall, TIMP-1 and MMP-9 regulated the balance between degradation and deposition of ECM, which were involved in airway remolding.\textsuperscript{28} On the other hand, clinical investigations showed that serum TIMP-1 and MMP-9 levels were markedly increased and significantly negatively related to FEV\textsubscript{1}/FVC or FEV\textsubscript{1}/% predicted in COPD patients.\textsuperscript{29,30} These studies identified TIMP-1 and MMP-9 as serum biomarkers of SAR and SAO in COPD. Therefore, the positive relationship between serum TIMP-1, MMP-9 and PAI-1 found in our study indicated that serum PAI-1 might be involved in the EMT and SAR process in COPD.

Besides, serum suPAR only showed positive correlations with serum TIMP-1 and CRP in our study, which is consistent with previous findings that increased uPAR expression in the small airway epithelium is involved in the EMT process in COPD patients.\textsuperscript{31} These findings implied that suPAR might act synergistically with PAI-1 in SAR and systemic inflammation in COPD.

Finally, we performed multivariable linear analysis for PAI-1 and found that FEV\textsubscript{1}/FVC, CRP, and TIMP-1 were the independent parameters associated with PAI-1, which confirmed the relationship between PAI-1 and lung function decline, systemic inflammation, and SAO in COPD.
In this study, we only enrolled 135 subjects because of the rigorous inclusion and exclusion criteria; this would limit the P-value in the correlation analysis. Meanwhile, we evaluated the relationship between PAI-1 levels and SAR only based on the correlation analysis with four indirect indicators TIMP-1, MMP-9, FEV/FVC, and MMEF25–75/Pre. To avoid these limitations, future studies should enroll more COPD and healthy subjects and collect more evidences of SAR, such as high-resolution computed tomography, biopsy, or other direct indicators.

**Conclusion**

In conclusion, our findings first revealed the hypothesis that PAI-1 might be associated with lung function decline, systemic inflammation, and SAO in COPD. A previous study showed that airway inflammation and chronic airway remodeling induced by ovalbumin were attenuated in PAI-1 gene knockout mice and wide-type mice receiving intratracheal administrations of small interfering RNA against PAI-1.\(^3\) In combination with this study, we suggested that PAI-1 might be introduced into clinical practice as a therapeutic target of COPD in the future. However, larger, in-depth clinical and laboratory studies are needed to confirm our findings and explore how elevated PAI-1 play roles in COPD.

**Acknowledgments**

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**Author contributions**

Hao Wang and Fuqiang Wen designed this research, and all authors contributed toward subjects recruit, data collection, statistical analysis, drafting, and critically revising the paper, gave final approval of the version to be published, and agree to be accountable for all aspects of the work.

**Disclosure**

The authors declare no conflicts of interests in this work.

**References**


