Platelet-activating factor mediates acid-induced lung injury in genetically engineered mice

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Adult respiratory distress syndrome (ARDS) is an acute lung injury of high mortality rate, and the molecular mechanisms underlying it are poorly understood. Acid aspiration–induced lung injury is one of the most common causes of ARDS, characterized by an increase in lung permeability, enhanced polymorphonuclear neutrophil (PMN) sequestration, and respiratory failure. Here, we investigated the role of platelet-activating factor (PAF) and the PAF receptor (PAFR) gene in a murine model of acid aspiration–induced lung injury. Overexpression of the PAFR gene in transgenic mice enhanced lung injury, pulmonary edema, and deterioration of gas exchange caused by HCl aspiration. Conversely, mice carrying a targeted disruption of the PAFR gene experienced significantly less acid-induced injury, edema, and respiratory failure. Nevertheless, the efficiency of PMN sequestration in response to acid aspiration was unaffected by differences in PAFR expression level. The current observations suggest that PAF is involved in the pathogenesis of acute lung injury caused by acid aspiration. Thus, inhibition of this pathway might provide a novel therapeutic approach to acute lung injury, for which no specific pharmaceutical agents are currently available.


Introduction

Adult respiratory distress syndrome (ARDS) is an acute lung injury with a mortality rate of 40–70% despite intensive care using currently available drugs. Acid aspiration–induced injury is one of the most common causes of ARDS. Potential mechanisms causing acid aspiration–associated lung injury include HCl-induced damage to the alveolar-capillary membrane, and polymorphonuclear neutrophil (PMN) adhesion, activation, and sequestration, leading to pulmonary edema and deterioration of gas exchange (1–3).

Platelet-activating factor (PAF) is a proinflammatory phospholipid mediator that has various biologic effects, including cell adhesion, endothelial cell activation, and the production of cytokines and eicosanoids via activation of G protein–coupled PAF receptor (PAFR) (4–13). Considering its potent biologic activity, PAF is potentially involved in the development of inflammatory disorders, including acute lung injury. To examine the pathophysiologic role of PAF in the pathogenesis of diseases, previous studies have depended on various types of PAFR antagonists, although there exist significant problems regarding the specificity of these agents. For example, some PAFR antagonists have inhibitory effects on histamine via interaction with its G protein–coupled receptor (14, 15). To elucidate the pathophysiologic role of PAF in vivo without using potentially nonspecific PAFR antagonists, we have recently developed genetically engineered mice (16, 17).

The purpose of this report was to investigate the exact role of PAF in a murine model of acid aspiration–induced lung injury. To address this question, we used 2 different mutant mice established in our laboratory: transgenic mice overexpressing the PAFR gene (PAFR-Tg) (16) and controls (PAFR-Con), and PAFR gene–disrupted mice (PAFR-KO) (17) and controls (PAFR-WT). We observed that overexpression of the PAFR gene exaggerated the acute lung injury induced by acid aspiration, whereas disruption of the PAFR gene significantly attenuated the injury, indicating that the PAFR, and presumably PAF, may mediate important features of acid-induced lung injury.

Methods

Mice. PAFR-Tg mice were established as described previously (16). Male PAFR-Tg mice were mated to BDF1 females. The genetic background was C57BL/6 × DBA/2 hybrid. Northern blot analysis demonstrated the expression of the PAFR transgene in trachea, lung, heart, skeletal muscle, eye, skin, and aorta (16).

PAFR-KO mice were established by gene targeting (17). Mice heterozygous for the PAFR mutant allele with the genetic background of the C57BL/6J × 129/Ola hybrid were mated. Offspring were genotyped at 4 weeks of age. For genotyping, genomic DNAs were isolated from biopsied tail and subjected to 35 cycles of PCR amplification (10 seconds at 98°C; 2 seconds at 65°C; 30 seconds at 74°C). The primers were as follows: forward 5’-TATGGCTGACCTGCTCTTCCTGAT-3’ and reverse 5’-TATTGGGCACTAGGTTGGTGGAGT-3’ for

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detecting the intact PAFR allele; and forward 5'-GCCT-GCTTGGCAATATCATGGTGGAAAAAT-3' and reverse 5'-GCGATGCCGACTCAGACGGGACGGCGGATA-3' for detecting the disrupted PAFR allele. The former set of primers amplified a 287-bp DNA fragment, and the latter PCR product consisted of 194 bp.

The animals were maintained on a light-dark cycle with light from 0900 to 2100 hours at 25°C. Mice were fed with a standard laboratory diet and water ad libitum. Mutant mice and their littermate controls (13–24 weeks old) were used in the current study.

Protocols. One minute before aspiration challenge, 2 deep inhalations (3 times tidal volume) were delivered to standardize volume history, and measurements were made as baseline. Then, anesthetized and mechanically ventilated mice received 2 mL/kg HCl (pH = 1.5) intratracheally, followed by a bolus of air (30 mL/kg). In the saline-treated group, animals received saline instead of HCl in the same manner and served as controls. In all groups, measurements were made at 30-minute intervals for 2 hours. In some animals, the observation period was extended up to 5 hours. To assess the development of lung injury physiologically, lung elastance (EL; a reciprocal of lung compliance) was measured as described previously (18–20). Briefly, we measured the tracheal pressure (Ptr), flow, and volume (V). EL and lung resistance (RL) were calculated by adjusting the equation of motion:

$$ EL = \frac{K}{V/dt} + RL $$

where K was a constant.

Assessment of respiratory failure. At the end of the experiment, blood samples for gas analysis were obtained from the left ventricle. The measurements of PaO₂, PaCO₂, and pH were then made to assess the extent of respiratory failure (blood gas analyzer, Compact 3; AVL Medical Systems, Schaffhousen, Switzerland).

Assessment of pulmonary edema. At the end of the experiment, the lung wet/dry weight ratios (W/D) were calculated to assess pulmonary edema. After the trapped blood was drained from the excised lungs, the lung wet weight was measured. Then, we heated the lungs at 90°C to constant weight in a gravity convection oven for 72 hours and weighed the residuum as the lung dry weight.

Bronchoalveolar lavage fluid. At the end of the experiment, bronchoalveolar lavage (BAL) was performed (1 mL of PBS × 5) in each group. After BAL fluid (BALF) was centrifuged at 450 g for 10 minutes, the total and differential cell counts of the BALF were determined from the cell fraction. The supernatant was stored at –70°C until the protein was determined. The concentration of protein was measured by Lowry’s method using BSA as a standard.

Histology. At the end of the experiment, the lungs were removed intact and frozen with liquid nitrogen (22). A constant transpulmonary pressure of 3 cm H₂O was maintained during freezing by delivering constant flow into the trachea. Frozen lungs were fixed in Carnoy’s solution (60% ethyl alcohol, 30% chloroform, and 10% acetic acid) at –70°C for 18 hours. Progressively increasing concentrations of ethanol at –20°C were then substituted for the Carnoy’s solution until 100% ethanol was reached. The tissue was maintained at –20°C for 4 hours, warmed to 4°C for 12 hours, and then allowed to reach and remain at room temperature for 2 hours. After fixation, the tissue blocks obtained from mid-sagittal slices of the lungs were embedded in paraffin. Blocks were cut 4-μm thick using a microtome. Slides were stained with hematoxylin-eosin.

Data analysis. Comparisons of data among each experimental group were carried out with ANOVA (Scheffe’s test). Data are expressed as mean ± SEM. P values less than 0.05 were taken as significant.

Table 1

<table>
<thead>
<tr>
<th>RL (Percent baseline)</th>
<th>PaCO₂ (mmHg)</th>
<th>pH</th>
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<tbody>
<tr>
<td>Control (saline)</td>
<td>100.0 ± 2.4</td>
<td>30.9 ± 3.7</td>
</tr>
<tr>
<td>Tg (saline)</td>
<td>101.5 ± 4.1</td>
<td>30.0 ± 4.0</td>
</tr>
<tr>
<td>Control (HCl)</td>
<td>120.7 ± 2.6</td>
<td>38.3 ± 3.1</td>
</tr>
<tr>
<td>Tg (HCl)</td>
<td>152.8 ± 4.0</td>
<td>58.8 ± 9.6</td>
</tr>
<tr>
<td>+/- (saline)</td>
<td>104.4 ± 1.2</td>
<td>31.5 ± 6.3</td>
</tr>
<tr>
<td>+/- (HCl)</td>
<td>128.9 ± 6.6</td>
<td>41.5 ± 4.0</td>
</tr>
<tr>
<td>+/- (saline)</td>
<td>108.4 ± 1.8</td>
<td>29.8 ± 2.0</td>
</tr>
<tr>
<td>+/- (HCl)</td>
<td>146.3 ± 4.0</td>
<td>59.8 ± 9.6</td>
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*P < 0.05 vs. saline-treated groups. **P < 0.05 vs. HCl-treated controls.

Figure 1

The time course of response in lung elastance (EL) in HCl- and saline-treated groups (n = 3 for each group). After HCl aspiration, all animals died at 2–2.5 hours in PAFR-Tg and at 3–3.5 hours in controls, whereas all animals survived in the PAFR-KO group.
Results

Figure 1 demonstrates the time course of changes in lung elasticity in HCl- and saline-treated groups. After HCl administration, the survival periods of PAFR-Tg and control mice were 2 hours and 3 hours, respectively, whereas PAFR-KO mice were all alive throughout the experiment and their survival was significantly prolonged ($P < 0.02$ vs. the other groups; $\chi^2$ analysis). In terms of changes in EL and RL, HCl-treated PAFR-Tg mice demonstrated a significantly greater response than the other groups, whereas the HCl-induced responses in PAFR-KO mice were significantly reduced compared with their WT controls (Figure 2 and Table 1).

HCl aspiration induced deterioration of gas exchange, which was not observed in the saline-treated groups. HCl-induced hypoxemia was most prominent in PAFR-Tg mice, whereas it was significantly attenuated in PAFR-KO mice (Figure 3). After HCl treatment, increases in PaCO$_2$ and decreases in pH were observed in PAFR-Tg mice, although there were no differences in PaCO$_2$ or pH levels between PAFR-KO mice and saline-treated groups (Table 1).

To assess the severity of lung edema, lung W/D was measured in each group (Figure 4). In control groups, HCl challenge significantly increased W/D compared with all saline-treated groups. The W/D in HCl-treated PAFR-Tg was significantly greater than that in any other group, whereas the W/D in HCl-treated PAFR-KO was markedly decreased and fell to virtually the same level as for the saline-treated groups.

HCl administration increased the amount of protein and the number of PMNs in BALF, indicating HCl-induced protein leakage and PMN infiltration. The amount of BALF protein was significantly greater in HCl-treated PAFR-Tg than that in its control, and HCl-induced protein leakage was significantly attenuated in PAFR-KO mice (Figure 5). Total cell counts and differential cell fractions are shown in Table 2. Of note, HCl aspiration elicited no significant differences in PMN infiltration between PAFR-Tg and its control or between PAFR-KO and PAFR-WT (Figure 6).

Figure 7 illustrates the lung histology after HCl administration. The HCl-induced lesions included alveolar thickening, distortion, protein leakage, and cellular infiltration and were most prominent in the PAFR-Tg mice. In contrast, the alveolar architecture was well preserved, and the histological changes were minimal in the PAFR-KO mice.

Discussion

The results of the current study suggest that the PAFR and its physiological ligand, PAF, are crucially involved in the pathogenesis of acid aspiration–induced lung injury. Overexpression of the PAFR gene enhanced pulmonary edema and deterioration of gas exchange caused by HCl aspiration. Disruption of the PAFR gene significantly attenuated the acid-induced edema and respiratory failure. However, PMN sequestration was not regulated by the PAFR gene. These observations indicate that PAF and the PAFR gene may mediate important features of acid-induced lung injury.

Aspiration of gastric contents is reported to be associated with a 26–36% incidence of ARDS (23, 24). Acute lung injury induced by massive acid aspiration is char-

Table 2

<table>
<thead>
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<th>Total cell counts and cell fractions in BALF 2 hours after aspiration in the experimental groups</th>
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<tbody>
<tr>
<td>Control (saline)</td>
</tr>
<tr>
<td>------------------</td>
</tr>
<tr>
<td>Total cell counts ($\times 10^5$)</td>
</tr>
<tr>
<td>Macrophages (%)</td>
</tr>
<tr>
<td>PMNs (%)</td>
</tr>
<tr>
<td>Lymphocytes (%)</td>
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$^A$P < 0.01 vs. saline-treated groups.
characterized by increased permeability in the lung, decreased lung compliance, enhanced PMN sequestration, and respiratory failure. Aspirated HCl may evoke direct damage to the alveolar-capillary membrane and promote PMN adhesion, activation, and sequestration. It has been reported that thromboxane (TX) synthesis and generation of oxygen radicals associated with PMN activation are potential mechanisms to induce pulmonary edema and respiratory failure (1, 2). It has been demonstrated that inhibition of either TX or H2O2 attenuates acid aspiration–induced increase in protein in BALF in rats (2, 25). After acid aspiration, there is generation of eicosanoids, including TXA2, from activated tissues (26, 27). It is speculated that activation and adhesion of PMNs might be the initial action, leading to generation of radical oxygen species and increases in permeability. The current findings suggest that PAF has a main role in PMN activation, but not in PMN sequestration. Potentially, leukotriene B4, a potent neutrophil chemotactant, has a dominant role in PMN sequestration (28). The current observation that neither overexpression nor disruption of the PAFR gene affected PMN sequestration may be explained by the possibility that acid-induced release of chemotactic mediators, including leukotriene B4, would not be affected by relative expression of the PAFR gene.

Thromboxanes and leukotrienes are potent mediators that are generated from arachidonic acid by cyclooxygenase and 5-lipoxygenase, respectively. A key enzyme for the production of these inflammatory mediators, including eicosanoids and PAF, is phospholipase A2. Especially, cytosolic PLA2 (cPLA2) is of importance, as it is activated by submicromolar concentration of Ca2+ and by phosphorylation via mitogen-activated protein kinases (29–33). There exists a possibility that cPLA2 per se is related to the formation of PAF (34), as the cPLA2 hydrolysis of alkyl-phospholipids produces lyso-PAF, which is further converted to PAF by the action of acetyltransferase. Potential interaction between PAF and eicosanoids may further upregulate the development of acute lung injury (20, 35).

After HCl treatment, the survival rate (100%) and levels of W/D, PaCO2, and pH in PAFR-KO mice were the same as for saline-treated controls. However, we observed that acid aspiration–induced alterations in EL, PaO2, BALF protein, and PMNs were not completely eliminat-
ed in PAFR-KO mice, although they were markedly attenuated. These observations suggest that factors other than PAF also play a role and contribute to physiological alteration, whereas PAF is closely related to the development of acute lung injury. It has been postulated that adhesion molecules and cytokines are also involved in this mechanism (22, 36). Considering that there are yet no drugs to manage pulmonary edema and improve survival rates, these mediators are possible targets to develop agents. The present study indicates that the specific intervention of the PAFR could make an important contribution to management of acid-induced lung injury, whereas overexpression of the PAFR gene exaggerated these phenomena. The current observations suggest that PAF might be involved in the pathogenesis of acute lung injury caused by acid aspiration. Taken together, specific antagonism of the PAFR might provide a novel and potential therapeutic approach to acute lung injury, for which no drugs are currently available.

**Figure 7**
Photomicrograph of lung tissues from PAFR-Tg (a and b), PAFR-WT (c and d), and PAFR-KO (e and f) mice 2 hours after HCl administration. Hematoxylin-eosin stain. Scale bar in a represents 200 μm in a, c, and e and 50 μm in b, d, and f.
Acknowledgments

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