PROTEOLYTIC CLEAVAGE OF EALFIN BY 20S PROTEASOME MAY CONTRIBUTE TO INFLAMMATION IN ACUTE LUNG INJURY

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ABSTRACT
Rationale We hypothesise that ealfin levels in acute lung injury (ALI) decrease over time due, in part, to proteolytic degradation as observed in other lung diseases.

Objectives The aim of this study was to characterise temporal changes in ealfin concentration in patients with ALI and to evaluate whether a decrease in ealfin levels is due to elevated protease activity.

Methods Bronchoalveolar lavage fluid (BALF) was obtained from patients with ALI within 48 h of onset of ALI (day 0), at day 3 and at day 7. Ealfin levels were quantified by ELISA. Ealfin susceptibility to proteolytic cleavage by ALI BALF was assessed by Western blot and by high-performance liquid chromatography–mass spectrometry.

Measurements and main results Ealfin levels were found to be significantly increased at the onset of ALI compared with healthy volunteers and fell significantly by day 7 compared with day 0. In contrast, levels of secretory leucocyte protease inhibitor did not decrease over time. This decrease in ealfin was due to cleavage by the 20S proteasome which was significantly increased in ALI BALF. Incubation of ALI BALF with the proteasome inhibitor epoxomicin confirmed that 20S proteasome protease activity was responsible for proteolytic cleavage of ealfin, resulting in diminished anti-elastase activity. In addition, free neutrophil elastase activity significantly increased in ALI BALF from day 0 to day 7.

Conclusions Ealfin concentrations fall within the pulmonary compartment over the course of ALI as a result of proteolytic degradation. This loss of ealfin may predispose people, in part, to excessive inflammation in ALI.

INTRODUCTION
Acute lung injury (ALI) is a major cause of morbidity and mortality in critically ill patients for which there is no effective pharmacological treatment.1 Ealfin is a 6 kDa serine protease inhibitor that has been isolated from lung secretions with concentrations in healthy lungs ranging from 1.5 to 4.5 μM and is induced by pro-inflammatory mediators such as interleukin 1β, tumour necrosis factor α, defensins, neutrophil elastase (NE) and lipopolysaccharide.2–6 Ealfin is released by proteolytic cleavage from its precursor protein, trappin-2, to form the C-terminal 57-amino-acid mature ealfin.7 8 The antiprotease activity of ealfin resides within this C-terminal domain with specificity for NE and proteinase 3 and a transglutaminase substrate binding motif (GQDPVK) is present at the N-terminus which allows it to cross-link extracellular matrix proteins.9 10 In addition, ealfin has antibacterial and immunomodulatory activities, which implicate it as an effector molecule in the host innate immune response.11–13

In the healthy lung, antiproteases are present at higher concentrations than their cognate proteases, thus providing the lung with a powerful anti-inflammatory screen. In ALI, this protease–antiprotease balance is tipped in favour of proteases leading to dysregulated extracellular protease activity.14 Consequently, this protease burden can damage the alveolar epithelial–capillary endothelial barrier resulting in the production of pulmonary oedema within the alveolar space.15 As a result of inflammation and elevated protease activity, protective antiprotease levels may be compromised. A decrease in plasma ealfin was found to correlate with altered ealfin gene expression and was associated with increased acute respiratory distress syndrome (ARDS) risk.16–18

As a result it is important to evaluate ealfin levels and activity in the lungs of patients with ALI as it has previously been demonstrated that elevated protease activity in certain lung diseases can degrade and inactivate host defence molecules such as ealfin which subsequently affects their biological properties.19–21 In this study, we investigated...
temporal levels of elafin in ALI bronchoalveolar lavage fluid (BALF) and correlated changes in levels to protease-mediated inflammation in ALI. We hypothesise that elafin levels in ALI decrease over time due, in part, to proteolytic degradation as observed in other lung diseases.

MATERIALS AND METHODS

 Acquisition of BALF
ALI BALF samples were acquired from the randomised clinical trial of Hydroxymethylglutaralyl-Coenzyme A Reductase Inhibition for ALI as described previously. BALF from healthy volunteers and from intubated and mechanically ventilated patients ‘at risk’ of ALI as described previously were used as control groups. ALI patient aetiology is outlined in the online supplement.

Determination of elafin, SLPI and proteasome levels in BALF
Elafin, secretory leukocyte protease inhibitor (SLPI) and proteasome levels in BALF were determined by ELISA as previously described.

Western blot analysis of recombinant elafin incubated with ALI BALF
Recombinant human elafin (50 ng) was incubated with 10 µl ALI BALF in tracheobronchial secretion in a final volume of 20 µl for 24 h at 37°C. In some experiments, ALI BALF was preincubated for 1 h at 37°C with a range of protease inhibitors, as indicated in the figure legends, before adding elafin. In other experiments, elafin and SLPI were incubated with 20S proteasome for 10 min–24 h at 37°C. Samples were separated by Tricine sodium dodecyl sulfate polyacrylamide gel electrophoresis (SDS-PAGE) on a 17.5% polyacrylamide gel and elafin and SLPI detected by Western blotting (see online supplement).

20S Proteasome and NE activity assay
20S Proteasome and NE activity in BALF were measured using the fluorogenic substrates LLVY-aminomethylcoumarin (LLVY-AMC) and Ala-Ala-Pro-Val-aminomethylcoumarin (AAPV-AMC), respectively (see online supplement).

HPLC mass spectrometry
Cleavage of elafin by 20S proteasome was assessed by incubating recombinant elafin (500 ng) with recombinant 20S proteasome (5 µg) in 20S buffer for 0–60 min. Samples were then analysed by reverse-phase high-performance liquid chromatography (HPLC) coupled with electrospray mass spectrometry as previously described. Experimental procedures are detailed in the online supplement.

Evaluation of the anti-elastase activity of 20S proteasome-cleaved elafin
To evaluate the antiprotease activity of 20S proteasome-cleaved elafin, we tested the ability of cleaved elafin to inhibit one of its target proteases, NE, using the substrate AAPV-AMC (see online supplement). Results are expressed as relative fluorescence units (RFUs).

STATISTICAL ANALYSIS
All data were analysed with GraphPad Prism V.5.0 (GraphPad Software Inc, San Diego, California, USA). Descriptive statistics of continuous variables were expressed as mean (±SEM) or median (IQR) depending on the normality of their distribution. Unpaired t test or Mann–Whitney test were used as appropriate for comparison between two groups. For comparison between three or more groups, data were analysed by means of one-way analysis of variance or Kruskal–Wallis test using Dunn’s multiple comparison test when significance was indicated (at the p<0.05 level). Statistical significance is presented as *p<0.05, **p<0.01 and ***p<0.001 in all figures.

RESULTS

Levels of elafin and SLPI in ALI BALF
As shown in figure 1A, BALF levels of elafin were significantly increased at the onset of ALI (30.66 ng/ml (16.61–56.01)) compared with healthy volunteers (0.5 ng/ml (0.37–0.67)). In addition, levels of elafin were significantly increased at the onset of ALI compared with the ventilated ‘at risk’ control group (0.87 ng/ml (0.44–1.4)). Levels of SLPI (figure 1B) were also significantly increased at the onset of ALI (105 ng/ml (53.5–224.4)) compared with the ventilated ‘at risk’ control group (0.1 ng/ml (0.06–0.15)) and healthy volunteers (0.12 ng/ml (0.08–0.15)). Temporal measurements of elafin were assessed in patients with ALI at the onset of ALI (day 0), day 3 and day 7 of disease. As shown in figure 1C, levels of elafin fell significantly by day 3 (12.73 ng/ml (6.89–39.3)) and day 7 (10.34 ng/ml (5.74–30.64)) compared with day 0 (30.66 ng/ml (16.61–56.01)). Unlike elafin, there was no significant change in the levels of SLPI by day 3 (95.67 ng/ml (58.01–243.9)) and day 7 (97.12 ng/ml (70.94–189.2)) compared with day 0 (105 ng/ml (53.5–224.4)) (figure 1D).

Effects of ALI BALF incubation on recombinant elafin integrity
Exogenous elafin was incubated with healthy control BALF and ALI BALF (figure 2). Elafin was resistant to proteolytic cleavage in healthy volunteer BALF with a band corresponding to intact elafin only detected in these samples (figure 2A). In contrast, exogenous elafin was susceptible to proteolytic cleavage in BALF with the presence of an upper band representing intact elafin and a lower band corresponding to cleaved elafin fragments (figure 2B).

Identification of protease family involved in cleavage of elafin in ALI BALF
ALI BALF was preincubated with a range of non-specific protease inhibitors targeting a number of protease families for 1 h before adding recombinant elafin. After 24 h of incubation at 37°C, samples were analysed by Western blot. As shown in figure 3A, preincubation with E64, leupeptin (cysteine protease inhibitors) and pepstatin (aspartic protease inhibitor) had no effect on elafin cleavage. In addition, preincubation with the specific matrix metalloproteinase (MMP) inhibitors (GM6001 and phosphoramidon) had no effect on elafin degradation, suggesting that an MMP is not involved in the proteolytic cleavage of elafin in ALI BALF. However, preincubation with the overlapping serine protease inhibitors, antipain and chymostatin, was found to successfully inhibit elafin degradation. Preincubation of ALI BALF with elastase inhibitors (SLPI, Alpha 1 Antitrypsin and MeOSuc-Ala-Ala-Pro-Val-chloromethylketone) did not inhibit degradation of elafin, suggesting that NE is not involved in ALI BALF-induced elafin cleavage (figure 3B). In addition, preincubation with anti-thrombin had no effect, suggesting that a number of the coagulation cascade proteases are not involved. Only the serine protease inhibitors TLCK and pefabloc successfully inhibited the proteolytic degradation of elafin, which indicated that a serine protease may be involved in the cleavage of elafin.
20S Proteasome is involved in mediating the proteolytic degradation of elafin

Overall, our findings indicated that a mixture of trypsin-like and chymotrypsin-like protease inhibitors abrogate the cleavage of elafin. The 20S proteasome exhibits trypsin-like and chymotrypsin-like enzymatic activity and biologically active 20S proteasome has been detected in the alveolar space of patients with ARDS and in the lungs in a rodent model of lung injury.25–27 To investigate if 20S proteasome was responsible for the cleavage of elafin, we examined whether epoxomicin—a highly specific proteasome inhibitor—could inhibit cleavage of elafin. As shown in figure 4A, inhibition of elafin cleavage was observed with all concentrations of epoxomicin. A time-course incubation of 40 ng exogenous elafin with 400 ng 20S proteasome was performed. As shown in figure 4B, incubation of recombinant elafin with active 20S proteasome resulted in the appearance of elafin cleavage fragments, generated in a rapid time-dependent manner with cleavage of elafin occurring just after 30 min with full-length elafin degradation observed after 120 min. In contrast, incubation of recombinant SLPI (100 ng) with 400 ng of 20S proteasome revealed only partial degradation of SLPI even after 24 h (figure 4C). Our results to this point suggest that 20S proteasome is involved in the proteolytic degradation of elafin by ALI BALF as active 20S proteasome cleaved recombinant elafin in vitro. In addition, the specific proteasome inhibitor, epoxomicin, inhibited the cleavage of elafin.

Measurement of 20S proteasome concentrations and activity in ALI BALF

20S Proteasome concentrations from patients with ALI and healthy subjects were determined by ELISA. As shown in figure 5A, 20S proteasome concentrations were significantly increased by day 0 (342±79 ng/ml) and day 3 (271±110 ng/ml) ALI BALF compared with healthy BALF (66±2 ng/ml). However, there was no significant difference between day 7 ALI BALF (75±17 ng/ml) and healthy BALF. 20S Proteasome activity levels in BALF from patients with ALI and healthy subjects were...
determined by cleavage of the fluorogenic substrate LIVY-AMC. As illustrated in figure 5B, 20S proteasome activity levels were significantly increased on day 0 (19053±6777 ΔRFU), day 3 (9171±7980 ARFU) and day 7 ALI BALF (4444±2309 ΔRFU) compared with healthy subjects (97±73 ΔRFU).

Analysis of 20S proteasome–elafin cleavage sites

Products from 20S proteasome–elafin incubations were analysed by HPLC and mass spectrometry. As before, elafin (500 ng) was incubated with 20S proteasome (5 μg) for 0, 15, 30 and 60 min. One component of substantial area was detected in the elafin/20S proteasome samples at 15 min and later (data not shown) but not in the 0 min control (figure 6A). The measured monoisotopic mass was 2463.33 Da, in agreement with that calculated for the P1-24 fragment of elafin. Tandem mass spectrometric sequencing of the peptide established that this peptide fragment was P1-24, confirming that the proteasome attacks elafin by hydrolysing the peptide bond between Ala24 and Met25, as illustrated in figure 6B. Another elafin fragment (P1-23) was also detected in our samples, although this was present at lower levels than P1-24, thus suggesting that the primary cleavage site is at Ala24–Met25.

Evaluation of the antiprotease activity of 20S proteasome-cleaved elafin

We evaluated the ability of 20S proteasome-cleaved elafin to inhibit NE activity. As before, elafin (100 ng) was incubated with 20S proteasome (1 μg) and then incubated with a fixed amount of NE (500 ng). NE activity was measured using the NE substrate AAPV-AMC. As illustrated in figure 7A, 20S proteasome-cleaved elafin lost its ability to inhibit NE activity in contrast to intact elafin, which completely abolished NE activity. 20S Proteasome inactivated with epoxomicin prior to incubation with elafin did not inhibit elafin’s anti-NE activity. Therefore, 20S proteasome-cleaved elafin loses its antiprotease activity against its target serine protease NE. We confirmed that 20S proteasome did not turn over the AAPV-AMC substrate (data not shown). We also evaluated free NE activity in ALI BALF. As shown in figure 7B, free NE activity significantly increased in ALI BALF on day 7 (590±319 RFU) compared with day 0 (92 ±38 RFU), indicating that proteasome-mediated cleavage of elafin correlates to the decreased anti-NE screen in the ALI lung over time.

DISCUSSION

ALI is associated with a high mortality rate of up to 40%. There is an urgent need to develop effective therapeutics for the treatment of this condition. The biological properties of elafin have been well described since its discovery, highlighting elafin as a potential therapeutic for the treatment of inflammatory lung conditions. Due to the multi-factorial nature of ALI, which involves a complex network of pro-inflammatory
cytokines and protease activities, protective host defence proteins such as elafin may be left vulnerable to enzymatic cleavage and inactivation. In keeping with previous reports, we showed elafin and SLPI levels to be significantly increased in patients with ALI compared with healthy volunteers.\textsuperscript{31} To ensure that this increase in antiprotease levels observed at the onset of ALI was not induced by mechanical ventilation, we measured elafin and SLPI levels in BALF from intensive care unit patients who were mechanically ventilated but did not fulfill criteria for ALI. Elafin and SLPI levels within this ‘at risk’ control group were similar to those found in healthy volunteers, confirming that the increase in elafin levels at onset of ALI was not as a result of mechanical ventilation.

We extended these findings further by assessing the temporal measurements of anti-protease levels in ALI BALF over the course of 7 days. In contrast to SLPI, elafin levels fell significantly by day 3 and 7 compared with baseline ALI BALF (day 0). To our knowledge, these data are the first to demonstrate a significant decrease in the temporal levels of elafin within the pulmonary compartment in patients with ALI. Recent temporal measurements in whole blood samples from patients with ARDS revealed levels of plasma elafin which declined with the clinical progress of ARDS.\textsuperscript{18} This decrease in plasma elafin was found to correlate with downregulated elafin gene expression in the acute stage of ARDS.\textsuperscript{17,18} It is therefore plausible that the decreased levels of elafin over time observed in ALI BALF may, in part, be mediated by decreased elafin expression. However, we now demonstrate that elafin is susceptible to proteolytic cleavage in the ALI lung.

Recent research suggests that the 20S proteasome is present and active in the extracellular space, although its function is as yet unclear.\textsuperscript{32,33} Extracellular, biologically active 20S proteasome has been detected in the alveolar space of patients with ALI\textsuperscript{25,26} and in a rodent model of lung injury.\textsuperscript{27} In addition, alveolar proteasome concentrations have been found to be markedly higher in patients with ALI, and in the circulation of patients with sepsis compared with healthy subjects.\textsuperscript{25,34} In agreement with previous data,\textsuperscript{25} we detected increased 20S proteasome concentrations in BALF obtained at the onset of ALI (day 0) compared with healthy BALF samples. Temporal analysis revealed that the levels of BALF proteasome decreased over the course of ALI so that by day 7 there was no significant difference compared with healthy BALF. In contrast, we detected significantly increased 20S proteasome activity levels in all ALI BALF compared with healthy subjects. However, similar to what was observed for concentrations, a temporal decrease in proteasome activity was also evident in ALI BALF. It is interesting to note the discrepancy between proteasome activity and levels in
Figure 7 20S Proteasome-cleaved elafin loses its antiprotease activity and free neutrophil elastase (NE) activity increases in acute lung injury (ALI) bronchoalveolar lavage fluid (BALF). (A) 20S Proteasome-cleaved elafin loses its ability to inhibit activity of the serine protease NE. 20S proteasome inactivated with epoxomicin prior to addition of elafin retained elafin’s anti-NE activity. NE activity was determined using the AAPV-AMC substrate and readings taken over time. Results are expressed as relative fluorescence units (RFUs). (B) NE activity in BALF was measured using AAPV-AMC substrate. The rate of substrate hydrolysis was monitored at 37°C over time and the results were expressed as the change (Δ) in RFU. NE activity increased significantly in day 7 ALI BALF compared with day 0 BALF. *p<0.05.

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healthy and day 7 ALI BALF. Despite similar levels of proteasome in both sets of samples, there is significantly more proteasome activity in day 7 ALI BALF compared with healthy control samples. On the basis of our results and the limited evidence to date, it is possible that a proteasome inhibitor may be present in healthy BALF.

Our data conflict with a previous study which reported a 17-fold decrease of proteasomal activity detected in patients with severe ARDS compared with healthy subjects despite concentrations of proteasome being significantly higher in ARDS BALF compared with healthy control BALF. This decreased activity in patients with ARDS was suggested to be due to the presence of an unknown inhibitor present in the alveolar space. To date, no extracellular proteasome inhibitors have been identified but the discovery of physiological inhibitors of the intracellular 20S proteasome suggests that it is possible for extracellular proteasome inhibitors to be present in the lung. Regardless, the finding that preincubation of ALI BALF with epoxomicin—a specific proteasome inhibitor—abolished proteasome activity and inhibited ALI BALF-induced cleavage of elafin indicates 20S proteasome activity is responsible for the proteolytic degradation of elafin. Further clarification of the role of 20S proteasome in mediating ALI BALF-induced proteolytic cleavage of elafin was achieved by incubating exogenous elafin with active 20S proteasome resulting in the appearance of elafin cleavage fragments generated in a time-dependent manner. In contrast to native, full-length elafin, these 20S proteasome-cleaved elafin fragments were unable to inhibit activity of NE. Pretreating ALI BALF with epoxomicin prevented the loss of elafin’s anti-elastase activity. Interestingly, we demonstrated that NE activity in ALI BALF significantly increased from day 0 to day 7, indicating that the proteasome cleavage of elafin may contribute in some part to increased NE activity in ALI. Free NE activity is well accepted as contributing directly to inflammation in ALI, resulting in microvascular injury, endothelial damage and interstitial oedema. To date, clinical trials utilising naturally occurring NE inhibitors have not been entirely successful due to problems with oxidative inactivation and proteolysis. However, elafin is currently in phase II trials for postoperative inflammation and may prove to be useful for treatment of ALI in future trials. It is possible that other mechanisms may contribute to this increased NE activity in day 7 ALI apart from proteasome-mediated elafin cleavage, including increased oxidation of other endogenous protease inhibitors such as α1 antitrypsin and SLPI or increased neutrophil necrosis. Further studies will be required to delineate fully the reasons for increased NE activity over time in the ALI lung.

In conclusion, this study shows elafin levels to decrease significantly over time in the pulmonary compartment of patients with ALI as a consequence of proteolytic degradation by the 20S proteasome. This decrease may contribute, at least in part, to increased NE activity in ALI over time. Further research is warranted to investigate the cellular source of origin of the 20S proteasome, its mechanism of transport into the extracellular space, regulation, expression, and above all, its biological role within the lung. The antiprotease deficiency as a result of cleavage by 20S proteasome suggests that antiprotease augmentation in patients with ALI may be clinically beneficial in attenuating the excessive unregulated NE activity that persists in ALI. Elafin’s multifunctional properties make it an attractive candidate as a therapeutic modality for ALI, although there is still a need for further research regarding its vulnerability to proteolytic degradation in vivo in conditions characterised by a protease burden such as ALI.
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