Coagulation factors in the airways in moderate and severe asthma and the effect of inhaled steroids

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ABSTRACT
Background: There is evidence of activation of the extrinsic coagulation cascade in the asthmatic airway, and both plasma and locally derived factors may be involved. The hypothesis that the normal haemostatic balance of healthy airways sampled by sputum induction favours fibrin formation in asthmatic airways, and that inhaled corticosteroids (ICS) and plasma exudation influence this balance, was tested.

Methods: ELISA and activity assays were used to measure α2-macroglobulin (an index of plasma leakage) and coagulation factors in hypertonic saline-induced sputum of 30 stable subjects (10 controls, 10 with moderate asthma and 10 with severe asthma). Additionally, the moderate cohort were weaned off their ICS, followed by further sputum induction 5 days after cessation of steroids.

Results: ICS wean induced a significant rise in plasminogen (median (interquartile range (IQR)): 13.92 (6.12–16.17) vs 4.82 (2.14–13.32) ng/ml, 95% CI 0.003 to 8.596, p = 0.0499) and tissue plasminogen activator (tPA; 5.57 (3.57–14.35) vs 3.88 (1.74–4.05) ng/ml, 95% CI 0.828 to 9.972, p = 0.0261) levels in sputum, such that tPA in untreated moderate asthma was significantly (p = 0.0029) higher than normal (2.14 (0.0–2.53) ng/ml). Subjects with severe asthma had significantly more α2-macroglobulin (p = 0.0003), tissue factor (p = 0.023), plasminogen activator inhibitor (p = 0.0091), thrombin-activatable fibrinolysis inhibitor (p = 0.0031) and fibrin degradation products (p = 0.0293) in their sputum than control subjects.

Conclusion: Untreated moderate asthma is associated with increased fibrinolysis that is corrected by ICS. Severe asthma and high dose corticosteroid therapy is associated with a profibrinogenic, antifibrinolytic environment in the airways. This study suggests that inhibition of fibrin deposition in severe asthma may be a therapeutic approach.

Asthma is characterised by airway inflammation and variable tissue remodelling.1 It is influenced by inhaled corticosteroid (ICS) therapy, yet many of those with asthma remain symptomatic, some with severe manifestations of the disease, as structural changes lead to airflow obstruction that may be irreversible.2 Plasma exudation from the bronchial microvasculature contributes to bronchial obstruction by fibrin deposition in mucus plugs, bronchial wall thickening, epithelial shedding, thickening of the basement membrane, hypertrophy of the smooth muscle and mucus secretion. Fibrin deposition in distal airways is also associated with increased bronchial hyper-responsiveness.3 We further reported that fibrin formation by bronchial epithelial cells in vitro is independent of plasma proteins, and is essential for bronchial epithelial repair.4 Exposure of plasma to cell-bound tissue factor (TF), the principal activator of the extrinsic coagulation cascade, initiates fibrin clot formation within minutes, although the rate of fibrin formation is determined by components of the intrinsic coagulation cascade.5 Thus, early, physiologically desirable, fibrin formation for normal wound healing, initiated by TF-bearing cells, may propagate into exudated bulk plasma, generating excessive and physiologically undesirable fibrin that could lead to fibrosis, mucus plug formation, airway narrowing and bronchial hyper-reactivity (BHR).5–6

Levels of TF in induced sputum are higher in patients with mild asthma compared with healthy controls,6 and after allergen challenge.10 A TF-dependent increase in coagulation and a plasminogen activator inhibitor (PAI-1)-dependent decrease in fibrinolysis is a feature of alveolar fibrin formation in acute respiratory distress syndrome (ARDS) and idiopathic pulmonary fibrosis.11 There has been recent interest in the role of platelets contributing to the coagulation response in patients with acute lung injury22; however, their role in the regulation of the extrinsic coagulation cascade in the lung, and in those with asthma is not known. Activation of the coagulation cascade has both fibrin-dependent and fibrin-independent influences on tissue fibrosis. Fibrin-dependent effects include neutralisation of surfactant, binding and accumulation of transforming growth factor β (TGFβ) and provision of a matrix for fibroblast proliferation. Independent effects include activation of serine proteases, notably thrombin, factor VIIa (FVIIa) and FXα, that activate protease-activated receptors (PARs), present on fibroblasts and bronchial epithelial cells, to stimulate fibrotic and inflammatory processes further in the lung6,11–13 and also to enhance TF expression further.14 Thus increased levels of TF are self-amplifying when coagulation is initiated.

In the normal lung, the haemostatic balance is antithrombotic and favours fibrinolysis.6 We sought to test the hypothesis that the normal haemostatic balance changes in favour of coagulation in moderate and severe asthma, and that ICS and plasma exudation influence this balance.

METHODS

Patients
Healthy controls and patients with moderate and severe asthma were recruited from the community and from the asthma clinic at Queen Alexandra Hospital, Portsmouth, UK. Asthma was confirmed...
by objective assessments of variable airflow obstruction and/or BHR, and together with lung function and exhaled nitric oxide measurements (FeNO), these are described in the online appendix. Subjects were 18–70 years old and had stable symptoms with no need for increased asthma treatment within 6 weeks of study entry. Current smokers, subjects with a smoking history of >10 pack years, other coexisting lung disease or significant comorbidity were excluded. “Moderate” asthma was defined as mild persistent disease with low or medium daily dose ICS (200–1000 μg beclomethasone dipropionate (BDP) or equivalent) under the Global Initiative for Asthma (GINA) criteria,15 with additional asthma medications permitted. “Severe” asthma was defined as GINA severe persistent disease with high daily dose ICS (>1000 μg of BDP or equivalent), with additional asthma drugs (including oral corticosteroids) permitted.

**Study design**

After initial screening and 2 week run in, patients underwent induced sputum and venous blood sampling. Furthermore, “moderate” subjects underwent tapered ICS withdrawal (200 μg BDP equivalent per week, see online appendix) 1 week after the first sputum induction. Continuation of long-acting β2 adrenoceptor agonists (LABAs) and reliever medications was permitted. Subjects completed twice-daily peak expiratory flow (PEF) (Wright mini peak flow meter, Clermont-Ferrand International, Harlow, UK) and diary card monitoring with weekly review until 5 days after complete steroid cessation when a second sputum induction was performed.

Ethical approval was granted by the Reading & Berkshire Research Ethics Committee, and all subjects provided informed consent. Sample size calculations are discussed in the online appendix.

**Sputum induction and processing**

Sputum induction was performed using an ultrasonic nebuliser (Devilbiss Ultraneb 2000, Sunrise Medical, UK) and hypertonic saline, and processed with 0.05% (w/v) dithiothreitol (DTT) (as described in the online appendix).

α2-Macroglobulin and coagulation factors in induced sputum

α2-Macroglobulin was used as a marker of plasma exudation16 17 and assayed by an in-house ELISA described in the online appendix. TF, tissue factor pathway inhibitor (TFPI), tissue plasminogen activator (tPA) and plasminogen activator inhibitor-1 (PAI-1) were analysed by sandwich ELISA using commercially available kits (American Diagnostica, Stamford, Connecticut, USA). Fibrinogen, FVII, FX, FXIII, plasminogen and thrombin-activatable fibrinolysis inhibitor (TAFI) were analysed by enzyme immunoassay using antibody pairs according to the manufacturer’s instructions (Affinity Biologicals, Ancaster, ON, Canada). D-dimers were assayed by sandwich ELISA (Hyphen Biomed, Andreys, France). Fibrin degradation products (DFPs) were measured using the Thrombo-Wellcotest (Remel Europe, Dartford, UK). Enzyme activity for thrombin and plasmin was analysed using specific substrates, S2238 and S2251, respectively (Chromogenix, Milan, Italy). Quantification was with the values for normal reference plasma (Precision Biologic, Dartmouth, NS, Canada). All standards were prepared with 0.05% (w/v) DTT.

**Statistical analyses**

The Kruskal–Wallis test compared median levels across the three independent subgroups. Comparisons between subgroups were made using a Mann–Whitney U test, and 95% CIs for the difference between medians calculated. Repeated measurements on the moderate asthma cohort were analysed using a paired t test. Associations between parameters were analysed using a pooled r value from the Spearman rank correlation coefficient after heterogeneity testing with a χ² test. Analysis was performed using GraphPad software, San Diego California, USA.

**RESULTS**

Baseline patient characteristics are shown in table 1. In the moderate group, one patient failed the second sputum induction (after withdrawal of steroids); therefore, nine participants remained in the moderate untreated group, with nine paired sputum samples (presteroid and poststeroid withdrawal) available for analysis.

**Blood and airway inflammation**

Sputum differential cell counts are presented in table 2 and blood results in online appendix table 1. Median values for sputum and blood eosinophilia were significantly higher in the severe group compared with the control group (sputum, 11.2 vs 0.0, 95% CI 1.52 to 31.24, p = 0.0001; blood, 0.6 vs 0.1, 95% CI 0.10 to 0.70, p = 0.0095). Patients with moderate treated asthma had significantly greater sputum neutrophilia than those with severe asthma (51.8 vs 18.2, 95% CI 10.0 to 45.6, p = 0.0054). ICS wean resulted in a non-significant increase in the proportion of sputum eosinophils relative to prewean values (1.97 vs 0.62, 95% CI −6.11 to 4.91, p = 0.404), but a significant sputum eosinophilia compared with controls (1.97 vs 0.00, 95% CI 0.76 to 10.4, p = 0.0015). FeNO levels did not differ between the groups and were not influenced by withdrawal of ICS (18 vs 28.5, 95% CI 3.50 to 29.5, p = 0.104: online appendix table 1).

Sputum α2-macroglobulin levels were significantly higher in the severe cohort compared with the other groups (control, 95% CI 741.3 to 4447.7, p = 0.0003; moderate treated, 95% CI 884.3 to 4676.0, p = 0.0002: fig 1 and online appendix table 2) and were not affected by ICS withdrawal (fig 1 and online appendix table 3). Over all groups there was a significant positive correlation with sputum eosinophils (r = 0.676, p<0.001: online appendix table 4).

Comparison of α2-macroglobulin levels in induced sputum with levels in reference normal plasma (1000 μg/ml) allowed a plasma dilution factor for the samples to be calculated. Thus, plasma proteins were present in induced sputum as a percentage of normal reference values that was: (mean (SD)) controls 0.031% (0.021%), moderate treated 0.015% (0.013%), moderate untreated 0.068% (0.135%) and severe 0.38% (0.45%). Knowledge of these values allowed us to consider the provenance, either plasma exudation or local cellular synthesis, of the coagulation factors present in the airways.

**The extrinsic coagulation cascade**

The full results for all assays are provided in online appendix tables 2 and 3, with a relevant summary provided below.

**Fibrinogenesis: TF, FVII and FX**

TF (fig 2A) levels were significantly higher in the severe cohort compared with controls (11.85 vs 3.95, 95% CI 1.49 to 16.05, p = 0.023) and treated moderate asthma (11.85 vs 5.13, 95% CI 1.3 to 16.21, p = 0.0167), with no significant difference after weaning ICS (5.915 vs 4.02, 95% CI −9.72 to 2.44, p = 0.102).
TF was significantly and positively correlated with sputum eosinophil counts ($r = 0.550$, $p = 0.008$).

FVII (fig 2B) levels were significantly higher in controls compared with the severe cohort (2.05 vs 0.00, 95% CI 0.091 to 3.045, $p = 0.025$) and those with untreated moderate asthma (2.05 vs 0.07, 95% CI 0.297 to 3.061, $p = 0.0029$). There was a significant reduction in FVII levels following withdrawal of ICS (1.14 vs 0.07, 95% CI 0.10 to 4.697, $p = 0.0429$; fig 2B, online appendix table 3). FX (fig 2C) levels were significantly decreased in severe asthma compared with controls (0.910 vs 6.00, 95% CI 0.65 to 9.46, $p = 0.0058$), with no effect following ICS withdrawal (2.62 vs 3.43, 95% CI 2.67 to 6.03, $p = 0.451$).

Fibrinolysis: plasminogen, tPA, plasmin, PAI-1, TAFI and FDPs

Plasminogen (fig 3A) was significantly lower in treated moderate asthma compared with controls (4.82 vs 12.99, 95% CI 2.62 to 13.62, $p = 0.0115$) and ICS withdrawal resulted in a significant increase in levels (4.82 vs 13.92, 95% CI 0.003 to 8.596, $p = 0.0499$).

Levels of tPA (fig 3B) were significantly higher in severe and untreated asthma cohorts when compared with controls (6.26 vs 2.14, 95% CI 1.599 to 11.975, $p = 0.0046$; and 5.57 vs 2.14, 95% CI 1.907 to 10.831, $p = 0.0015$, respectively), and there was a significant increase in levels following ICS withdrawal (3.88 vs 5.57, 95% CI 0.828 to 9.972, $p = 0.0261$; online appendix table 3).

Significantly higher levels of PAI-1, TAFI and FDPs (fig 3C,D,F) were observed in severe asthma compared with controls (95% CI 1.12 to 12.77, $p = 0.0091$; 0.450 to 6.751, $p = 0.0031$; and 0.01 to 30.1, $p = 0.0301$, respectively). Furthermore, TAFI was significantly higher in the severe cohort as compared with those with treated asthma (95% CI 0.852 to 8.049, $p = 0.0052$). In the severe group, FDPs levels correlated positively with sputum eosinophil count ($r = 0.667$, $p = 0.003$) and tPA ($r = 0.493$, $p = 0.006$).

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### Table 1  Patient characteristics

<table>
<thead>
<tr>
<th>Group</th>
<th>Control</th>
<th>Moderate</th>
<th>Severe</th>
</tr>
</thead>
<tbody>
<tr>
<td>No. of subjects</td>
<td>10</td>
<td>10</td>
<td>10</td>
</tr>
<tr>
<td>Sex (M/F)</td>
<td>6/4</td>
<td>3/7</td>
<td>5/5</td>
</tr>
<tr>
<td>Age (years)</td>
<td>36.0 (29.5–49.0)</td>
<td>56.0 (39.0–62.0)</td>
<td>54.0 (42.5–53.5)</td>
</tr>
<tr>
<td>Smoking history:</td>
<td></td>
<td></td>
<td></td>
</tr>
<tr>
<td>Ex</td>
<td>0</td>
<td>2</td>
<td>5</td>
</tr>
<tr>
<td>Never*</td>
<td>10</td>
<td>8</td>
<td>5</td>
</tr>
<tr>
<td>Pack years</td>
<td>0</td>
<td>0.75</td>
<td>2.1</td>
</tr>
<tr>
<td>Atopy (+/−)*</td>
<td>1/9</td>
<td>5/5</td>
<td>8/2</td>
</tr>
<tr>
<td>ICS dose (BDP equivalent, μg)†</td>
<td>–</td>
<td>400 (200–400)</td>
<td>2000 (1800–4000)</td>
</tr>
<tr>
<td>Oral prednisolone (mg daily)</td>
<td>–</td>
<td>–</td>
<td>4.0 (0–7.5)</td>
</tr>
<tr>
<td>Additional medications:</td>
<td></td>
<td></td>
<td></td>
</tr>
<tr>
<td>Long-acting β-agonist</td>
<td>–</td>
<td>6</td>
<td>8</td>
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<tr>
<td>Leukotriene receptor antagonist</td>
<td>–</td>
<td>1</td>
<td>3</td>
</tr>
<tr>
<td>Long-acting antimuscarinic</td>
<td>–</td>
<td>–</td>
<td>3</td>
</tr>
<tr>
<td>Oral theophylline</td>
<td>–</td>
<td>–</td>
<td>6</td>
</tr>
<tr>
<td>Oral antihistamine</td>
<td>–</td>
<td>–</td>
<td>4</td>
</tr>
<tr>
<td>FEV1% predicted‡</td>
<td>103 (92–117)</td>
<td>90 (87–108)</td>
<td>60.5 (55–79)</td>
</tr>
<tr>
<td>FEV1/FVC ratio‡</td>
<td>0.78 (0.71–0.82)</td>
<td>0.76 (0.74–0.8)</td>
<td>0.62 (0.53–0.76)</td>
</tr>
<tr>
<td>Baseline FeNO (ppb)</td>
<td>15.5 (9–26)</td>
<td>18 (13–24)</td>
<td>21 (8.5–42)</td>
</tr>
</tbody>
</table>

All figures are presented as the median (interquartile range). *$p<0.05$, control group vs severe group by Fisher exact test. †$p<0.05$ by Mann–Whitney U test. ‡$p<0.05$, both control and moderate group vs severe group by Mann–Whitney U test.

BDP, beclomethasone dipropionate; F, female; FeNO, fraction of exhaled nitric oxide; FEV1, forced expiratory volume in 1 s; FVC, forced vital capacity; ICS, inhaled corticosteroid; M, male; ppb, parts per billion.

<table>
<thead>
<tr>
<th>Parameter</th>
<th>Control</th>
<th>Moderate treated</th>
<th>Moderate untreated</th>
<th>Severe</th>
</tr>
</thead>
<tbody>
<tr>
<td>Sputum weight (g)</td>
<td>3.125 (1.72–5.14)</td>
<td>3.04 (1.53–3.54)</td>
<td>2.94 (0.77–2.69)</td>
<td>2.68 (1.54–6.32)</td>
</tr>
<tr>
<td>Cells/g of sputum (×10⁶/g)</td>
<td>0.76 (0.51–2.18)</td>
<td>0.57 (0.28–1.88)</td>
<td>1.58 (0.51–2.76)</td>
<td>1.13 (0.40–3.2)</td>
</tr>
<tr>
<td>Squamous cells (%)</td>
<td>16.28 (8.35–28.69)</td>
<td>19.8 (10.3–31.7)</td>
<td>12.0 (9.83–22.8)</td>
<td>13.8 (4.59–21.7)</td>
</tr>
<tr>
<td>Macrophages (%)</td>
<td>49.65 (23.02–70.11)</td>
<td>43.1 (35.5–58.5)</td>
<td>39.9 (29.7–66.5)</td>
<td>52.6 (28.1–73.6)</td>
</tr>
<tr>
<td>Neutrophils (%)*</td>
<td>35.42 (18.38–65.43)</td>
<td>51.8 (29.2–64.9)</td>
<td>45.4 (24.7–59.7)</td>
<td>18.2 (12.0–43.1)</td>
</tr>
<tr>
<td>Eosinophils (%)†</td>
<td>0.0 (0.0–0.36)</td>
<td>0.62 (0.28–5.58)</td>
<td>1.97 (0.93–11.8)</td>
<td>11.2 (1.25–45.5)</td>
</tr>
<tr>
<td>Lymphocytes (%)</td>
<td>1.31 (0.45–2.37)</td>
<td>1.26 (0.55–1.5)</td>
<td>1.11 (0.44–2.0)</td>
<td>1.3 (1.07–1.96)</td>
</tr>
<tr>
<td>Epithelial cells (%)</td>
<td>3.69 (1.46–5.35)</td>
<td>1.07 (0.77–2.69)</td>
<td>1.38 (0.69–2.15)</td>
<td>1.08 (0.61–1.68)</td>
</tr>
</tbody>
</table>

All data are presented as the median (interquartile range). *$p<0.01$, moderate treated group vs severe group. †$p<0.001$, control group vs moderate untreated group.
DISCUSSION

We have compared plasma exudation and coagulation activity in the sputum of healthy controls, patients with moderate persistent asthma, before and after withdrawal of ICS, and patients with severe persistent asthma. ICS withdrawal for 5 days resulted in significantly decreased FVII, increased plasminogen and tPA levels compared with pre-weaning values, and evidence of a more fibrinolytic environment in the airways of patients with untreated moderate asthma compared with controls. Conversely, patients with severe asthma had significantly higher levels of $\alpha_2$-macroglobulin, TF, PAI-1 and TAFI compared with those with treated moderate asthma, indicative of a change to a profibrinogenic and antifibrinolytic environment in severe asthma. We believe this is the first study to investigate the haemostatic balance in the airways of patients with severe asthma and following withdrawal of ICS in moderate asthma.

The significantly higher levels of $\alpha_2$-macroglobulin in the airways of patients with stable severe asthma are indicative of ongoing bronchial inflammation despite treatment with inhaled and/or oral steroids, and the strong correlation of $\alpha_2$-macroglobulin with sputum eosinophils supports this notion.
Discussion of the study design, and cohort characteristics, including sputum differential cell counts and FeNO measurements, is provided in the online appendix.

**Fibrinogenesis**

The higher levels of TF in the airways of patients with severe asthma, in the absence of detectable difference in TFPI levels, indicates a potentially procoagulant environment. Taking into account the plasma dilution factor, most of the TF we detected in the airways is likely to derive from local cellular sources, including monocytes, macrophages,18 eosinophils19 and possibly apoptotic bronchial epithelial cells20 that are features of severe asthma.

Levels of FVII were significantly lower in both untreated and severe asthma. This raises the possibility that FVII may be reduced by a mechanism associated with asthma pathogenesis, rather than therapy. Our unpublished in vitro evidence indicates that FVII and FX rapidly bind to bronchial epithelial cells, supporting the notion that these factors are lower in severe asthma because coagulation is initiated on cell surfaces by TF via reactions dependent on catalytic phospholipids in the cell membranes, and our samples have examined supernatant, not cell-bound FVII or FX. A strong negative correlation of these factors with observed TF levels in this study (FVII, \( r = -0.610, p = 0.002 \); FX, \( r = -0.634, p = 0.001 \)) lends further support for this hypothesis. Therefore, our observation may reflect binding and consumption of FVII and FX in the coagulation process, rather than their absence, which would completely abolish coagulation. Whilst not measured in this current study, levels of FXa activity have been shown to correlate with mucosal thickness and collagen deposition in a mouse model of asthma, and blockade with FXa inhibitor reduced airway hyper-responsiveness.21

Thrombin activity is normally undetectable in plasma, but was detected in all sputum samples (fig 2D), indicating local activation of prothrombin in the airways. Considering the plasma dilution factor, thrombin in the healthy airway is likely to be largely (94%) derived from locally expressed prothrombin, activated by FXa. However, prothrombin expression by bronchial epithelial cells remains to be confirmed by reverse transcription–PCR.

Levels of fibrinogen in our controls and subjects with asthma were similar to those reported elsewhere,22 23 although this is the...
first study to report levels in stable severe asthma and a positive correlation of fibrinogen with D-dimers (r = 0.417, p = 0.050) was observed. The levels in the control group may equate to derivation by plasma exudation, although a local source is not ruled out.4 However, values in the asthma groups were lower than those that would be derived from unfiltered plasma, indicating that fibrinogen is either deposited in the tissue matrix before it arrives at the epithelium22 or rapidly cleaved by thrombin and deposited as insoluble fibrin on epithelial cell surfaces.

FXIII levels in the severe group (0.46 (0.0–3.1) ng/ml) were lower than predicted (~36 ng/ml) by the plasma dilution factor. This indicates that FXIII may be adsorbed by the fibrin clot that forms as FXIII cross-links fibrin monomers, and/or that FXIII is proteolytically degraded, for example by neutrophil elastase which is shown to regulate FXIII within fibrin clots.23

**Fibrinolysis**

tPA is expressed by a number of cells in the airway including mast cells, macrophages, fibroblasts26 and bronchial epithelial cells.26 We have demonstrated higher levels of tPA in the severe and untreated asthma groups, which may indicate a disease-related increase in tPA expression. D-dimers were detected in all samples, indicating fibrin formation and fibrinolysis even in normal healthy airways, as previously reported.27 Despite significantly higher levels of tPA in untreated moderate and severe asthma, there were no significant differences in D-dimer levels between the groups which may have been degraded by local neutrophil elastase activity.28 Fibrinolysis is negatively regulated by PAI-1 and TAFI, both of which were significantly higher in the severe group as compared with controls, and calculation of a PAI-1/tPA molar ratio demonstrated a raised ratio in the severe cohort (control group median 0.873, severe 1.455) which suggests an antifibrinolytic environment in the airways, related to asthma severity. The higher FDP levels in severe subjects compared with controls is likely to represent overall upregulation of the external coagulation cascade with fibrin accumulation.

Levels of plasminogen detected in sputum are lower than those predicted from plasma dilution, indicative of binding of plasminogen to fibrin and cell surfaces, which accelerates its conversion to plasmin by tPA. Plasminogen expression by cells outside of the liver, including keratinocytes (reviewed by Irigoyen et al29), has been reported, but to our knowledge there are currently no reports of plasminogen expression by cells in the airway. Concentrations of tPA and PAI-1 in induced sputum are orders of magnitude higher than predicted from plasma exudation alone, and most must come from local cellular sources. TAFI was detected at levels that could be expected from plasma exudation, and TAFI expression by cells other than hepatocytes and adipocytes has not been described.30

Neutrophils,31 mast cells,32 macrophages, fibroblasts and the bronchial epithelium33 are potential local sources of PAI-1. A 4G/5G polymorphism in the PAI-1 promoter region has been reported, but to our knowledge there are currently no reports of plasminogen expression by cells in the airway. Concentrations of tPA and PAI-1 in induced sputum are orders of magnitude higher than predicted from plasma exudation alone, and most must come from local cellular sources. TAFI was detected at levels that could be expected from plasma exudation, and TAFI expression by cells other than hepatocytes and adipocytes has not been described.30

The role of corticosteroids

Plasminogen levels were significantly lower in those with treated moderate asthma compared with controls and rose significantly on cessation of ICS, with no change in plasma exudation, indicating that ICS may suppress local plasminogen expression. A similar suppressive effect of ICS on local tPA expression, possibly by bronchial epithelial cells,24 was demonstrated in the same group. Lower plasminogen and tPA levels would tend to reduce fibrinolysis and support fibrin-dependent epithelial repair in the airway, revealing a possible mechanism by which ICS contribute to epithelial continuity in moderate asthma.

PAI-1 and TAFI expression is enhanced by corticosteroids,30 35 and when oral and inhaled corticosteroid doses are converted to equivalent units (see online appendix) a significant correlation of PAI-1 with corticosteroid dose (r = 0.46, p = 0.04) was found for the moderate and severe groups. This raises the possibility that corticosteroids contribute to the increased expression of PAI-1 and TAFI in severe asthma and excessive fibrin formation that may be contributing to airway narrowing and BHR.3, 35

**Conclusions**

Defective fibrin formation and/or instability, and increased degradation of the fibrin matrix may contribute to the pathogenesis of moderate asthma, an effect that can be reversed by treatment with ICS.

Conversely, the levels of coagulation factors in severe asthma indicate a procoagulant, antifibrinolytic environment in which excessive fibrin formation and accumulation may contribute to irreversible airway narrowing and subepithelial bronchial fibrosis. Since PAI-1 and TAFI control the clearance of fibrin in the lung, and a stimulatory effect of PAI-1 on neutrophilic inflammation has been reported,31 36 our findings suggest that these molecules may be possible therapeutic targets in severe asthma.

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**Competing interests:** None.

**Ethics approval:** A favourable local research ethics committee opinion was obtained for the study.

**Provenance and peer review:** Not commissioned; externally peer reviewed.

**REFERENCES**

Treatment with low-dose lansoprazole may reduce frequency of COPD exacerbations in older patients

Colds caused by respiratory viruses can trigger exacerbations of chronic obstructive pulmonary disease (COPD). This Japanese study investigated whether lansoprazole reduces the frequency of COPD exacerbations by mitigating the effects of respiratory viruses.

One hundred and three patients with COPD (mean age 75 years, 95% male), but without evidence of gastro-oesophageal disease, were randomised to receive lansoprazole 15 mg once daily or no treatment on top of their usual COPD medications. Patients were followed for 1 year during which time they recorded a daily score for symptoms of a common cold and attended hospital fortnightly. The authors used a healthcare utilisation definition for an exacerbation of COPD.

Direct comparison of the two groups using standard parametric tests revealed no difference in the frequency of common colds between the two groups, although multivariate analysis suggested that lansoprazole use may be associated with a lower risk of frequent colds (>5/year).

There was a significant reduction in COPD exacerbations in the lansoprazole group (mean 0.34 exacerbations per year in the lansoprazole group vs 1.18 in the control group).

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