Expression and activation of TGF-β isoforms in acute allergen-induced remodelling in asthma

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ABSTRACT

Background
Airway wall remodelling and inflammation are features of chronic asthma. Transforming growth factor β (TGF-β) has been implicated in these processes.

Objectives
We determined the effect of allergen challenge on airway inflammation and remodelling and whether TGF-β isoforms and the Smad signalling pathways were involved.

Methods
Thirteen atopic asthmatics underwent inhalational challenge with 0.9% saline (SC), followed by allergen (AC) 3-4 weeks later. After both challenges, fiberoptic bronchoscopy was undertaken in order to obtain bronchial biopsies and tissue samples were processed for immunohistochemistry and examined by microscopy.

Results
FEV₁ fell after AC (-28.1 % ± 0.92, mean ± SEM, at 30 minutes) with a late response at 7 hours (-23.0% ± 1.23). AC caused an increase in neutrophils (p=0.016) and eosinophils (p=0.01) in the bronchial mucosa when compared with SC. Sub-basement membrane (SBM) thickness did not change after AC, but tenascin deposition in SBM was increased (p=0.02). Intranuclear (activated) Smad 2/3 and Smad 4 detected by immunohistochemistry were increased after AC in epithelial and subepithelial cells of bronchial biopsies. No inhibitory Smad (Smad 7) protein was detected. TGF-β isoforms 1, 2 and 3 were expressed predominantly in bronchial epithelium both after saline or allergen, but only TGF-β2 expression was increased after AC (p=0.03). Using double-immunostaining, an increase in TGF-β2-positive eosinophils (p=0.01) and neutrophils (p=0.04), but not in TGF-β1 positive eosinophils and neutrophils, was also found after AC.

Conclusions
TGF-β2 may contribute to the remodelling changes in allergic asthma following single allergen exposure; further detailed studies will be needed.

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Abbreviations:
SBM: subbasement membrane
TGF-β: transforming growth factor-beta
ECM: extracellular matrix
FEV₁: forced expiratory volume in one second
MBP: major basic protein
AC: allergen challenge
SC: saline challenge
INTRODUCTION

Airway wall remodelling in asthma comprises of structural changes within the airways, including an increase in subbasement membrane thickening, collagen deposition, smooth muscle hypertrophy/hyperplasia, angiogenesis and mucous gland hypertrophy. These features have been correlated with an increased loss of lung function observed in patients with asthma and with the development of progressive fixed airflow obstruction in some patients with asthma.

Although airway wall remodelling is now considered a characteristic feature of chronic asthma in adults, there is also evidence that remodelling can start at an early stage of the disease in children with asthma and may occur rapidly after an acute allergic response. An increase in the number of activated submucosal myofibroblasts has been reported within 24 hours after allergen challenge of patients with mild asthma. An increase in the number of submucosal fibroblasts characterized by collagen synthesis and in tenascin deposition within the subbasement membrane may also occur in response to acute allergen challenge in patients with atopic asthma. These studies indicate that airway wall remodelling is a dynamic process that could result from repeated incremental activation of the epithelial-mesenchymal-trophic unit as a result of repeated allergen exposures.

The mechanisms by which these rapid initial events in airway wall remodelling and inflammation are regulated have not been elucidated. Transforming growth factor β (TGF-β) is an important fibrogenic and immunomodulatory factor that can function either as a pro- or an anti-inflammatory cytokine, and therefore TGF-β may be involved in airway wall remodelling of asthma. TGF-β protein is expressed by inflammatory cells such as neutrophils and eosinophils, and also by structural cells such as epithelial cells, fibroblasts, and smooth muscle cells. TGF-β blockade with an anti-pan TGF-β antibody decreased the development of airway remodelling in a murine model of repeated allergen challenge.

TGF-β signaling pathway involves the phosphorylation of downstream Smad proteins, comprising of the receptor-regulated Smad (R-Smad, such as Smad 2, 3), the Co-mediator Smad (Co-Smad or Smad 4), and the inhibitory Smad (I-Smad, such as Smad 7). Activated Smad complexes translocate to the nucleus to upregulate the transcription of many target genes. Upregulation of R-Smad and Co-Smad has been found in the lungs of sensitized ovalbumin-exposed mice, indicating that the Smad-mediated signalling may be important in the pathophysiology of allergic pulmonary diseases. TGF-β can upregulate the production of extra-cellular matrix proteins, tenascin and fibronectin from lung fibroblasts, and the production of connective tissue growth factor (CTGF) from airway smooth muscle cells. Three different TGF-β isoforms (β1, β2, and β3) have been described although TGF-β1 has been the most extensively studied. High levels of TGF-β1 and β2 have been found in bronchoalveolar lavage fluid from patients with asthma following segmental allergen challenge. Of the TGF-β isoforms, only TGF-β2 was found to be increased in airway biopsies of patients with severe asthma compared to subjects with mild asthma; this increase was mainly localised to eosinophils.

In the present study, we examined the short term effect of an acute allergen challenge on the inflammatory and remodelling airway features in mild asthma. We have hypothesised that features of airway wall remodelling may be induced within 24 hours of an acute allergen exposure and that the expression and activation of TGF-β could be important in the pathogenesis of these changes.
METHODS
Patient population
Thirteen mild atopic non-smoking asthmatic patients (34.7 ± 2.4 years; 6 male; FEV₁: 92 ± 2.8% of predicted) were recruited from our outpatient clinic or by advertisement. All subjects had a history of mild allergic asthma together with documented bronchial hyperresponsiveness (metacholine PC₂₀ < 8mg/ml). None of the subjects had used inhaled or systemic corticosteroids. The protocol was approved by the Local Ethics Committee of the Royal Brompton & Harefield NHS Trust and National Heart & Lung Institute, London, UK. Informed consent was obtained from all subjects.

Skin prick testing
Atopy was defined by positive skin reaction (wheal diameter of over 3 mm above the negative control response) to extracts of Dermatophagoides pteronyssinus, cat fur and mixed grass pollen (Aquagen® from ALK, Reading, UK). The extract producing the largest wheal was used for allergen challenge.

Saline and allergen challenges
Patients stopped the use of short-acting β₂ agonists for at least 8 hours prior to the challenges. A challenge with 0.9% NaCl (saline) was first performed, followed one month later by a challenge with allergen between 0800 and 1000 am. Challenges were performed by inhalation of 5 breaths of solution from a breath-activated dosimeter (Model MB3, Mefar, Bovezzo, Italy). Each solution was administered from a hand-held nebulizer attached to the breath-activated dosimeter with a delivery time of 1 s per breath (aerodynamic mass median diameter of 3.5 µm; output of 9µl/breath). FEV₁ was recorded using a dry wedge spirometer (Vitalograph, Buckingham, UK) and the highest value was taken as the post-saline value. One month later, an allergen challenge (AC) was performed. Fresh dilutions of freeze-dried allergen extract (Aquagen SQ; AlK, Reading, UK) were made up with 0.9% saline from a stock solution of 100000 U/ml. After baseline and post-saline FEV1 measurements, starting with an allergen concentration of 250 U/ml, 5 breaths of serially increasing concentrations of allergen were inhaled from a hand-held nebuliser. The challenge was terminated when >15% fall in FEV₁ from the post-saline value was observed. After AC, FEV₁ was measured also at 20, 30, 45, and 60 min and thereafter in duplicate at 30 min intervals up to 10 h. A late asthmatic response was defined as a fall in FEV₁ of greater that 15% from the post-saline value on at least three occasions, between 4 and 10 h.

Bronchial biopsies
The biopsies were embedded in OCT (optimal cutting temperature) medium and then snap-frozen in precooled isopentane in liquid nitrogen and stored at -70°C. Cryostat sections (6 µm) were obtained from biopsies, placed on poly-L-lysine-coated microscope slides and stained with haematoxylin and eosin (for measurement of subbasement membrane thickness).

Staining of eosinophils and neutrophils: Fixed sections in acetone were washed immunostained for eosinophils and neutrophils in bronchial mucosa using
immunoperoxidase 3% in methanol for 30 minutes (Vectstain kit PK4002, Vector, Peterborough, UK) and incubated with mouse monoclonal antibodies to major basic protein (MBP; dilution 1:100; Cat No MON6008-1, Monosan, The Netherlands) or to neutrophil elastase (NE; dilution 1:50; DAKO Cytomation, Cat No M0756, DAKO, Denmark), for 1 hour at room temperature. Sections were washed in PBS and incubated 45 minutes with biotinylated mouse IgG secondary antibody, made up in PBS containing horse serum. 3,3'-diaminobenzidine (Sigma D5905, Sigma-Aldrich Company Ltd, Gillingham, UK) was used for brown staining. Sections were finally counterstained in haematoxylin (20% in tap water) and dehydrate through alcohols to xylene.

**Staining of TGF-β isoforms:** TGF-β isoforms were also detected following the same method as above, in tissue sections, using TGF-β1, β2 and β3 specific (Santa Cruz Biotechnology, Santa Cruz, CA, USA).

To determine specific cells types expressing TGF-β, double immunostaining labelling techniques with immunoperoxidase (for cell markers) and rhodamine-conjugated (for TGF-β isoforms) were applied. Briefly, endogenous peroxidase activity was blocked using 3% H2O2. Sections were washed in PBS and then incubated with primary anti-neutrophil elastase antibody or anti-MBP antibody in PBS containing 0.1% bovine albumin (BSA). IgG secondary antibody was then applied for 45 minutes. 3,3'-diaminobenzidine (Sigma D5905, Sigma-Aldrich Company Ltd, Gillingham, UK) was used for brown staining. Sections were then incubated overnight at 4°C with TGF-β1 or TGF-β2 polyclonal antibody diluted in PBS/saponin buffer. Sections were washed thoroughly in PBS/saponin buffer before incubation with Rhodamine Red-X-conjugated donkey anti-rabbit IgG (Jackson Immunoresearch, Stratech Scientific Ltd, Cambridgshire, UK) for 45 min at room temperature in dark. After washing with buffer nuclei were counterstained with DAPI (4',6 diamidino-2-phenylindole) (Vectashield, Vector Laboratory, Peterborough, UK). For negative control preparations, the primary antibody was replaced by non-specific mouse immunoglobulin. Neutrophils and eosinophils stained brown and TGF-β1 and β2 stained as fluorescent red. Pictures of sections from the same area were taken to co-localize both staining.

**Staining of ECM proteins (tenascin, procollagen-III) and Smad proteins (2-3, 4 and 7):** Tissue sections were pretreated with PBS and then incubated with antibody against tenascin (Mouse monoclonal, dilution 1:50, Monosan MON7025, Monosan, The Netherlands), procollagen-III (Rabbit polyclonal, dilution 1:50, Chemicon AB764P, Chemicon International, Hampshire, UK) overnight in the refrigerator; and Smad 2/3 (Rabbit polyclonal, dilution Upstate Cat No. 07-408, Upstate, UK), Smad 4 (Rabbit polyclonal, dilution 1:50, Upstate Cat No. 06-693, Upstate, UK ), and Smad 7 (Santa Cruz, Cat No 7004, dilution 1:50, Santa Cruz, CA, USA) for 1 hour at room temperature. After extensive washing, sections were incubated with Rhodamine and nuclei were counterstained with DAPI (4',6 diamidino-2-phenylindole) (Vectashield, Vector Laboratory, Peterborough, UK).

**Tissue quantification**

The subbasement membrane was identified under light microscopy and the thickness (µM) was calculated by dividing the area of subbasement membrane by the length using a computer analysis system (KS-300, Zeiss, Germany). Eosinophil and neutrophil counts were expressed as number of cells staining positively for major basic protein or neutrophil elastase respectively per 0.036mm². Measurements were performed down to a depth of 100µm from the luminal surface
in areas where the epithelium was present. The same method was used to determine cells expressing TGF-β1, TGF-β2 and TGF-β3 staining. Immunofluorescence pictures were visualized using a confocal microscope (Leica TCS SP, Heidelberg, Germany). All the slides had their images recorded using the same settings for brightness and contrast, which allowed for comparison and analysis. The presence of SBM proteins (tenascin and procollagen-III) was calculated using Scion Image Analysis software package (2000, Scion Corporation, Maryland, USA) by measuring the fluorescence intensity. The density (number of pixels) in the SBM area (µm²) was calculated and divided by the SBM length to obtain thickness (µm) per density (pixels) as a measure of expression of the matrix proteins in the subbasement membrane.

Smad 2/3, 4 and 7 positive cells were counted using confocal microscope. A cell was considered as positive when SMAD protein was expressed in the nucleus. Counts are expressed as % of intranuclear positively-stained cells of total cells in the biopsy area down to a depth of 100µm from the luminal surface where with epithelium intact.

Double immunostaining for MBP or NE and TGF-β1 or TGF-β2 was calculated by measuring the % of eosinophils or neutrophils staining positively for TGF-β1 or TGF-β2 in the biopsy area down to a depth of 100µm from the luminal surface where the epithelium was present.

These measurements were performed by one investigator (AT) who was not aware of the treatment received by the patient as the slides were coded by another investigator (TO).

**Statistical Analysis**
Data are presented as means ± SEM. Paired Student’s t-test was used to analyse changes in the numbers of positive cells, SBM thickness and density after saline and allergen challenge. A p value of < 0.05 was accepted as significant.
RESULTS

FEV\textsubscript{1} and mucosal eosinophils and neutrophils

All patients experienced a biphasic asthmatic response after allergen challenge. The fall in FEV\textsubscript{1} after allergen inhalation was maximal at 30 minutes (-28.1 \% \pm 0.92, p<0.001), followed by a partial recovery and with a late fall maximal at 7 hours (-23.02 \% \pm 1.23, p<0.01). Following saline challenge, there was no significant change over the ensuing 10 hours.

There was a significant increase in neutrophils and eosinophils when comparing biopsies after saline or allergen challenge: neutrophils (from 6.2 \pm 1.0 to 11.6 \pm 1.8 cells per 0.036mm\textsuperscript{2}; p=0.016) and eosinophils (from 2.7 \pm 0.7 to 8.6 \pm 1.4 cells per 0.036mm\textsuperscript{2}; p=0.01). The increase in neutrophils was significantly correlated with the increase in eosinophils (r=0.607, p=0.001).

SBM thickness and density

There was no difference in SBM thickness after the two challenges (8.1 \pm 0.4 vs. 8.33 \pm 0.4 \textmu m). The deposition of the ECM protein, tenascin, as measured by the thickness multiplied by density in the SBM, was increased after allergen challenge (thickness x density, p=0.02) (Figure 1). No significant changes in procollagen III deposition was found.

Smad protein and TGF-\beta isoform expression

Positive Smad 2/3 and Smad 4 cells were localised predominantly in the epithelium although there were also some positively-stained subepithelial cells (Figure 2). There were significantly higher number of cells expressing Smad 2/3 and Smad 4 in the nuclei of cells located predominantly in the epithelium after allergen challenge than after saline challenge (Smad 2/3: 4.38 \pm 1.5 \% positive cells after saline challenge, 13.00 \pm 3.6 \% positive cells after allergen challenge, p=0.03; Smad 4: 4.4 \pm 1.08 \% positive cells after saline challenge to 16.07 \pm 3.4 \% positive cells after allergen challenge, p=0.01).

No positively-stained Smad 7 cells were detected. In order to ensure that the Smad 7 staining antibody was able to detect Smad7, we stained airway smooth muscle cells transfected with an adenovirus vector expressing double-positive Smad7. We were able to demonstrate positive staining for Smad7 using our anti-Smad7 antibody (data not shown).

There was positive staining with the antibodies to the 3 TGF-\beta isoforms (1, 2 and 3). The predominant TGF-\beta staining was found in the epithelial cells, particularly in the apical pole. However, there was a significant difference in TGF-\beta2 positive cells after allergen challenge. The counts were as follows: TGF-\beta1 from 14.01 (\pm 1.46) \% positive cells after saline challenge to 15.06 (\pm 1.61) \% positive cells after allergen challenge (p>0.05); TGF-\beta2: from 11.3 (\pm 1.9) to 17.1 (\pm 1.5) \% positive cells (p=0.03); and TGF-\beta3: from 8.31 (\pm 1.53) to 8.69 (\pm 1.8) \% positive cells (p=0.76) (Figure 3).

Expression of TGF-\beta1 and TGF-\beta2 in eosinophils and neutrophils

Using double immunostaining TGF-\beta1 and TGF-\beta2 were found in both eosinophils and neutrophils present in the airways mucosa (Figure 4). The eosinophils and neutrophils positively stained for TGF-\beta1 did not change after the allergen challenge (7.4 (\pm 0.9) \% of TGF-\beta1 positive eosinophils after saline challenge to 8.1 (\pm 1.12) \% of TGF-\beta1 positive eosinophils after allergen challenge (p=0.21); and from 6.1 (\pm 0.76) \% of TGF-\beta1 positive neutrophils after saline challenge to 7.0 (\pm 0.93) \% of TGF-\beta1 positive neutrophils after allergen challenge (p=0.43). However, TGF-\beta2 positive eosinophils and neutrophils significantly increased after allergen challenge: 5.7 (\pm 0.7) \% of TGF-\beta2 positive eosinophils after saline challenge to 9.4 (\pm 1.1) \% of TGF-\beta2 positive eosinophils after allergen challenge (p=0.01).
challenge to 9.3 (±1.16) % of TGF-β2 positive eosinophils after saline challenge (p=0.016); and from 3.27 (±0.6) % of TGF-β2 positive neutrophils after saline challenge to 5.3 (±0.75) % of TGF-β2 positive neutrophils after allergen challenge (p=0.04) (Fig 5).

We found no significant correlations between the degree of fall in FEV₁ in the early or late phase response and the number of positively-staining Smad2/3 or Smad4 epithelial cells, or the number of positively-staining TGFβ1- or 2-staining eosinophils or neutrophils measured after allergen challenge.
Discussion

The novel finding in this study is the increase in expression of TGF-β2 expressing cells particularly in the airway epithelial cells, and also in eosinophils and neutrophils following acute allergen challenge, but without an increase in expression of TGF-β1 and TGF-β3 in the epithelium or infiltrating leucocytes. Further support for a role for TGF-β2 comes from the demonstration that there is an increased expression and activation of Smad 2/3 and Smad 4 expression following allergen challenge, expressed principally in the epithelium. Our data indicate that TGF-β2 may be responsible for the increased deposition of tenascin observed after single allergen challenge, although the contribution of TGF-β1 and TGF-β3 to the process may still be important.

Examination of bronchial biopsies from atopic asthma subjects undergoing allergen challenge revealed that within 24 hours after allergen exposure, there was no significant change in subbasement membrane thickness. However, tenascin deposition within the sub-basement membrane was increased, although procollagen-III deposition did not change significantly. This implies that changes in airway wall remodelling may occur within 24 hours of allergen challenge, confirming a previous study in which similar measurements were made.

The epithelial-mesenchymal trophic unit, comprising of the airway epithelium, the subbasement membrane, and adjacent fibroblasts, has been proposed to be responsible for the changes in airway wall remodelling observed in asthma. We demonstrate that there is an increased expression of TGF-β2 and activation of the TGF-β signalling pathway following allergen challenge. Activation of the TGF-β2 activating signalling pathway may occur in the airway epithelial cells, and we presume that this may be the cause of increased tenascin deposition observed in the extracellular matrix. Indeed, both TGF-β1 and TGF-β2 can increase the expression of tenascin and fibronectin in the extracellular matrix surrounding BEAS2B human bronchial epithelial cell line.

There is circumstantial evidence for a role for TGF-β in asthma, particularly under conditions of the experimental allergen challenge. For example, increased levels of TGF-β have been measured in broncho-alveolar lavage fluids of asthmatic subjects following allergen challenge. TGF-β pathway signalling involves the phosphorylation of downstream Smad proteins, which is comprised of the receptor-regulated Smad (R-Smad, such as Smad 2, 3), the co-mediator Smad (Co-Smad or Smad 4), and the inhibitory Smad (I-Smad, such as Smad 7). Activated Smad complexes translocate into the nucleus to upregulate the transcription of target genes. Rosendahl et al. have reported from examination of human airway biopsies and of murine allergen-challenged models that Smad signalling in bronchial epithelial cells could play a determinant role in the pathophysiology of allergic respiratory diseases. Our present data also support a previous observation that demonstrated upregulation of R-Smad (Smad 2) in asthmatic lungs induced by allergen.

In our study, TGF-β1 expression did not change after allergen challenge, but there was significant baseline expression of TGF-β1, TGF-β2 and TGF-β3 in biopsies obtained from mild atopic asthma patients. Because we did not compare this expression with that found in normal healthy subjects, we are unable to determine whether this represents an increase in mild asthma. However, other studies suggest that in mild asthma, TGF-β1 expression is increased. With regard to TGF-β2, this isoform has been demonstrated to be the most prominent in biopsies with severe asthma, and this was result of the increased expression in
infiltrating eosinophils. In addition, this increase in TGF-β2 expression was associated with an increase in subbasement membrane thickness in the patients with the most severe asthma.

One limitation of our study is that we studied only one time point (24 hours) and cannot therefore tell whether the deposition of tenascin observed at 24 hours is only the beginning of the process. Relevant to this is our inability to detect any evidence of the inhibitory Smad (Smad 7) activation, even though our antibody was able to detect Smad 7. However, we do not know whether Smad 7 activation had occurred earlier or would occur later, although it is likely that all the Smad family members would be expected to be activated simultaneously. Whether the lack of Smad 7 activation represents a defect in asthma, which could underlie an increased functional effect of TGF-β activation in asthmatics undergoing allergen challenge, is not known. Nakao and colleagues have reported that the baseline immunoreactive expression of Smad 7 in bronchial epithelial cells of asthmatic subjects was significantly less compared to that of non-asthmatic subjects, and that the expression correlated inversely with the degree of subbasement membrane thickness.

We found that there was both an increase in submucosal eosinophils and neutrophils following allergen challenge. Although much emphasis has been previously placed on the post-allergen eosinophilia, little attention has been given to the neutrophilic inflammation which was as prominent as the eosinophilia. We also found that both eosinophils and neutrophils were capable of expressing TGF-β1 and TGF-β2, with negligible expression of TGF-β3. In addition, in both neutrophils and eosinophils, there was an increase in the percentage of these cells expressing TGF-β2 following allergen challenge, as determined by the double-immunostaining techniques we used. The capacity for both eosinophils and neutrophils to express TGF-β isoforms, particularly after allergen challenge, supports the possibility that inhibition of both eosinophilic and neutrophilic inflammation may lead to prevention of airway wall remodelling. Neutrophilic inflammation is a particularly important feature of severe asthma. Treatment with anti-IL5 antibody, that reduced the number of submucosal eosinophils in patients with asthma, was associated with a reduction in TGF-β levels in bronchoalveolar lavage fluid, indicating that TGF-β may be induced by IL-5. In-vitro studies of epithelial cells also indicate that neutrophil elastase could induce TGF-β secretion from the extracellular matrix, further emphasising the potential contribution of neutrophils to airway wall remodelling in asthma. Human mast cells chymase also possesses similar properties, and could be another stimulus for TGF-β release following allergen challenge. Finally, in a murine asthma model, therapeutic treatment with a blocking anti-TGF-β antibody reversed peribronchiolar extracellular matrix deposition.

In conclusion, we have shown that allergen challenge of patients with mild asthma induces the deposition of tenascin in the subbasement membrane within 24 hours associated with neutrophil and eosinophil inflammation, activation of the TGF-β associated Smad pathway, and upregulation of TGF-β2 expression in epithelial cells, neutrophils and eosinophils. These changes in tenascin deposition and TGFβ2 expression are small; however, repeated episodes of allergic inflammation induced by continuous or repeated allergen exposures could lead to significant cumulative increases. Our data also implicate a potential importance for TGFβ2 expression and activation in the increase in tenascin deposition. However, further studies are needed to confirm this.
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Competing interests
The authors declare no competing interests.

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References


Legend to figures

Figure 1
Panel A: Individual changes in SBM thickness (µm) and in expression of tenasin and pro-collagen III in SBM measured as thickness (µm) x density (pixels). Dotted lines are mean ± SEM.
Panel B: Representative confocal microscopy pictures showing increase of tenasin immunoreactiviy (red) in SBM from two subjects (A, B) after saline (A1, B1) and allergen challenge (A2, B2). Sections were counterstained with DAPI (blue) to demonstrate nuclear staining. Original magnification: x 400

Figure 2
Panel A: Changes in % of total cells from biopsy sections with nuclear positive staining for Smad 2/3 and Smad 4 proteins after saline and allergen challenges. Dotted lines represent mean ± SEM.
Panel B: Representative immunofluorescence confocal microscopy showing an increase in Smad 2-3 (A) and Smad 4 (B) proteins (red staining) in bronchial tissue after saline (1) and allergen (2) challenge in the nucleus (arrows).

Figure 3
Expression of TGF-β1, TGF-β2 and TGF-β3 in bronchial mucosa.
Panel A: Changes in the % of total cells with positive staining for TGF-β1, TGF-β2 and TGF-β3 after saline and allergen challenge. Dotted lines represent mean ± SEM.
Panel B: Representative pictures of bronchial tissue taken with light microscopy immunostained for TGF-β1 (a), TGF-β2 (b) and TGF-β3 (c). Original magnification: x400.

Figure 4
Localization of TGF-β2 staining in MBP and neutrophil elastase positive cells in airway mucosa
Top panel: Section under confocal microscopy from atopic asthma subject with TGF-β2 positive staining shown as red fluorescence.
Bottom panel: Similar sections demonstrating either major basic protein positive (MBP) staining (eosinophils) or neutrophil elastase (NE) staining (neutrophils) as brown staining. Arrows show the same cells (either eosinophil or neutrophil with TGF-β2 positive staining). Original magnification: x400.

Figure 5
Expression of TGF-β2 in eosinophils and neutrophils. Change in % of eosinophils and neutrophils from biopsy sections positively stained for TGF-β2. Cells were counted in the area down to a depth of 100µm from the luminal surface.
Fig 1A
Fig 2A
Fig 5