



# Effects of cigarette smoke extract on human airway smooth muscle cells in COPD

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**ABSTRACT** We hypothesised that the response to cigarette smoke in airway smooth muscle (ASM) cells from smokers with chronic obstructive pulmonary disease (COPD) would be intrinsically different from smokers without COPD, producing greater pro-inflammatory mediators and factors relating to airway remodelling.

ASM cells were obtained from smokers with or without COPD, and then stimulated with cigarette smoke extract (CSE) or transforming growth factor- $\beta$ 1. The production of chemokines and matrix metalloproteinases (MMPs) were measured by ELISA, and the deposition of collagens by extracellular matrix ELISA. The effects of CSE on cell attachment and wound healing were measured by toluidine blue attachment and cell tracker green wound healing assays.

CSE increased the release of CXCL8 and CXCL1 from human ASM cells, and cells from smokers with COPD produced more CSE-induced CXCL1. The production of MMP-1, -3 and -10, and the deposition of collagen VIII alpha 1 (COL8A1) were increased by CSE, especially in the COPD group which had higher production of MMP-1 and deposition of COL8A1. CSE decreased ASM cell attachment and wound healing in the COPD group only.

ASM cells from smokers with COPD were more sensitive to CSE stimulation, which may explain, in part, why some smokers develop COPD.



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## Introduction

Chronic obstructive pulmonary disease (COPD) is a leading cause of morbidity and mortality worldwide and results in an economic and social burden that is both substantial and increasing [1, 2]. In COPD a number of changes occur in the lungs, namely the development of persistent inflammation and irreversible airflow limitation [3].

Airflow limitation is caused by three interrelated processes: thickening (remodelling) of the small airway walls, loss of small airways, and emphysema. However, small airway remodelling is considered to have the greatest influence on airflow limitation [4–6]. Furthermore, it is likely that the small airway remodelling is the primary pathological insult in COPD. In a study using micro-computed tomography analysis of COPD lung tissue, it was found that remodelling and loss of terminal bronchioles preceded emphysematous changes microscopically [7]. The small airway remodelling in COPD consists of folded mucosa, thickening of basement membrane and deposition of connective tissue, as well as increased airway smooth muscle (ASM) mass, especially in severe COPD [6, 8, 9]. The connective tissue consists of an intertwined framework of extracellular matrix (ECM) proteins, and the specific ECM proteins are known to be altered in the airways of patients with COPD [10, 11].

In the developed world, the main risk factor for the development of COPD is cigarette smoking. Through the use of both *in vivo* and *in vitro* models the effects of smoking on the aetiology of COPD is beginning to be understood. Most studies address the paradigm that the aetiology of COPD is cigarette smoke-induced inflammation leading to tissue damage; however, previous research suggested that airway remodelling may be induced independently of inflammation [12]. It has previously been found that fibroblasts from patients with COPD produced an excessive amount of fibronectin and perlecan or reduced proteoglycans (decorin and biglycan) in response to cigarette smoke extract (CSE), in comparison to cells from people without COPD [12, 13]. Epithelial cells in COPD also respond differently to CSE [14], but whether COPD ASM cells respond differently to CSE is not known.

In this study we hypothesised that the response to cigarette smoke in ASM cells from people with COPD would be intrinsically different to that in ASM cells from people without COPD, specifically in the production of pro-inflammatory mediators and factors relating to airway remodelling.

## Material and methods

### Study subjects

Subject information was obtained regarding diagnosis, smoking history and lung function. Subjects with a diagnosis of asthma, infectious diseases or interstitial lung disease were not included. Samples were obtained from subjects who were classified as follows according to severity of airflow limitation [15]. 1) Non-COPD:  $n=21$ , forced expiratory volume in 1 s (FEV<sub>1</sub>)/forced vital capacity (FVC)  $\geq 70\%$  and FEV<sub>1</sub>  $\geq 80\%$ . 2) COPD:  $n=20$ , FEV<sub>1</sub>/FVC  $< 70\%$ . Full details are provided in the supplementary material (table S1). All study subjects or their next of kin provided written informed consent. Approval of all the experiments using human lung tissues was provided by the Ethics Review Committee of the South West Sydney Area Health Service, St Vincent's Hospital Sydney, and the University of Sydney Human Research Ethics Committee (all Sydney, Australia).

### Cell culture and sample preparation

Human ASM cells were obtained from human lung by a method modified from one described previously [16]. Human ASM cells were microdissected from approximately sixth-order or greater bronchi, and were initially cultured in growth medium comprised of DMEM (Invitrogen, Carlsbad, CA, USA) supplemented with 5% fetal bovine serum (DKSH, Melbourne, Australia), 1% antibiotics (Invitrogen) and 25 mM HEPES (Invitrogen). All the cells tested negative for the presence of mycoplasma before they were set up for experiments, and were used between passages 2 and 7. ASM cells were seeded in six-well or 96-well culture plates (BD Biosciences, North Ryde, Australia) at a density of  $1 \times 10^4$  cells·cm<sup>-2</sup> in growth medium and incubated at 37°C/5% CO<sub>2</sub> for 72 h. Cells were starved in quiescing medium consisting of DMEM supplemented with 0.1% fetal bovine serum, 1% antibiotics and 25 mM HEPES for 24 h before stimulation with CSE or 10 ng·mL<sup>-1</sup> of transforming growth factor (TGF)- $\beta$ 1 (R&D Systems, Minneapolis, MN, USA) in quiescing medium. After stimulation, supernatants from human ASM cells (in six-well plates) were collected, and cells (in 96-well plates) were lysed with 0.016 mM ammonium hydroxide (NH<sub>4</sub>OH) at 37°C for 20 min then washed with 0.05% PBS-Tween (Tw)20 (vol/vol). These supernatants and cell-free ECM plates were stored at -20°C until analysis.

### Cigarette smoke extract

We used Marlboro Red cigarettes (Philip Morris, Victoria, Australia), and each cigarette contained 1.1 mg of nicotine, 15 mg of tar, and 15 mg of carbon monoxide. CSE was prepared by a method modified from

one described previously [12]. Briefly, one Marlboro cigarette was bubbled through 25 mL of DMEM at a constant rate and this solution was regarded as 100% concentration CSE. The 100% CSE was freshly generated for each experiment, and diluted to final working concentration and used within 30 min.

#### **Cell viability and toxicity assays**

Human ASM cells were seeded in 96-well plates as described previously, and cells were stimulated with serial dilutions of CSE from 0.05% to 50%. The mitochondrial activity of living cells was tested by Thiazolyl blue tetrazolium bromide (MTT) (Sigma Aldrich, St Louis, MO, USA) assay. The membrane integrity of cells was tested by a lactate dehydrogenase (LDH) (Sigma Aldrich) assay, based on the amount of cytoplasmic LDH released into the medium. After stimulation for 72 h, MTT and LDH release were measured using a spectrophotometer (Spectramax M2; Molecular Devices, Sunnyvale, CA, USA) setting with absorbance 570 nm/690 nm and 490 nm/690 nm, respectively.

#### **Chemokine ELISA**

Human ASM cells were seeded in six-well plates as described previously, and cells were stimulated with different concentration of CSE or 10 ng·mL<sup>-1</sup> of TGF-β1 for 72 h. The concentrations of CXCL8 (interleukin (IL)-8), CXCL1 (GROα), CCL2, CCL5 and CXCL10 in the supernatants from human ASM cells were measured by using commercial human CXCL8/IL-8, CXCL1/GROα, CCL2/monocyte chemoattractant protein-1, CCL5/RANTES, and CXCL10/inducible protein-10 ELISA kits (R&D Systems) according to the manufacturer's instructions. The absorbance was read at 450 nm/570 nm using a spectrophotometer (Spectramax M2).

#### **Transcription factor nuclear factor-κB and activator protein-1 activity assay**

Human ASM cells were seeded in six-well plates as described previously, and cells were stimulated with CSE (5% and 10%) or 10 ng·mL<sup>-1</sup> of TGF-β1 for 60 min and then nuclear extracts were collected. The activities of nuclear factor (NF)-κB and activator protein (AP)-1 of each sample were assessed using the TransAM ELISA kits (Active Motif, Carlsbad, CA, USA) according to the manufacturer's instructions. The absorbance was read at 450 nm/655 nm using a spectrophotometer (Spectramax M2).

#### **Real-time PCR array**

Human ASM cells obtained from smokers with (n=3) and without COPD (n=3) were stimulated *in vitro* with 5% CSE, 10% CSE or 10 ng·mL<sup>-1</sup> of TGF-β1. RNA was collected at 48 h then purified using the Isolate RNA mini kit (Bioline, London, UK), and mRNA was converted to cDNA using M-MLV reverse transcriptase (Invitrogen). Equal amounts of cDNA of each sample from the same group were pooled; the gene expression was then tested using TaqMan array human extracellular matrix and adhesion molecules 96-well plates according to the manufacturer's instructions (Invitrogen). Real-time PCR was performed using the StepOne Plus detection system and data were collected and analysed by StepOne software (Applied Biosystems, Melbourne, Australia). The relative abundance of mRNA was calculated using the ΔΔCt method [17], and results were normalised to 18S rRNA.

#### **Matrix metalloproteinase ELISA**

Human ASM cells were seeded in six-well plates as described previously, and cells were stimulated with different concentration of CSE or 10 ng·mL<sup>-1</sup> of TGF-β1 for 72 h. The concentrations of total matrix metalloproteinase (MMP)-1, -2, -3, -10 and -12 were measured using human MMP ELISA kits (R&D Systems) according to the manufacturer's instructions, and the reading was performed using a Luminex analyser (Luminex 200 System; Luminex, Brisbane, Australia). MMP-1 enzyme activity was measured using human active MMP-1 fluorescent assay kit (R&D Systems), and the relative fluorescence units were determined using a fluorescence plate reader (Spectramax M2) setting with excitation wavelength 320 nm and emission wavelength 405 nm.

#### **ECM ELISA**

Human ASM cells were seeded in 96-well plates as described previously, and cells were stimulated with different concentrations of CSE or 10 ng·mL<sup>-1</sup> of TGF-β1 for 72 h. Cell-free ECM plates were used to measure the deposition of protein in the ECM by ELISA according to the previously modified method [16]. Primary antibodies used for detecting ECM proteins were rabbit polyclonal anti-human COL5 antibody (2 μg·mL<sup>-1</sup>) (Abcam, Cambridge, UK), rabbit polyclonal anti-human collagen VIII alpha 1 (COL8A1) antibody (2 μg·mL<sup>-1</sup>) (Novus Biologicals, Littleton, CO, USA), mouse monoclonal anti-human fibronectin antibody (2 μg·mL<sup>-1</sup>) (Millipore, Billerica, MA, USA), and mouse monoclonal anti-human perlecan antibody (2 μg·mL<sup>-1</sup>) (Invitrogen). Rabbit IgG (Dako Cytomation, Glostrup, CA, USA) isotype control antibody and mouse IgG1 κ isotype control antibody (BD Biosciences) were used at the same concentration as the primary

antibodies. For measurement of COL8A1 and perlecan, the biotinylated goat anti-rabbit antibody  $0.5 \mu\text{g}\cdot\text{mL}^{-1}$  and biotinylated chicken anti-mouse antibody  $0.8 \mu\text{g}\cdot\text{mL}^{-1}$  were used, respectively.

#### **Western blots**

Human ASM cells were seeded in six-well plates as described previously, and cells were stimulated with CSE (5% or 10%) or TGF- $\beta$ 1 for 72 h. The supernatants from each sample were collected to assess the soluble COL1 and soluble fibronectin using Western blots. Proteins were size fractionated on 10% polyacrylamide gels, transferred to polyvinylidene fluoride membranes and blocked in 5% (wt/vol) skim milk solution for 1 h. The membranes were incubated with primary antibody ( $8.8 \mu\text{g}\cdot\text{mL}^{-1}$  of mouse monoclonal anti-COL1 antibody (Sigma Aldrich) or  $1 \mu\text{g}\cdot\text{mL}^{-1}$  of mouse monoclonal anti-human fibronectin antibody (Millipore) in 2% bovine serum albumin/TBS-Tw) for 2 h, followed by incubation with secondary antibody ( $2.6 \mu\text{g}\cdot\text{mL}^{-1}$  of rabbit anti-mouse Ig-horseradish peroxidase antibody (Dako Cytomation) in 2% bovine serum albumin/TBS-Tw) for 1 h. Immunoblot detection was performed using Immobilon Western Chemluminescent HRP Substrate (Millipore) and bands were analysed using Kodak image station 4000 MM (Eastman Kodak Co., New York, NY, USA). The amount of protein present in each sample was determined as the densitometric density.

#### **Immunohistochemistry**

The preparation of immunohistochemical samples and immunohistochemical methods have been described previously [18]. The airway sections were treated to minimise nonspecific background staining, and incubated with primary rabbit polyclonal anti-human COL8A1 antibody  $1 \mu\text{g}\cdot\text{mL}^{-1}$  (Abcam) and rabbit IgG isotype control antibody  $1 \mu\text{g}\cdot\text{mL}^{-1}$  (Dako Cytomation). The conjugated secondary antibody was labelled polymer anti-rabbit (EnVision+System-HRP; Dako Cytomation) and the tissue staining was visualised with substrate chromogen, liquid DAB (Dako Cytomation). 10 images of each section (one section per subject) were taken and immunostaining was quantified using Fiji software (ImageJ) [19]. Full details are provided in the supplementary material.

#### **Cell attachment assay**

96-well culture plates were exposed to growth medium for 72 h and to quiescing medium for 24 h then exposed to quiescing medium with different concentration of CSE (0.05% to 10%) for 72 h. Human ASM cells were seeded on these treated plates at a density of  $5 \times 10^4 \text{ cells}\cdot\text{cm}^{-2}$  in quiescing medium for 2 h. Cell attachment was detected by a toluidine blue attachment assay as previously described [20]. The relative number of attached cells was measured using spectrophotometry at an absorbance of 595 nm (Spectramax M2). Full details are provided in the supplementary material.

#### **Wound healing assay**

An Oris cell migration assembly kit (Platypus Technologies, Madison, WI, USA) was used to perform the wound healing assay. The wound was created on the treated 96-well black plate and cells were labelled with cell tracker green CMFDA (Invitrogen). The labelled human ASM cells were seeded on the wounded black plate at a density of  $5 \times 10^4 \text{ cells}\cdot\text{cm}^{-2}$  in growth medium. After adhesion for 24 h the stoppers were removed, and incubation was continued for 4 h. The wound healing value was measured using a fluorescence plate reader (Wallac VICTOR<sup>2</sup>; Perkin Elmer, Waltham, MA, USA) read from the bottom with excitation wavelength set at 485 nm and emission wavelength at 535 nm. Full details are provided in the supplementary material.

#### **Statistical analysis**

Data analysis was performed using GraphPad Prism 5.0 software (GraphPad Software, San Diego, CA, USA). All the data are presented as mean  $\pm$  SEM. One-way ANOVA, two-way ANOVA plus Bonferroni post-test or Mann–Whitney test were used as appropriate to determine the statistical significance. A p-value  $\leq 0.05$  was considered significant.

## **Results**

### **Cytotoxicity of CSE on ASM cells**

As high concentrations of CSE are known to be cytotoxic we used two toxicology assays, assessment of mitochondrial activity *via* MTT and membrane integrity *via* LDH release, in order to ensure that the concentrations of CSE we used had no toxic effects on human ASM cells. CSE at a concentration  $\geq 15\%$  substantially reduced viability (fig. 1); therefore, we used 0.05% to 10% CSE as stimulation in subsequent experiments.

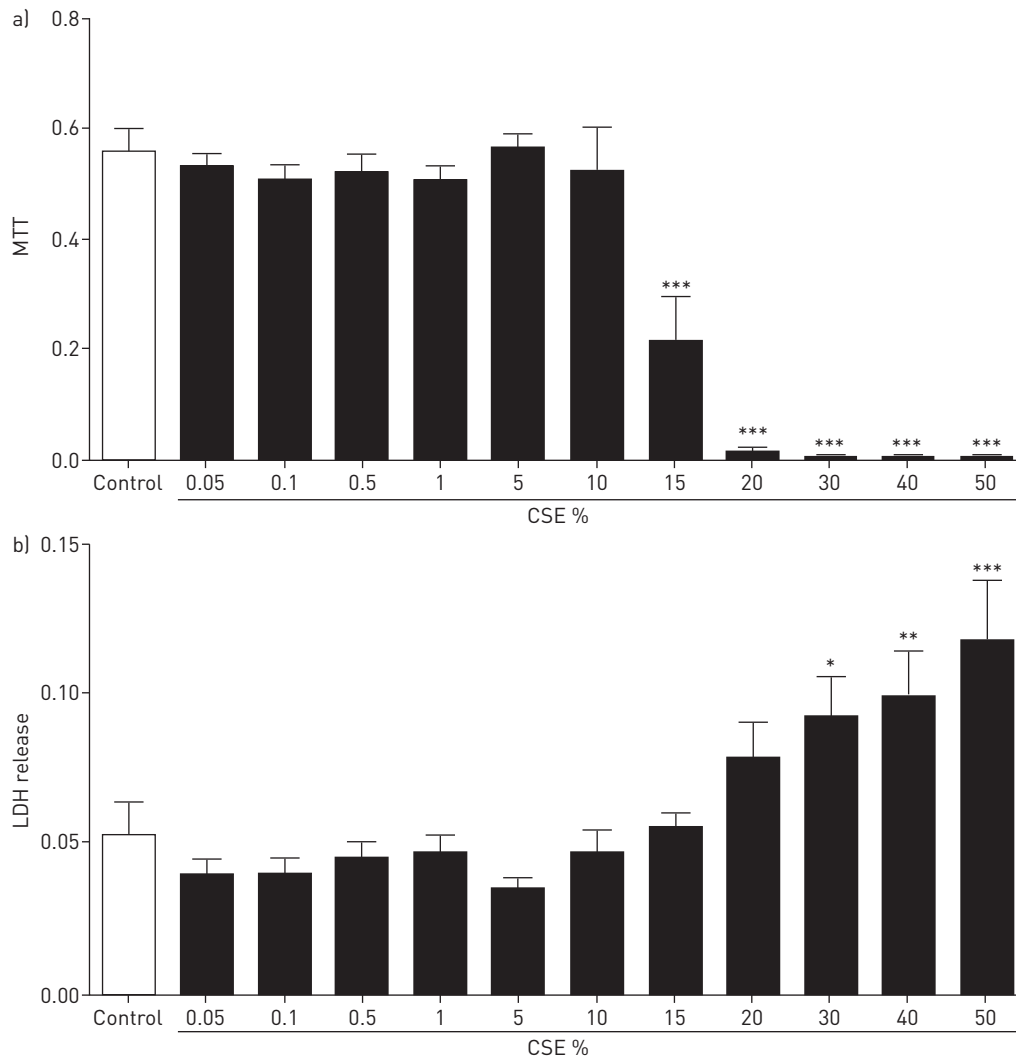


FIGURE 1 The effect of cigarette smoke extract (CSE) on cell viability. The cytotoxicity of CSE on a) the mitochondrial activity and b) lactate dehydrogenase (LDH) release of human airway smooth muscle (ASM) cells was measured by Thiazolyl blue tetrazolium bromide (MTT) and LDH assays at an absorbance of 570 nm/690 nm and 490 nm/690 nm, respectively. Human ASM cells were stimulated with serial dilution of CSE for 72 h (n=4). Data are presented as mean  $\pm$  SEM. One-way ANOVA plus Bonferroni post-test was used to determine statistical significance. \*:  $p < 0.05$ ; \*\*:  $p < 0.01$ ; \*\*\*:  $p < 0.001$ , compared with control.

### CSE induces chemokines

As CSE is known to induce CXCL8 release from human ASM cells [21], we used this output to validate our *in vitro* model. CXCL8 release was increased by 10% CSE and 10 ng·mL<sup>-1</sup> of TGF- $\beta$ 1 in both non-COPD and COPD groups (fig. 2a and b). The release of CXCL1 was increased by 10% CSE in the non-COPD group and increased by both 5% and 10% CSE in the COPD group; however, TGF- $\beta$ 1 did not induce the release of CXCL1 in either group (fig. 2c and d). There were no differences in basal or maximum production of CXCL8 and CXCL1 between ASM cells from these two groups. Our results show that the release of CCL5 and CXCL10 from ASM cells were too low to be detected (the detection limit of these assays are 15.625 pg·mL<sup>-1</sup> and 31.35 pg·mL<sup>-1</sup>, respectively). We found that CCL2 was produced by ASM cells, but this was not induced by CSE or TGF- $\beta$ 1, and there were no differences between the non-COPD and COPD groups (data not shown).

### CSE effects transcription factors DNA binding activity

Our results show that neither CSE nor TGF- $\beta$ 1 increased the activity of transcription factor NF- $\kappa$ B (fig. S1a and b); however, 5% CSE increased the activity of transcription factor AP-1 in only the non-COPD group (fig. S1c and d).

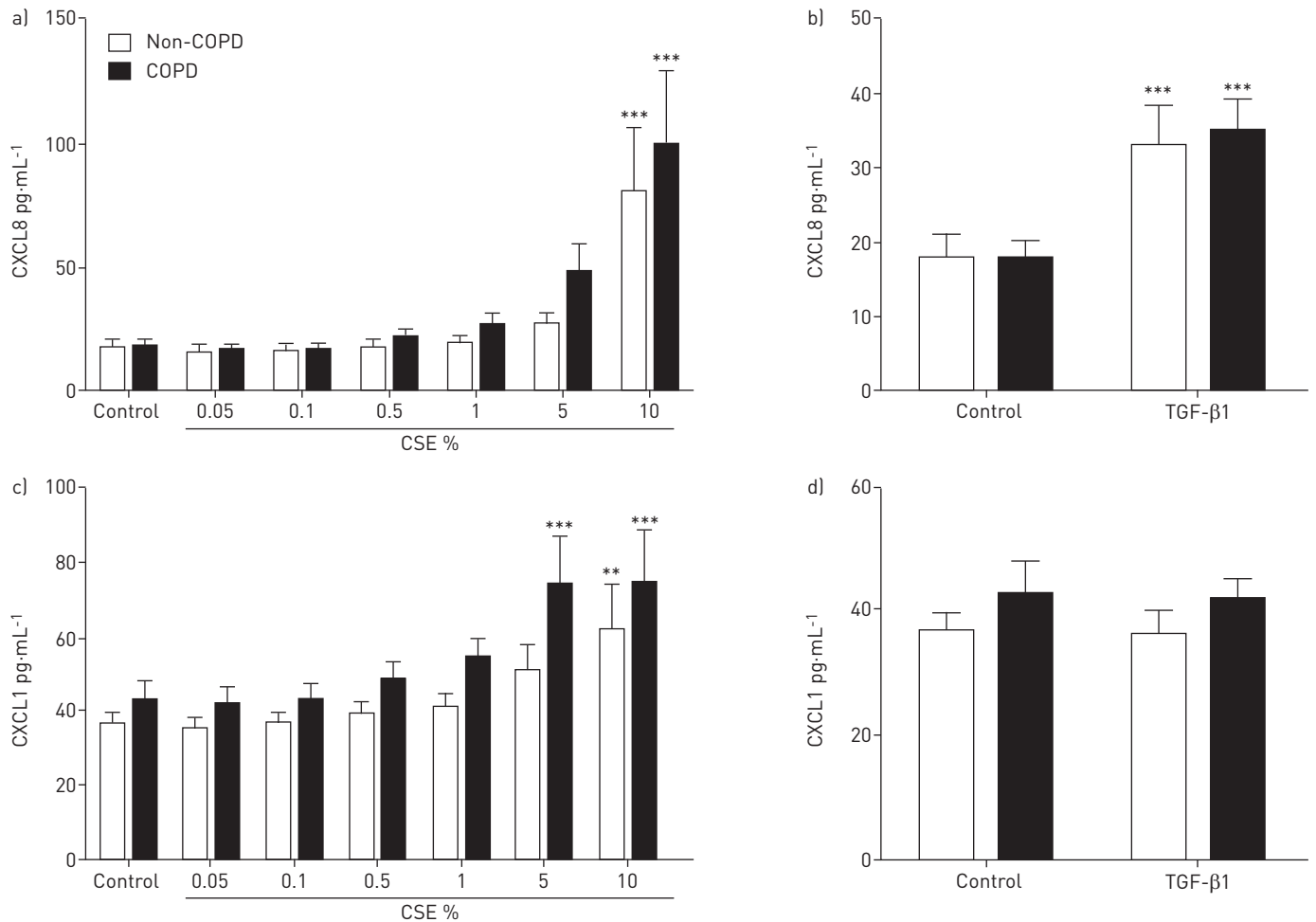


FIGURE 2 Release of CXCL8 and CXCL1 from human airway smooth muscle (ASM) cells. The concentrations of a, b) CXCL8 and c, d) CXCL1 in supernatant from human ASM cells from subjects with (n=9) and without (n=9) chronic obstructive pulmonary disease (COPD) after 72 h stimulation with cigarette smoke extract (CSE) or transforming growth factor (TGF)-β1 were measured by ELISA. Data are presented as mean ± SEM. Two-way ANOVA plus Bonferroni post-test was used to determine statistical significance. \*\*: p<0.01, \*\*\*: p<0.001, compared with control.

#### Gene expression of ECM and adhesion molecule-related genes

To investigate the potential effects of CSE on airway remodelling we used a PCR-based array as a screening tool. The gene expression of 70 ECM proteins, adhesion molecules and MMPs were assessed in response to CSE and TGF-β1 stimulations (fig. S2). CSE upregulated more ECM- and adhesion molecule-associated genes than TGF-β1, and there were differences in MMPs gene expression between CSE and TGF-β1 stimulations. Using a  $\geq 2$  cut-off line and/or  $>1.5$  fold differences between COPD and non-COPD group, MMP-1, -2, -3, -10 and -12, and COL5, COL7 and COL8A1 were further investigated.

#### CSE induces MMPs

To verify the MMP changes, we measured the release of MMP-1, -2, -3, -10 and -12 from human ASM cells. After stimulation with either CSE or TGF-β1, the concentration of total MMP-12 in the supernatants from human ASM cells was too low to be detected (data not shown). The concentration of total MMP-1 was increased by 10% CSE in the non-COPD group and increased by 5% and 10% CSE in the COPD group (fig. 3a and b). Neither CSE nor TGF-β1 affected the production of MMP-2 (fig. 3c and d). 10% CSE increased the release of MMP-3 and MMP-10 in both groups, whilst TGF-β1 increased the release of MMP-3 and MMP-10 in the COPD group only (fig. 3e-h).

#### The production of active MMP-1

To further investigate the effect of CSE on the activity of MMP-1, we measured the active form of MMP-1 from human ASM cells. The concentration of active MMP-1 was increased by 10% CSE in both groups, while TGF-β1 decreased the release of active MMP-1 in the non-COPD group only (fig. 4).

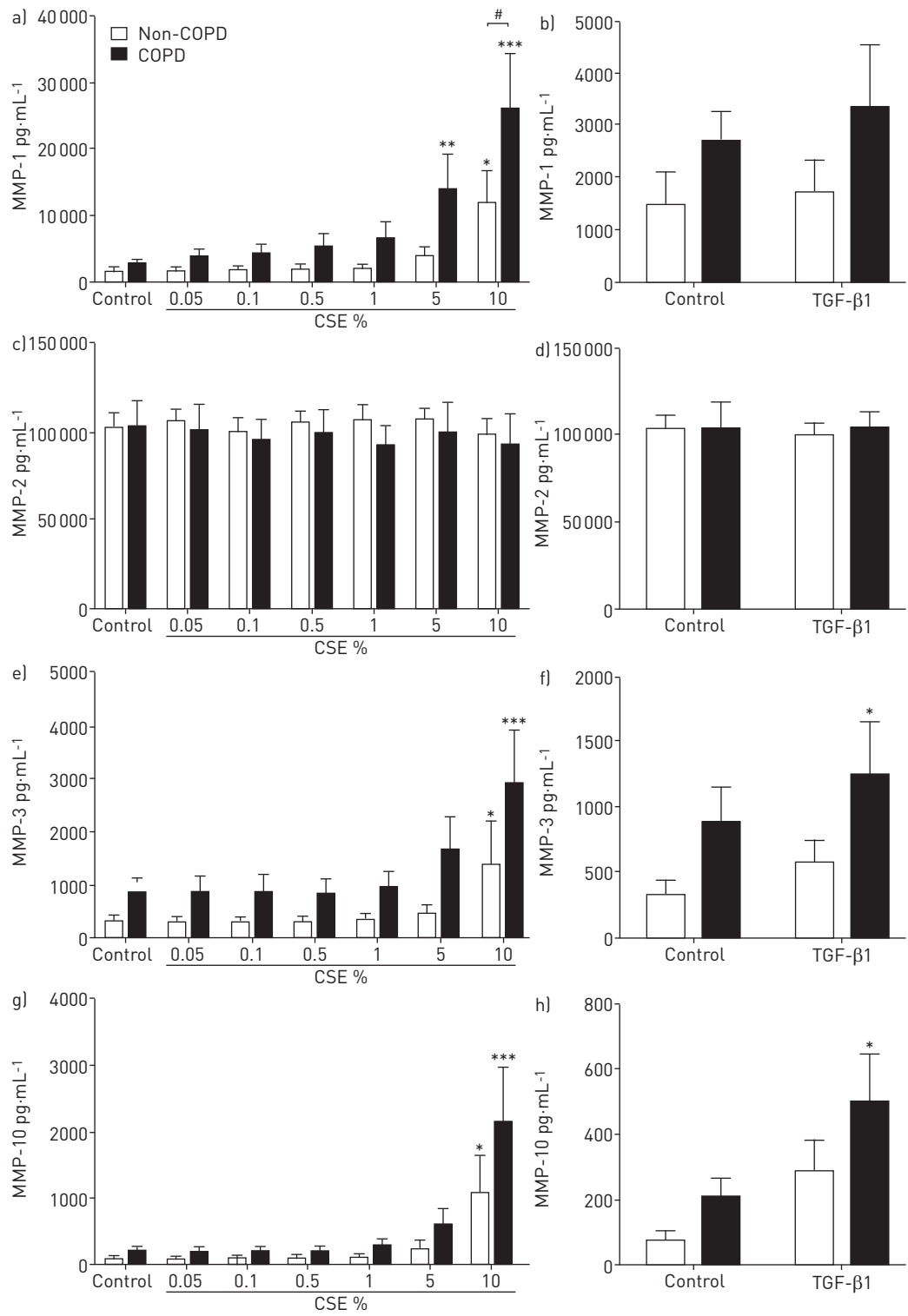


FIGURE 3 Production of matrix metalloproteinases (MMPs) from human airway smooth muscle (ASM) cells. The concentrations of a, b) MMP-1, c, d) MMP-2, e, f) MMP-3, and g, h) MMP-10 in supernatant from human ASM cells from subjects with (n=8) and without (n=7) chronic obstructive pulmonary disease (COPD) after 72 h stimulation with cigarette smoke extract (CSE) or transforming growth factor (TGF-β1) were measured by ELISA. Data are presented as mean ± SEM. Two-way ANOVA plus Bonferroni post-test was used to determine statistical significance. \*: p<0.05; \*\*: p<0.01; \*\*\*: p<0.001, compared with control. #: p<0.05, comparison between two groups.



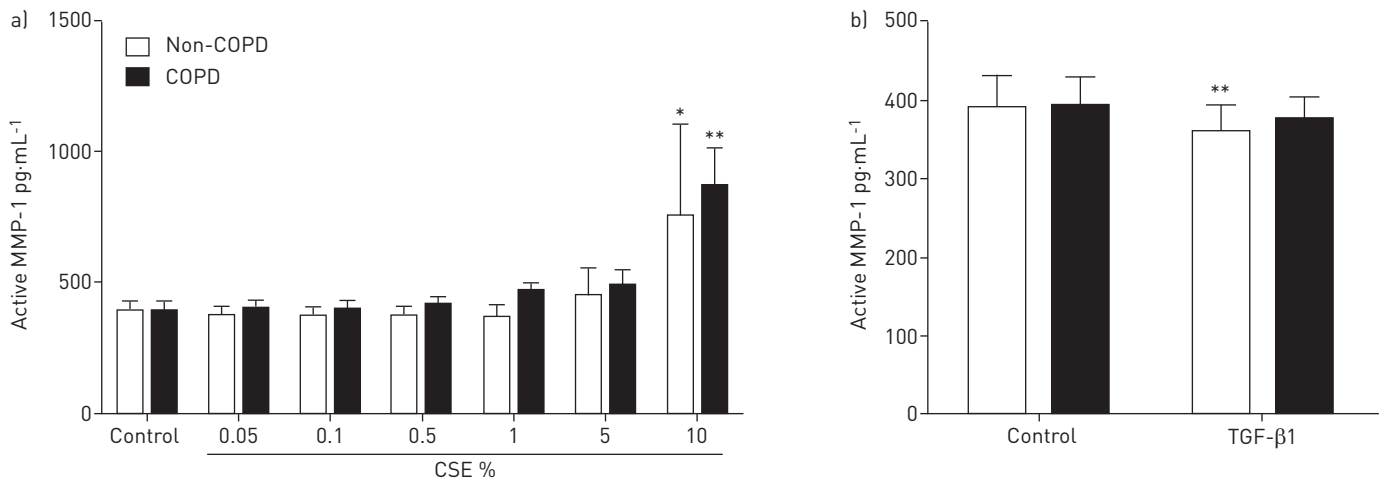


FIGURE 4 Production of active matrix metalloproteinase (MMP)-1 from human airway smooth muscle (ASM) cells. The concentrations of active MMP-1 in the supernatants from human ASM cells from subjects with (n=5) and without (n=5) chronic obstructive pulmonary disease (COPD) after 72 h stimulation with cigarette smoke extract (CSE) or transforming growth factor (TGF)-β1 were measured by ELISA. Data are presented as mean ± SEM. Two-way ANOVA plus Bonferroni post-test was used to determine statistical significance. \*: p < 0.05; \*\*: p < 0.01, compared with control.

#### The deposition of ECM proteins

To verify the alteration of ECM proteins after stimulation with CSE or TGF-β1, we assessed the deposition of COL5, COL7 and COL8A1 in the ECM. The deposition of COL7 was too low to be detected (data not shown). CSE did not alter the deposition of COL5 (fig. S3). The deposition of COL8A1 was increased by 0.5%, 1%, 5% and 10% CSE only in the COPD group, and there was significantly more COL8A1 induced by 1% and 5% CSE from the COPD cells compared to the non-COPD cells (fig. 5a). TGF-β1 increased the deposition of COL8A1 in the COPD group only (fig. 5b). CSE did not alter the deposition of fibronectin from either group which is in stark contrast to previous findings in fibroblasts [12], and CSE inhibited the deposition of perlecan in human ASM cells from both groups (fig. S4).

#### The release of ECM proteins

To verify the alteration of soluble ECM proteins after stimulation with CSE or TGF-β1, we assessed the soluble COL1 and soluble fibronectin in the supernatants using Western blots. Our results showed that neither CSE nor TGF-β1 significantly increased the release of COL1 (fig. S5). Our results also showed that CSE did not induce the release of soluble fibronectin; however, TGF-β1 significantly increased the release of fibronectin (fig. S6).

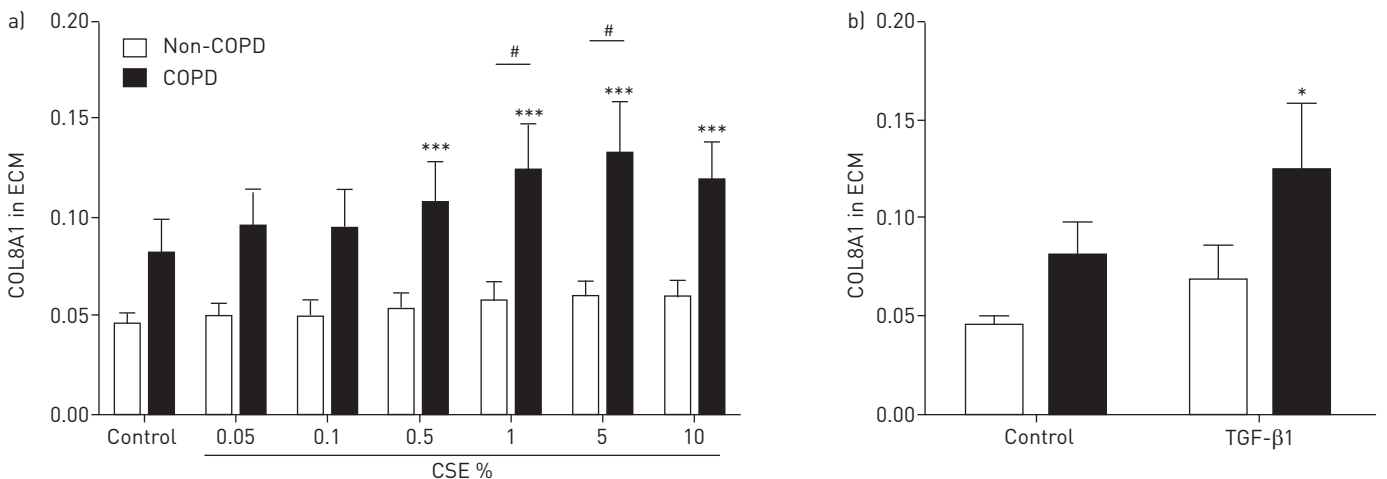


FIGURE 5 The deposition of collagen VIII alpha 1 (COL8A1) from human airway smooth muscle (ASM) cells. The deposited COL8A1 in the extracellular matrix (ECM) from human ASM cells from subjects with (n=7) and without (n=7) chronic obstructive pulmonary disease (COPD) after 72 h stimulation with cigarette smoke extract (CSE) or transforming growth factor (TGF)-β1 was measured by ECM ELISA at an absorbance of 450 nm/570 nm. Data are presented as mean ± SEM. Two-way ANOVA plus Bonferroni post-test was used to determine statistical significance. \*: p < 0.05; \*\*\*: p < 0.001, compared with control. #: p < 0.05, comparison between two groups.



**COL8A1 in airway bronchus**

As we found greater CSE-induced COL8A1 by the COPD ASM cells, we next investigated if our *in vitro* findings were reflective of COPD *in vivo*. Immunohistochemistry revealed COL8A1 was expressed in airway tissue from patients with (n=10) and without COPD (n=7). In the airways from both groups the COL8A1 appeared to be localised in the basement membranes, vascular walls and ASM bundles. The positive staining was controlled by threshold and isotype control staining. Using densitometric analysis (quantification of staining area) we found there was higher overall expression of COL8A1 in the COPD group (fig. 6).

**CSE inhibits cell attachment and wound healing**

To further investigate the effect of CSE on the function of human ASM cells, we assessed cell attachment of ASM cells from people with (n=5) and without (n=5) COPD. CSE significantly decreased the attachment

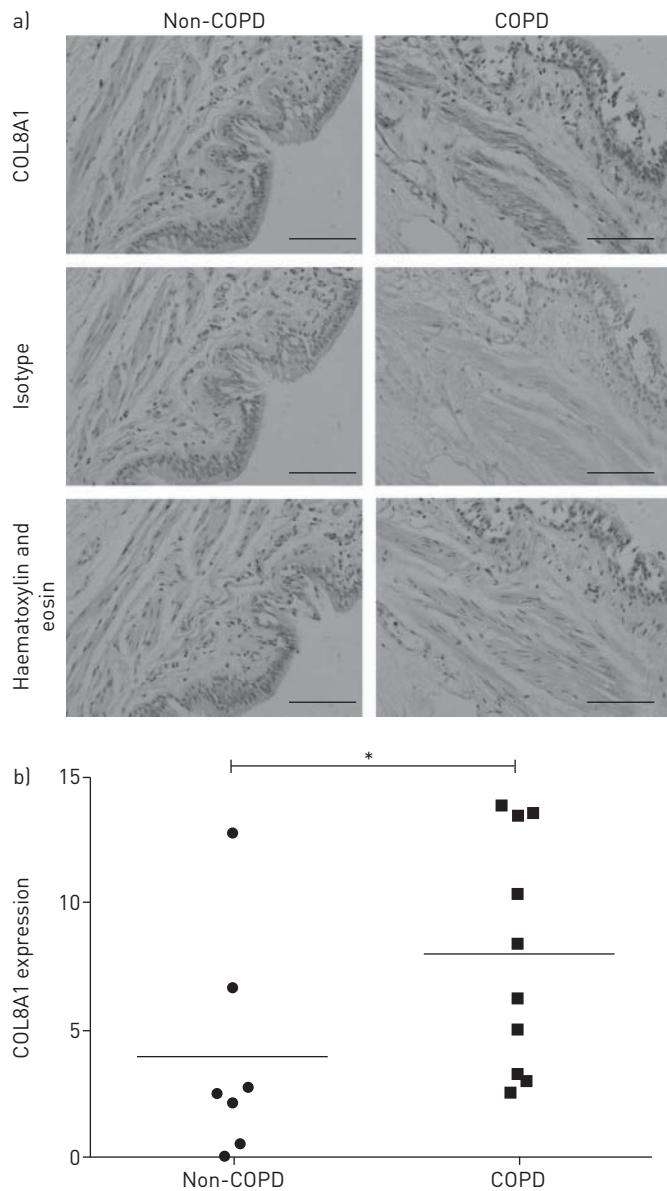


FIGURE 6 Collagen VIII alpha 1 (COL8A1) airway tissue staining. a) COL8A1 in airway bronchus from subjects with (n=10) and without (n=7) chronic obstructive pulmonary disease (COPD) was measured by immunohistochemistry. Specific staining was detected using a chemical chromophore, DAB (brown), and the cell nucleus was counterstained with haematoxylin (blue). Tissue structure was stained with haematoxylin and eosin. Scale bars=100 µm. b) Immunostaining of COL8A1 in non-COPD and COPD groups were quantified by positive staining area and corrected with isotype control. Data are presented as median. The Mann–Whitney test was used to determine statistical significance. \*: p<0.05.

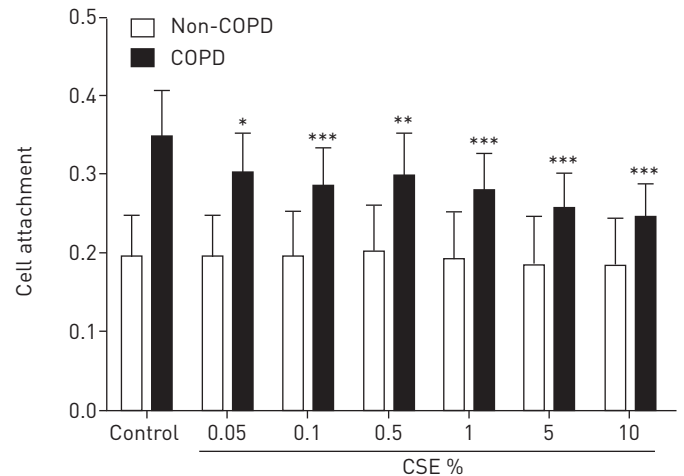


FIGURE 7 The effect of cigarette smoke extract (CSE) on 2-h cell attachment at an absorbance of 595 nm. Cell attachment on a CSE treated plate of human airway smooth muscle cells from subjects with ( $n=5$ ) and without ( $n=5$ ) chronic obstructive pulmonary disease (COPD) were measured by toluidine blue assay. Data are presented as mean  $\pm$  SEM. Two-way ANOVA plus Bonferroni post-test was used to determine statistical significance. \*:  $p<0.05$ ; \*\*:  $p<0.01$ ; \*\*\*:  $p<0.001$ , compared with control.

of ASM cells to culture plates from the COPD group only (fig. 7). Wound healing assays showed that high concentration of CSE significantly decreased the rate of wound healing in human ASM cells ( $n=6$ ) (fig. S7).

### Discussion

We have found differential responses to CSE in ASM cells from smokers with and without COPD. Specifically MMP-1 and the deposition of COL8A1 in ASM cells were increased by CSE, and these increases were higher in the COPD group. Our results also showed that CSE decreased ASM cell attachment to culture plates and wound healing specifically in cells isolated from smokers with COPD. These findings suggest that ASM cells from smokers with COPD are more sensitive to CSE stimulation which may explain, in part, the development of COPD in some smokers.

COPD is an inflammatory disease characterised by an increased number of neutrophils [9, 22], and increased amount of neutrophil chemokines (such as CXCL8 and CXCL1) in bronchoalveolar lavage fluid [23, 24]. CSE is a potent inducer of CXCL8 in ASM cells [21, 25]; however, whether hypersecretion of CXCL8 occurs in ASM cells isolated from patients with COPD as in other airway cells was not known [26, 27]. Therefore, we also measured CSE-induced CXCL8, and found CSE increased the release of CXCL8 from ASM cells yet without differences between cells from people with and without COPD. We also measured CSE-induced CXCL1 and found CSE increased the release of CXCL1 from ASM cells. Furthermore, ASM cells from COPD patients were more sensitive to CSE stimulation for the production of CXCL1. Both CXCL8 and CXCL1 have similar biological properties, in that they both have effects on the recruitment of neutrophils [28–30]. In our study we used ASM cells from smokers with and without COPD. The induction of CXCL8 and CXCL1 by CSE in ASM cells from both groups may reflect the observation that neutrophils are increased in COPD patients [31] and smokers without COPD [32]. Furthermore, as low concentrations of CSE induced CXCL1 in the COPD cells only, this suggested that these cells were hyperresponsive to CSE, and may explain why some smokers develop COPD and others do not.

Our results showed that high concentrations of CSE increased the release of both proMMP-1 and active MMP-1 (interstitial collagenase) from human ASM cells, and this seemed more pronounced in smokers with COPD than those without COPD. When we measured only active MMP-1 we found similar production between cells from both groups. This indicates that production of active MMP-1 in ASM may not be a key determinant of lung pathology in COPD, but may be related to processes common to both smokers with and without COPD. In other cells, CSE increased the production of MMP-1 from human epithelial cells and human lung fibroblasts which appear to be driven primarily through the extracellular regulated kinase-1/2 mitogen-activated protein kinase pathway [33, 34]. We also found human ASM cells constitutively produced high levels of MMP-2 (gelatinase A) and this was not increased by CSE, which is in contrast to findings in cigarette smoke-exposed fibroblasts [35].

Compared to other MMPs, MMP-3 (stromelysin 1) and -10 (stromelysin 2) have not been extensively studied to date. Our study showed that CSE increases the gene expression of MMP-3 and -10 in human

ASM cells, and that 10% CSE increases production of MMP-3 and -10. These results indicate that both MMP-3 and -10 may lead to different progression in smokers with COPD. Two genotyping studies indicated that MMP-3 polymorphisms associate with disease progression in COPD [36, 37]. Another study showed the expression of the MMP-10 gene was increased in both small airways and the parenchyma surrounding small airways in association with progression of COPD [38]. In addition, our results found MMP-12 (macrophage elastase) gene expression to be increased by CSE *in vitro*. However, the release of MMP-12 protein from CSE stimulated ASM was lower than the detection limit of the assay ( $9.2 \text{ pg}\cdot\text{mL}^{-1}$ ).

It has previously been shown that CSE increased the deposition of fibronectin and perlecan from COPD fibroblasts [12], so in this study we also measured their production by ASM cells. However, CSE did not affect the production of fibronectin in COPD ASM cells, and decreased the production of perlecan indicating that responses to CSE are cell type specific. Our array data showed that several collagens also changed in response to CSE stimulation, so we chose to evaluate COL5, COL7 and COL8. The protein level of COL5 did not change in response to CSE, and the tools to measure COL7 were unreliable. However, CSE induced greater COL8 production from COPD than from control ASM cells. There is little known about COL8 in COPD, especially not on the amount of COL8 in airways of people with and without COPD. In this study we found that the expression of COL8 was increased in COPD airways, particularly in and around the smooth muscle bundles, suggesting that the smooth muscle produces COL8 *in situ* and indicating that the deposition of COL8 from smokers with COPD is likely to contribute to the airway pathology. COL8 has a short triple helix and contains  $\alpha 1$  and  $\alpha 2$  chains, and each  $\alpha$  chain contains a collagenous domain, a short N-terminal non-triple-helical region (NC2) and a longer C-terminal non-triple-helical domain (NC1). As is known for other collagens, different regions of COL8 can have opposing biological effects. For example, the entire COL8 molecule increased aortic smooth muscle cell proliferation and migration [39], whilst the NC1 domain of COL8A1 inhibited the mitochondrial activity of bovine aortic endothelial cells [40]. COL8 has not been reported previously as a determinant of ECM in COPD, but exposure of pregnant mice to pollution resulted in increased COL8 in the tubular cells in the kidney of offspring [41]. This raises the question as to whether similar hereditary effects could occur in the offspring of pregnant mothers who smoke. In addition, the functional and long-term outcomes of such exposures have not been examined.

We have not investigated if any interaction between MMP-1 and COL8 occurs, and have not been able to find specific examples in the literature. It would be tempting to speculate that the increased MMP could degrade the collagen; however, we found that they both increased at the same time, showing that the net effect is collagen deposition.

In COPD there is impaired repair in the small airway walls and alveolar walls [6, 7]. Our results show that CSE only reduced cell attachment in human ASM cells from patients with COPD, and higher concentration of CSE also decreased the wound-healing rate of ASM cells. Those results may indicate an innate difference of ASM cells from smokers with and without COPD. One study about the effect of CSE on the function of the human lung has shown that CSE reduces the migration and contractile activity of normal human bronchial smooth muscle cells [42]. In another study it was shown that CSE impairs the wound healing of bovine bronchial epithelial cells *via* a reactive oxygen species dependent mechanism [43]. Another study has shown CSE inhibits the proliferation of human lung fibroblasts [44], but whether these fibroblasts were derived from patients with COPD or not was not clear.

To investigate if differences in transcription factor activity could account for the increased responsiveness of the COPD ASM cells to CSE we chose to measure the activity of the transcription factors NF- $\kappa$ B and AP-1, as these have previously been shown to be involved in the release of MMPs and cytokines from ASM cells [45, 46]. We found no evidence of increased transcription factor activity in COPD ASM cells. This suggests that the increased response to cigarette smoke in the COPD cells may occur due to epigenetic changes, as we have found in other cells in COPD [47].

In conclusion, we showed the differential production of chemokines, MMPs and collagens by human ASM cells from people with and without COPD in response to cigarette smoke stimulation. ASM cells isolated from subjects with COPD showed a higher response to cigarette smoke in the release of inflammatory mediators and factors associated with airway remodelling and cell behaviour. Our data from this manuscript suggests that the ASM in COPD is capable of responding to the soluble components of cigarette smoke. This would act to recruit neutrophils (through the release of chemokines) and potentially affect other airway cells through the altered deposition of ECM. Since the amount of smooth muscle is positively correlated with COPD severity these effects may be amplified in severe COPD. Therefore, our findings suggest that ASM cells from smokers with COPD contribute to the pathological development of this disease.

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