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Inflammation and remodelling in experimental models of COPD - Mechanisms and therapeutic perspectives

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Chapter 1

GENERAL INTRODUCTION

COPD

Chronic obstructive pulmonary disease (COPD) is defined as “a disease state characterized by airflow limitation that is not fully reversible. The airflow limitation is usually both progressive and associated with an abnormal inflammatory response of the lungs to noxious particles or gases” (1). Tobacco smoke exposure is the major risk factor for the development of COPD, although other forms of insult to the lungs as well as genetic predisposition may also contribute (1, 2). COPD has been predicted by the Global Burden of Disease Study to become the third leading cause of death worldwide in 2020 (3).

Innate and adaptive immune system in COPD

The inflammatory response in COPD consists of infiltration of various innate immune cells, including macrophages and neutrophils, as well as adaptive immune cells, such as T and B lymphocytes (4-6). Macrophages clear the lung of inhaled particles, bacteria and apoptotic cells by phagocytosis. After phagocytosis, the macrophages are removed by mucociliary clearance, which constitutes an important mechanism of defense against infection. However, the phagocytic capacity of alveolar macrophages may be decreased in COPD (7), whereas the epithelial cilia function is impaired by cigarette smoke, resulting in increased inflammation in the lung (8-10). In addition to the removal of foreign material from the lung, macrophages are also considered important regulators of immune responses in the lung due to their ability to release various pro-inflammatory cytokines and chemokines. Macrophages release tumor necrosis factor- α (TNF- α) and interleukin-1 β (IL-1 β), cytokines which induce local and systemic inflammatory responses as well as increased mucus production. Furthermore, the release of chemokines such as interleukin-8 (IL-8) and leukotriene B₄ (LTB₄), which are potent neutrophil chemoattractants, may potentiate neutrophilic inflammation in the lung (4-6, 11, 12). In addition, macrophages are a source of proteolytic matrix metalloproteinases (MMPs) as well as reactive oxygen species which inhibit antiproteases, both of which may contribute to the development of emphysema (4, 6, 11, 12).

Neutrophils in the lung are also a source of proteolytic enzymes, including neutrophil elastase, an elastolytic enzyme which is considered to play a key role in the development of emphysema. In addition to its role in the breakdown of lung tissue, neutrophil elastase has also been implicated in increased mucus production and decreased mucociliary clearance, as well as stimulation of IL-8 and pro-fibrotic transforming growth factor- β (TGF- β) release. Neutrophils also release IL-8 and LTB₄, thereby perpetuating neutrophilic inflammation.

Azurophilic granules in neutrophils contain the enzyme myeloperoxidase (MPO), which participates in the formation of hypochlorous acid, a strong oxidant which may inactivate antiproteases in the lung, thus further augmenting tissue breakdown by elastolytic enzymes. Additionally, released MPO may result in lipid peroxidation and the formation of reactive nitrogen species, leading to tissue injury (4, 6, 13, 14).

Various studies have demonstrated increased numbers of T and B lymphocytes in the lungs of COPD patients, suggesting a role for the adaptive immune system in this disease (6, 13, 15-17). The numbers of T cells in the parenchyma and peripheral airways were shown to correlate with the degree of emphysema and airflow obstruction, respectively, suggesting an important role for these cells in the pathogenesis of COPD (15, 16, 18). Although CD4⁺ and CD8⁺ T cells are both increased in the lung, CD8⁺ cells are the predominant type. CD8⁺ T cells may contribute to tissue breakdown in COPD by releasing cytotoxic perforins and granzymes (6, 19). CD8⁺ cells in sputum of COPD patients were shown to be highly activated and express high levels of perforins, associated with increased cytotoxic activity of these cells in vitro (20). In addition, the degree of CD8⁺ cell activation in peripheral blood of COPD patients correlated with disease severity (21). Due to their strength in numbers, CD8⁺ cells have received more attention in COPD research than the CD4⁺ cells. The role of CD4⁺ cells in COPD is still unclear, but it has been suggested that they contribute to the development of adaptive immune responses by priming and prolonging survival of CD8⁺ cells as well as by taking part in the activation and differentiation of B cells (19).

B cell follicles have been observed in both the parenchyma and the airways of COPD patients (17, 18, 22, 23). Although the small airway area occupied by these follicles correlates with airflow obstruction in COPD (18), the contribution of B cells to the pathogenesis of this disease is unknown (24). There is evidence that B cells produce immunoglobulins, which are directed against antigens in the lung (25). The nature of these antigens is presently unclear, however. It has been suggested that the immunoglobulins may be directed against microbial antigens, since there is increased colonisation in the lungs of COPD patients (18). Alternatively, the protein content of cigarette smoke may be a source of antigens, as well as protein adducts formed in the lung as a result of cigarette smoke exposure (17). In addition, immunoglobulins directed against elastin peptides have been found in plasma of COPD patients, indicating that the breakdown of extracellular matrix proteins may result in the formation of antigens (26).

Structural changes in the lung

The persistent inflammatory response in the lung may lead to tissue damage and result in structural changes in the lung. The major structural abnormalities which develop in COPD include airway remodelling and emphysema. Emphysema is the destruction of alveolar structures, which leads to impaired gas exchange in the lung. In addition, loss of elastic recoil due to a decreased number of alveolar attachments results in collapse of the airways, contributing to airflow limitation (5, 27). As mentioned above, various inflammatory cells have been implicated in the development of emphysema and a particularly important role is attributed to elastolytic enzymes, such as neutrophil elastase.

Airway remodelling in COPD is characterized by peribronchial fibrosis, mucus gland hypertrophy, goblet cell hyperplasia and increased airway smooth muscle mass (5). These structural abnormalities result in thickening of the airway wall and mucus plugging of the lumen, thereby contributing to airflow obstruction. Although some features of airway remodelling may be observed in the large airways, they are most pronounced in small airways (<2 mm diameter), which are considered the major site of airflow obstruction in COPD (18, 27-29).

The pro-fibrotic growth factor TGF- β and its downstream Smad signalling pathway may be importantly involved in fibrotic processes in COPD (5). Increased TGF- β expression has been found in the airway epithelium of smokers and COPD patients (11, 30, 31). In addition, cigarette smoke exposure was shown to increase TGF- β expression and to induce Smad signalling in rat tracheal explants and in mouse lung (32-34). Furthermore, exposure of latent (recombinant) TGF- β to cigarette smoke in a cell-free system resulted in the release of active TGF- β (35).

Mucus gland hypertrophy and goblet cell hyperplasia in COPD result in mucus hypersecretion, which has been associated with morbidity and mortality in this disease (36-38). In the peripheral airways of COPD patients increased expression of the mucin MUC5B is observed in the lumen, whereas MUC5A/C expression is increased in the epithelium (39). The increased presence of mucus in the airways, combined with a cigarette smoke-induced deficiency of the ciliary apparatus, impairs the mucociliary clearance, leaving the lung susceptible to microbial colonisation (5, 6, 8, 36).

Airway smooth muscle remodelling is discussed separately below.

In addition to the airway remodelling, structural changes in the pulmonary vasculature have also been observed in COPD (5). Pulmonary vascular remodelling in COPD is characterized by thickening of the vessel wall, due to proliferation of the intima and the thickening of the media. In addition, muscularization of microvessels, which lack a smooth muscle layer under healthy condition, has also been observed (5). Remodelling of the pulmonary vasculature may contribute to pulmonary hypertension, a co-morbidity of COPD, which is present in a large proportion of the patients and may manifest itself at rest or during exercise. The mechanisms underlying pulmonary vascular remodelling in COPD are not known, but may involve hypoxia or pulmonary inflammation, leading to endothelial dysfunction and increased expression of growth factors in the vessel wall (5, 40-45). In addition, pulmonary hypertension results in remodelling of the right ventricle, which becomes hypertrophied as a result of the increased afterload (41).

Extrapulmonary manifestations of COPD

COPD may be associated with a wide range of extrapulmonary manifestations, including cardiovascular disease, loss and dysfunction of skeletal muscle, diabetes, osteoporosis, anemia, increased gastro-oesophageal reflux and clinical depression and anxiety (46, 47). These may be the result of a spill-over of lung inflammation into the circulation, causing low grade systemic inflammation or, alternatively, systemic inflammation which does not originate from the lung (46-49).

Pharmacological treatment of COPD

Currently, inhaled β -adrenergic receptor agonists and anticholinergics constitute the main bronchodilator therapy, whereas inhaled corticosteroids are the major anti-inflammatory therapy in COPD (50). Although glucocorticosteroids may be used to reduce exacerbations (51) and some recent studies have indicated that long-term therapy with high doses of inhaled corticosteroids, with or without long-acting bronchodilators, may decrease inflammation and the rate of lung function decline in (subgroups of) COPD patients (52, 53), the effectiveness of inhaled corticosteroids on progression of COPD is still a topic of discussion (54). Thus, it appears that the sensitivity to glucocorticosteroids is reduced in most patients with COPD, which may involve inhibition of histone deacetylase-2 by oxidative stress (54). PDE4 inhibitors, such as roflumilast, are a novel class of anti-inflammatory agents bearing perspectives in (severe) COPD (55).

The role of airway smooth muscle in the pathogenesis of COPD

Airway smooth muscle area may be increased in the small airways of COPD patients (16, 18, 56, 57). This contributes to the increase in airway wall area which leads to airflow obstruction. Indeed, both airway smooth muscle mass (16) and total airway wall area (18) were shown to correlate with a decrease of FEV₁ in COPD. Airway smooth muscle from COPD patients was shown to have an increased capacity to generate force in vitro (58, 59); the generated force negatively correlated with both FEV₁/FVC ratio and FEV₁ in these patients (59). These data suggest that increased area and altered function of airway smooth muscle may contribute to airflow obstruction and hyperresponsiveness in COPD (27).

In addition to their role as contractile cells regulating airway diameter, airway smooth muscle cells may also act as synthetic cells, producing chemokines, growth factors and extracellular matrix proteins in response to various G-protein-coupled receptor agonists, growth factors, pro-inflammatory cytokines or cigarette smoke (28). The increased release of chemokines, such as IL-8, may contribute to airway inflammation, whereas increased deposition of extracellular matrix proteins and the release of growth factors may contribute to fibrosis as well as airway smooth muscle cell proliferation (28, 60-67).

Cell proliferation is a mechanism underlying the increased airway smooth muscle area in obstructive airways diseases (68, 69) and may be induced by the release of growth factors from structural cells of the airway wall, such as epithelial cells, macrophages or even airway smooth muscle cells themselves (60, 70, 71). Growth factors and other mitogens, including various G-protein-coupled receptor agonists, cytokines and extracellular matrix proteins, can induce a proliferative phenotype of airway smooth muscle cells (66, 72-81). This phenotype is characterized by an increased rate of proliferation as well as a decreased expression of contractile proteins and decreased contractile function (74, 76-80). Phenotypic modulation is a reversible process; removal of the proliferative stimulus or altered extracellular matrix expression can induce maturation of the cells to a normo- or even hypercontractile phenotype, resulting in airway smooth muscle tissue with an increased force-generating capacity (74, 77, 82, 83).

Mitogen-activated protein (MAP) kinase pathways have been shown to play a major role in growth factor-induced signalling and the induction of a proliferative, hypocontractile phenotype of airway smooth muscle (72, 75, 77, 78, 84, 85). Even though airway smooth muscle mass may be increased in COPD, the potential contribution of disease-relevant stimuli, such as cigarette smoke or

lipopolysaccharide (LPS), to airway smooth muscle proliferation and phenotype modulation by direct action on the airway smooth muscle cells has not yet been addressed.

In vitro models of COPD

Cigarette smoke

Cigarette smoke extract (CSE) is the cornerstone of in vitro COPD modelling. CSE is made by passing CS through cell culture medium or a physiological buffer solution. There is no standard for preparing CSE. The number of cigarettes, volume of cell culture medium and type of cigarette used vary between different laboratories. Furthermore, cigarettes may be burned with or without filter and the CSE may be used as is or filtered. Therefore, a detailed description of the procedure used to prepare the CSE is required for comparing results obtained in different laboratories.

Various cell types have been exposed to CSE in order to mimic CS exposure in the lung. These include structural cells such as fibroblasts, epithelial and airway smooth muscle cells (62-65, 86-88) as well as inflammatory cell types such as macrophages and neutrophils (89-91). Upon CSE exposure, functional responses and gene expression changes, as well as the underlying intracellular signalling mechanisms, have been evaluated in these cells. Experiments in various cell types, including bronchial epithelial and airway smooth muscle cells as well as macrophages and neutrophils, have indicated that CSE induces increased release of the neutrophil chemokine interleukin-8 (IL-8) (63-65, 86, 89, 90). Some of these studies revealed that oxidative stress and activation of NF- κ B and MAP kinase pathways play a role in this process (63, 64). In addition, experiments using CSE exposure of macrophages and neutrophils have indicated that TLR receptors may play a key role in CS-induced IL-8 release (89, 90). Various studies have also indicated that CSE may induce cell death (92-95). However, in the majority of these studies cells are exposed for prolonged periods of time (92, 93), whereas smokers' lungs are usually exposed for several minutes at a time. Effects of short, pulsatile CSE-stimulation of cells are still unknown.

CS instead of CSE has also been used to stimulate epithelial cells (96, 97), since these cells are directly exposed to inhaled air in the lung and this type of exposure may therefore be suitable for studying the effects of CS. Exposure of other airway wall cells, such as fibroblasts and airway smooth muscle cells, may possibly be more suitably modelled using CSE rather than CS, as in vivo components of CS may only reach these cells by diffusion through the airway wall.

Although most *in vitro* models have focused on CS or CSE exposure of cultured cells, airway or lung tissues have also been exposed to these stimuli (33, 98-100). A particularly elegant model using rat tracheal explants embedded in agar and exposed to CS on the epithelial side, thereby mimicking airway CS exposure *in vivo*, allows determination of the processes in the airway wall that are activated by CS without the involvement of immune cells (33, 101). Studies using this model have revealed that CS may induce pro-fibrotic changes in the airway wall in the absence of an inflammatory response. These findings were subsequently confirmed using an *in vivo* mouse model, showing that the pro-fibrotic gene expression in the airways is increased by CS before inflammatory cell numbers are increased (32).

Components of CS

Cigarette smoke is a complex mixture of over 5300 identified components (102). Exposure of cells to known components of CS, such as acrolein, formaldehyde or acetaldehyde, has been used to elucidate the contribution of these components to the release of pro-inflammatory cytokines and mucus production (103-106). This approach is indeed useful for identifying the contribution of individual CS components to the pathophysiology of COPD, but may be less suitable as a model as the contribution of a large number of other components is excluded.

Bacterial components

Stimulation of cells with bacteria (*P. aeruginosa*, *H. influenzae*) or bacterial components, such as LPS, as well as viruses (RSV, rhinovirus) has been used to model exacerbations of COPD. Airway infections have been associated with exacerbations of COPD, which contribute to the progression of the disease (107, 108). Increased levels of LPS have been demonstrated in BAL fluid from COPD patients, indicating that it may be a relevant stimulus for disease progression (109). LPS has been shown to induce IL-8 release from lung fibroblasts as well as from epithelial and airway smooth muscle cells (110-112). LPS has also been shown to induce connective tissue growth factor release or mucin production from bronchial epithelial cells, indicating a possible contribution of LPS to fibrotic processes and mucus hypersecretion in the airways (113, 114).

Elastase

Elastase is a proteolytic enzyme, which is released by activated neutrophils in the lung and is considered to be a major contributor to the breakdown of alveolar tissue, resulting in emphysema (115). However, several studies have shown that elastase may induce pro-inflammatory gene expression and IL-8 release from bronchial epithelial cells (115-119). In addition, neutrophil elastase has also been

shown to induce mucus production by these cells (120-122). Elastase was also found to lower the ciliary beat frequency of epithelial cells, suggesting a role in impaired bacterial clearance in COPD (123). Mechanisms underlying the effects of elastase treatment appear to be complex and may involve activation of protease-activated receptors (PARs) (124, 125) or the proteolytic release of epidermal growth factor (EGF) from lung fibroblasts (126, 127). Interestingly, similar to CS, elastase was found to activate TLRs, supporting the importance of these receptors in the pathophysiology of COPD (128, 129).

Animal models of COPD

Animal models of disease are used for studies, which for ethical reasons cannot be performed in humans. The major disadvantage of these disease models is that they are per definition based on species different from human. However, once this limitation is considered when interpreting results, animal models can be used to gain insight into specific pathophysiological processes and putative therapeutic interventions. Each animal model has its own specific advantages and disadvantages; therefore, the suitability of a particular model will highly depend on the aims of the study.

In the last 30 years, modelling of COPD in animals has largely been performed in small laboratory animals, such as mice, rats and guinea pigs. Mice offer the advantage of a wide range of molecular and genetic tools, such as antibodies and knock-out mice. On the other hand, the lung and airway structures, the distribution of mucus glands and goblet cells, the autonomic innervation of the airways and airway vascularisation in guinea pigs closely resemble that of humans whereas in mice and rats they do not (130). Despite the fact that several studies have used rats, these animals appear to be relatively resistant to the development of experimental COPD (131).

Cigarette smoke exposure

Cigarette smoke exposure is the major risk factor for the development of COPD and is therefore considered by many to be the ideal choice for modelling COPD (131). The protocols used for the exposure of animals vary greatly between the different studies. Thus, there is large variation in the length, frequency and number of exposures, the type of exposure (nose-only or whole-body exposure) and the type of cigarette used. Exposure of small animals to cigarette smoke has been performed in commercially available or home-made CS exposure apparatuses of varying degrees of complexity (131). The nose-only exposure system involves extensive handling and restraining of the animals, which may be

very stressful. Whole-body exposure results in deposition of CS particles on the pelt, which can be ingested during grooming; however, the animals are unrestrained and the exposure requires less animal handling. It has been demonstrated that animals suffered less weight loss after whole body exposure than after nose-only, suggesting this method may be preferred in order to avoid unwanted effects due to handling and restraining stress (131-133). Nevertheless, the nose-only cigarette smoke exposure has successfully been used for modelling COPD in animals (17, 134, 135) Because animals may change their breathing patterns as an avoidance reaction to inhalation of CS it has been suggested that monitoring of serum cotinine or blood carboxyhaemoglobin should be performed in order to quantify the exposure (131, 132, 136).

Despite these limitations and challenges, CS exposure has been shown to induce various features of COPD in animals, including pulmonary infiltration of macrophages and neutrophils, airway fibrosis and emphysema (32, 131, 134, 137-141). Although emphysema has consistently been reported after chronic CS exposure (131, 137-141), the degree varies between studies. This may be due to variation in the smoking protocol or species and strain differences (131, 142)

In guinea pigs, CS has also been shown to induce pulmonary hypertension (PH), associated with pulmonary vascular remodeling, both of which are observed in patients with COPD (135, 143-145). In addition, CS-induced PH has also been reported in rats but not in mice, although vascular remodeling does occur in the latter species (131, 146).

A major advantage of CS-models may be that the stimulus used is a major contributor to the development of COPD in human subjects. CS-exposure of experimental animals induces several pathological features of COPD. Another drawback is that CS-induced models are costly and time-consuming, as these models require 5 days per week CS-exposure for several hours, during 6 months. The severity of disease in these models is rated as GOLD stage 2, which indicates mild disease (131).

Elastase model of emphysema

This model consists of instillation of elastolytic enzymes in the lung resulting in tissue damage. A single instillation results in the development of emphysema. Early studies by Gross and colleagues (147), using papain instillation were instrumental in the establishment of the protease/anti-protease hypothesis and were a major contribution to the understanding of the role of α 1-antitrypsin deficiency in the development of emphysema.

Although intuitively the mechanism of elastase-induced emphysema should be simple, involving destruction of tissue structure by enzymatic digestion, several lines of evidence suggest a much more complex mechanism. The half life of instilled elastases in the lung was found to be as short as 45-50 min (148), whereas the enlargement of alveolar spaces is progressive over a period of days and continues therefore after the exogenously applied elastases are no longer present in the lung (149-151). In accordance, treatment with elastase inhibitors prior to or immediately after instillation of elastase results in inhibition of airspace enlargement, whereas treatment 4 or 8 h post instillation does not (152-155). Elastase instillation induces an inflammatory response in the lung. Thus, increased expression of TNF- α , IL-1 β , IL-6 and IL-8 as well as infiltration of macrophages and neutrophils have been reported, suggesting that inflammation might play a role in the development of elastase-induced emphysema (156, 157). Accordingly, mice lacking TNF- α - and/or IL-1 β - receptors are strongly protected against the development of elastase-induced emphysema (156, 158).

Major advantages of the elastase model are the technical ease of inducing disease by a single instillation of the enzyme in the lung and the ability to control disease severity by adjusting the amount of enzyme (131). This model is particularly suited to study potential mechanisms of emphysema and regeneration processes in the lung. However, the model is not suitable for studying airway remodeling.

Starvation-induced model of emphysema

Starvation has been shown to induce emphysematous changes in the lung parenchyma. The study which revealed this phenomenon for the first time was conducted in the Warsaw Ghetto during World War II, where autopsy findings showed that a high percentage of people who died from starvation (13.5%) had emphysema (159). Subsequently, a number of studies reproduced this observation in rats, by subjecting the animals to severe caloric restriction for periods of several weeks. These studies confirmed that starvation induces emphysema-like changes as well as changes in lung mechanics. Reduced number of alveoli, increased alveolar volume and decreased alveolar surface area were reported (160-164). More recently, a study in anorexic subjects has shown that long-term caloric restriction is associated with loss of lung tissue and that in anorexic subjects the body mass index correlates with the diffusion capacity of carbon monoxide in the lung (165). However, despite similarities between the emphysematous changes induced by starvation, and emphysema in COPD, there are major differences in the pathology of these two conditions. The abnormal enlargement of airspaces and alveolar wall destruction observed in COPD is permanent, whereas the emphysema-like changes induced by starvation are

being reversed to normal levels after sufficient caloric intake (166, 167). Moreover, in rats with elastase-induced emphysema, calorie restriction induces a further increase in emphysema severity which is reversed by refeeding back to levels observed in animals treated with elastase alone (167). Furthermore, starvation-induced emphysematous changes, unlike emphysema in COPD, are not associated with airflow obstruction (160). In addition, this model lacks inflammatory cell influx in the lungs and structural changes in airways and pulmonary vasculature, which are observed in COPD (131). These data suggest that the starvation-induced emphysematous changes are likely the result of abnormal lung maintenance or growth. Therefore, this animal model is not well suited to study COPD.

LPS-induced model of COPD

LPS is a component of the cell wall of gram-negative bacteria, which is present as a contaminant in cigarette smoke, air pollution and organic dusts (168, 169). Exposure to environmental dusts containing LPS has been shown to induce chronic airflow obstruction and has been associated with the development of COPD in farmers (170, 171). In animals, a single LPS exposure induces pulmonary inflammation, characterized by infiltration of neutrophils and macrophages, as well as airway hyperresponsiveness (172-176). In addition, an increased number of airway mucus cells has also been observed after a single LPS exposure (177-181). Animals chronically exposed to LPS develop pulmonary inflammation as well as structural changes in the lung which are characteristic for COPD. Chronic LPS exposure induces increased numbers of macrophages and neutrophils (180, 182-188). Furthermore, CD8(+) T-cells as well as peribronchial and perivascular lymphocytic aggregates containing both CD8(+) and CD4(+) T-cells and B-cells were found to be increased in mice after chronic LPS exposure (188). Interestingly, 12 weeks of twice-weekly LPS exposure resulted in increased gene expression of the pro-inflammatory cytokines TNF- α , IFN- γ and IL-18 as well as increased numbers of macrophages and lymphocytic aggregates after 1- and 8-week recovery periods (188). This indicates that, similarly to COPD (189, 190), chronic LPS exposure induces inflammatory responses, which persist even after the stimulus is no longer present.

In addition to inflammation, chronic LPS exposure has been shown to induce airway remodelling. Increased numbers of Goblet cells have been observed in airways of both mice and guinea pigs after repeated LPS exposure (180, 188). Chronic LPS exposure also resulted in increased peribronchial deposition of collagen and thickening of the airway wall in mice (182, 184-188, 191). Although 12 weeks of twice weekly LPS exposure resulted in increased airway smooth

muscle mass in mice (188), such changes were not observed in guinea pigs after 3 weeks of 3 times per week LPS exposure (180). The cause of this discrepancy could be the difference in the number of exposures, but species differences might also play a role. Chronic LPS exposure has also been shown to induce emphysema in mice, guinea pigs and hamsters (188, 192-195). Vernooij and colleagues showed that the airway remodelling and emphysematous changes are still present in mice 8 weeks after the LPS exposure had ceased (188). In addition, Brass et al. showed that airways of mice exposed to LPS for 4 or 8 weeks, followed by 4 weeks of recovery, still show submucosal thickening and increased collagen deposition, associated with pro-fibrotic gene expression (186, 191). The numbers of goblet cells in the airways of LPS-exposed mice, although still above control levels, appeared to be decreased after 8 weeks of recovery, indicating this may be a reversible feature of LPS-induced disease (188). Collectively, these data indicate that chronic LPS-exposure induces persistent inflammatory responses in the lung as well as alterations of airway and lung structure, which closely resemble the pathological changes observed in COPD.

Several studies have shown that the LPS-mediated effects are dependent on the Toll-like receptor 4 (TLR4). Thus, features of LPS-induced disease, including AHR, inflammation and airway remodelling, were considerably impaired in mice expressing deficient TLR4 (182) or lacking LPS-binding protein (183) or CD14 (196), which are both required for the activation of TLR4 by LPS. In addition, LPS-induced AHR, inflammation and airway remodelling were inhibited by treatment of mice with a TLR4 antagonist, E5564 (Eritoran) (187).

The complex mechanisms underlying the LPS-induced inflammation and structural changes in the lung are not fully known. LPS induces the release of pro-inflammatory cytokines including TNF- α , IL-1 β , IL-8, IL-6 and GM-CSF in the lung (175, 183, 188, 196-198). This leads to the influx and activation of inflammatory cells, which may contribute to the development of LPS-induced disease. In addition, LPS-induced activation of TLR4 in macrophages and/or neutrophils may perpetuate inflammation due to an additional IL-8 and TNF- α release (199, 200).

IL-1 appears to play an important role in LPS-induced pathophysiology. Thus, IL-1 receptor antagonist (IL-1Ra) was found to partially inhibit LPS-induced airway inflammation in rats, whereas IL-1R-knockout mice were protected against development of LPS-induced airway wall thickening and fibrosis, indicating a crucial role for this receptor in chronic LPS-induced airway inflammation and remodelling (184, 201).

Interestingly, TLR4 has also been shown to mediate CS-induced inflammation *in vivo*. Pulmonary infiltration of inflammatory cells as well as increase of pro-inflammatory cytokines in the BAL fluid after up to 5 weeks of CS exposure was strongly reduced in mice lacking a functional TLR4 (202, 203). In addition, IL-1R-knockout mice were similarly protected against CS-induced pro-inflammatory responses in the lung (202). These data indicate that signalling pathways required for the development of LPS-induced pathological changes also play an important role in the inflammatory responses to CS.

Major advantages of the LPS-induced COPD model are the wide range of pathological features which are induced over a relatively short period of time, requiring only 2-3 LPS exposures per week. Several studies have indicated that TLR4 may play a role in CS-induced inflammatory responses (89, 202, 203) as well as in the development of COPD (170, 171), suggesting that LPS is a relevant stimulus for modeling COPD. This model is also suitable for testing therapeutic interventions and is less labour intensive and less costly than CS-exposure. Like CS-exposure, chronic LPS is considered to induce relatively mild disease, up to now (131). Although LPS and CS may share common mechanisms to induce disease it is important to realize that there still may be differences in the etiology of disease.

Caenorhabditis elegans model of COPD?

Recently, one group has exposed *Caenorhabditis elegans* (*C. elegans*) to CS in order to study its effects on gene expression of the innate immune system while eliminating any involvement of the adaptive immune system, which is not present in this nematode (204). The authors showed that *C. elegans* tolerates up to 4 h of CS exposure, as this had no significant effect on the rate of mortality measured at 48 h post exposure. Furthermore, CS exposure had profound effects on gene expression and resulted in impaired bacterial clearance. Several genes, which had human orthologues were selected for further investigation using RNAi techniques. This approach led to the conclusion that CS-induced downregulation of *lbp-7* (human orthologue: fatty acid binding protein 5 (FAB5)) plays a major role in bacterial clearance in *C. elegans*. In order to determine the relevance of this finding for COPD, the authors demonstrated that FAB5 mRNA expression is increased by bacteria in primary human bronchial epithelial cells and that cells isolated from COPD patients show lower FAB5 mRNA levels than cells from healthy smokers.

Another recent study used a similar approach to investigate the role of hypercapnia - which may occur in COPD as a result of impaired gas exchange - in

gene expression as well as motility and muscle morphology (205). Exposure of *C. elegans* to increased CO₂ levels resulted in altered gene expression and decreased motility, which was associated with abnormal muscle fiber organization. These data support a role for hypercapnia in skeletal muscle wasting observed in COPD (46, 48, 49).

These two studies show that *C. elegans* can be used to study potential mechanisms involved in COPD. Exposure of *C. elegans* to conditions or factors relevant to human disease could be a powerful source of information due to its ease of use, low cost, fully sequenced genome and availability of RNAi tools, providing the obtained results can be translated to mammalian models and human disease. The use *C. elegans* as a model species for COPD does however have major drawbacks. Most importantly, since *C. elegans* is a nematode, it does not have lungs. Therefore, although *C. elegans* offers many possibilities for investigating functional implications of altered gene expression and could thus be used for studying specific processes related to human disease, it is not a disease model of COPD.

Cellular and molecular mechanisms

The understanding of cellular and molecular mechanisms underlying COPD is limited compared to other obstructive lung diseases such as asthma. A major goal of our investigations was to define specific mechanisms involved in inflammation and airway remodelling in this disease, using CSE- and LPS-exposed airway smooth muscle cells *in vitro*, as well as a newly developed guinea pig model of LPS-induced COPD *in vivo* and *ex vivo*. The *in vitro* studies were focused on signalling mechanisms involved in CSE-, LPS- and growth factor-induced changes in airway smooth muscle phenotype and function. The animal model studies addressed the role of the cholinergic system and the NO-arginase axis as two major, interrelated pathways that are likely to be involved in the inflammatory, structural and functional changes observed in COPD.

Cholinergic mechanisms

Increased cholinergic tone has been identified as the major reversible component of airflow limitation in COPD (206). Acetylcholine release from parasympathetic nerve endings results in contraction of airway smooth muscle and increased mucus release. This contribution of acetylcholine to airflow limitation is the basis for the use and effectiveness of anticholinergics as bronchodilators in COPD (207). However, recent studies have suggested that anticholinergics, in addition to bronchodilatation, may have other beneficial effects. Data from the UPLIFT

trial show that the long-acting anticholinergic tiotropium bromide reduces exacerbation frequency, all-cause mortality and the number of adverse respiratory and cardiac events in COPD patients (208, 209). In addition, a subgroup analysis indicated that tiotropium reduces the rate of lung function decline in patients with mild COPD (210), young patients (211) and in those not on other controller medication (212). Recent observations in an acute mouse model of CS exposure suggest that tiotropium may reduce pulmonary neutrophilia by reducing the expression of pro-inflammatory cytokines and chemokines in the lung (213). The effects of tiotropium on structural remodelling in COPD are still unknown.

However, tiotropium has been shown to inhibit allergen-induced airway eosinophilia as well as airway smooth muscle thickening and increased mucin expression, suggesting that acetylcholine contributes to airway inflammation and remodelling in asthma (214, 215). Muscarinic receptors are expressed in both inflammatory and structural cells in the lung (207). Muscarinic agonists have been shown to induce the release of pro-inflammatory chemokines from macrophages and epithelial cells (216, 217) and to augment CSE-induced IL-8 release from airway smooth muscle cells (62). In addition, the observation that muscarinic receptor stimulation induces proliferation and collagen synthesis in lung fibroblasts suggests a role for acetylcholine in fibrosis (218, 219). Interestingly, the acetylcholine synthesizing enzyme choline acetyltransferase (ChAT) is also widely expressed in the lung (207). The observations that muscarinic receptors are present on cells which have no parasympathetic innervation and that acetylcholine can be synthesized by non-neuronal cells indicate that non-neuronal acetylcholine may also be involved in lung physiology and pathophysiology. Indeed, ChAT expression is increased in lung fibroblasts from smokers and COPD patients, suggesting that non-neuronal acetylcholine may be involved in the pathogenesis of COPD (220).

Arginase

Arginase is an enzyme which converts L-arginine to L-ornithine and urea. It is involved in the urea cycle in the liver but is also expressed in extrahepatic tissues lacking a complete urea cycle (221). Two isoforms of arginase have been identified; arginase I is a cytosolic enzyme primarily expressed in the liver, whereas arginase II is a mitochondrial enzyme expressed in extrahepatic tissue (222). Both arginase isoforms are constitutively expressed in the lung, particularly in epithelial cells, (myo)fibroblasts and endothelial cells, as well as in macrophages and neutrophils (223-226). Arginase expression and/or activity was shown to be increased in animal models of asthma (221) as well as in

asthmatic patients (227-229). In a guinea pig model of allergic asthma, increased arginase activity was shown to contribute to allergen-induced airways obstruction, hyperresponsiveness, and inflammation (230). Increased arginase activity was found to result in a decreased L-arginine bioavailability for the enzyme nitric oxide synthase (NOS). This leads to a decreased production of the bronchodilatory, anti-inflammatory nitric oxide (NO) as well as an increased production of the pro-contractile, pro-inflammatory oxidant species, peroxynitrite (231-236). Decreased production of NO may contribute to airway obstruction, inflammation and hyperresponsiveness (235). Under inflammatory conditions, inducible NOS (iNOS) is upregulated and arginase-induced low L-arginine bioavailability results in concomitant production of NO and superoxide anions (O_2^-) by this enzyme, leading to rapid formation of peroxynitrite (237). Peroxynitrite production by iNOS may contribute to allergen-induced airway hyperresponsiveness as well as inflammation in allergic asthma (230, 238). Furthermore, peroxynitrite also induces MUC5A/C expression in airway epithelium and may therefore contribute to mucus hypersecretion (239). Arginase has also been shown to play a role in TGF- β -induced collagen synthesis, due to increased formation of the collagen precursor L-proline downstream of L-ornithine (222, 240, 241). Recent studies from our group indicate that increased arginase activity contributes to airway remodelling in asthma, which may involve increased synthesis of L-proline and polyamines downstream L-ornithine, as well as reduced production of NO (242).

Although many studies have indicated an important role of arginase in the pathophysiology of asthma, there are very few studies that have focused on its potential role in COPD. Only recently, increased arginase activity was demonstrated in BAL fluid from COPD patients (243). Remarkably, similar observations were made in sputum from patients with asthma or chronic bronchitis in the late 1970s (244, 245), but this was interpreted as leaking of hepatic arginase. Increased arginase gene expression in the lung has been observed in CS-exposed rats (246) and in LPS-exposed mice (191, 247). In addition, smoking was shown to further increase arginase gene expression and immunostaining in the airways of mild asthmatics (248). Increased arginase expression and/or activity has also been found in pulmonary endothelial cells and serum of patients with pulmonary arterial hypertension (249), which may occur as a co-morbidity of COPD (5, 41). Increased arginase activity may contribute to endothelial dysfunction in pulmonary arterial hypertension by causing reduced production of eNOS-derived vasodilatory NO (249). This is further supported by the observation that oral L-arginine or inhaled NO decrease pulmonary arterial pressures in this disease (250). Despite accumulating

evidence suggesting a potential increase of arginase activity or expression in COPD, no studies have thus far demonstrated a role for arginase in the pathophysiology of this disease.

TGF- β -activated kinase 1

TGF- β -activated kinase 1 (TAK1) is a serine/threonine kinase and a member of the MAP kinase kinase kinase family (MAP3K7). It was originally identified as a mediator of Smad-independent TGF- β signalling, but has since also been shown to play a key role in pro-inflammatory signalling pathways downstream of TLR, IL-1R and TNFR (Figure 1) (251-256). TAK1 is an important activator of the NF- κ B pathway, as well as the ERK 1/2 and p38 MAP kinase pathways (251-255, 257-260). Activation of TAK1 contributes to the initiation of immune responses by inducing the release of pro-inflammatory cytokines, such as IL-8, IL-6 and TNF- α , and has also been shown to promote proliferation or survival of B-cells, T-cells and neutrophils (257, 261-264). In addition to stimuli of the above mentioned receptors, TAK1 is also activated by diesel exhaust particles as well as osmotic stress and hypoxia, indicating a role in stress-induced signalling (265-267). Considering the major role of TAK1 in TLR-signalling and the evidence suggesting a role for TLRs in the pathophysiology of COPD as well as in CS-induced signalling and inflammation, an important role for TAK1 in the development of COPD can be envisaged. However, no direct evidence has yet been reported.

In addition to its role in inflammation, several studies have indicated a major involvement of TAK1 in embryonal development (268-271). Interestingly, aberrant TAK1 signalling was shown to result, among others, in impaired development of the lung as well as abnormal vascular development lacking smooth muscle (269, 270). TAK1 has also been implicated in tissue remodelling. Increased TAK1 expression was associated with cardiac hypertrophy resulting from pressure overload in mice or with myocardial infarction in rats (272, 273). Expression of activated TAK1 in the mouse myocardium is sufficient to induce cardiac hypertrophy (273). Despite this evidence for the role of TAK1 in the development of the lung and vascular tissue as well as the remodelling of cardiac muscle tissue, there are no reports addressing the potential role of TAK1 in airway smooth muscle function.

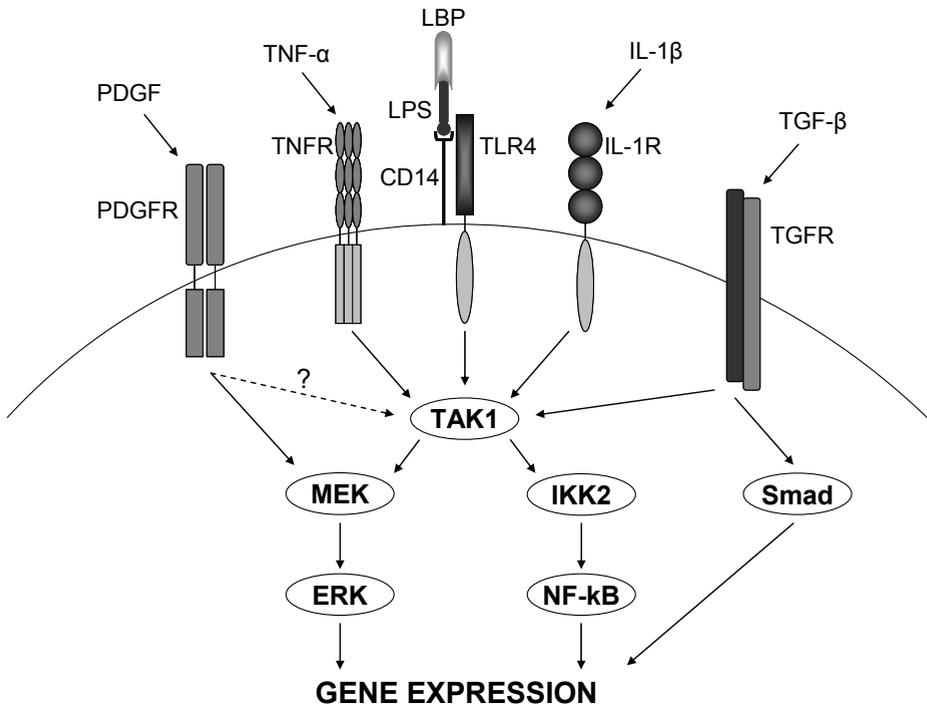


Figure 1: Pro-inflammatory and pro-fibrotic signalling in COPD.

Inhalation of cigarette smoke or other invoking factors of COPD, including lipopolysaccharide (LPS), leads to the release of pro-inflammatory cytokines and pro-fibrotic growth factors from structural and inflammatory cells in the lung. Pro-inflammatory cytokines, such as tumor necrosis factor- α (TNF- α) and interleukin-1 β (IL-1 β), activate TNF- and IL-1-receptors (TNFR and IL-1R), respectively. LPS binds LPS binding protein (LBP) and CD14, thereby activating Toll-like receptor 4 (TLR4). Subsequently, intracellular adaptor proteins of the three receptors activate interleukin-1 receptor associated kinase (IRAK) and TNF-receptor-associated factor (TRAF), leading to activation of transforming growth factor- β (TGF- β)-activated kinase 1 (TAK1). TAK1 initiates NF- κ B signalling via activation of I κ B kinase 2 (IKK2) and extracellular signal-regulated kinase (ERK) signalling via activation of mitogen activated protein kinase kinase (MEK). Among others, both signalling pathways are involved in the transcription of genes encoding for pro-inflammatory chemokines such as IL-8, which may perpetuate inflammation of the lung. Growth factors, including platelet-derived growth factor (PDGF), activate MAP kinase signalling via the small GTPase Ras and induce proliferative responses involved in airway remodelling. Although a link between PDGF and TAK1 has thus far not been demonstrated, we hypothesize that PDGFR stimulation may also lead to activation of TAK1. TGF- β activates the TGF- β receptor (TGFR), initiating both Smad and TAK1 signalling (either directly by TGFR or via the X-linked inhibitor of apoptosis (XIAP)), which may cause tissue remodeling by inducing proliferative and fibrotic responses.

Scope of the thesis

As indicated above, the objective of this thesis was to investigate mechanisms of pulmonary inflammation and remodeling in COPD by using *in vitro* and *in vivo* approaches. The focus of **Chapters 2 and 3** is phenotypic modulation of airway smooth muscle by stimuli involved in the pathogenesis of COPD. As discussed above, mitogens induce a proliferative, hypocontractile phenotype of airway smooth muscle. Proliferation of airway smooth muscle cells may cause increased airway smooth muscle mass, which has been demonstrated in COPD (18). Using cultured BTSM cells and tissue, the studies in **Chapter 2** explore the mitogenic capacity of CSE and LPS as well as the MAP kinase pathways potentially involved in the CSE- and LPS-induced responses. In **Chapter 3** the role of TAK1 in growth factor-induced ERK 1/2 signalling and modulation of airway smooth muscle cells and tissue to a proliferative, hypocontractile phenotype is investigated. The study was performed in BTSM cells and tissue as well as in primary human airway smooth muscle cells, the role of TAK1 being explored by the pharmacological TAK1 inhibitor LL-Z-1640-2 as well as the expression of dominant-negative TAK1. **Chapter 4** focuses on the potential role of airway smooth muscle cells as a source of pro-inflammatory cytokines in COPD. The studies described in this chapter investigate the role of TAK1, NF- κ B and ERK 1/2 pathways in CSE-induced IL-8 release by cultured human airway smooth muscle cells. Studies described in **Chapter 5** address the role of endogenous acetylcholine in pulmonary inflammation and remodelling using an animal model of COPD. For this purpose, a guinea pig model of LPS-induced COPD was developed. Animals were intranasally instilled, twice weekly for 12 weeks, with sterile saline or LPS and pre-treated with either saline or the long-acting anticholinergic drug tiotropium. The effect of tiotropium on LPS-induced neutrophilia, MUC5A/C expression and changes in hydroxyproline content, mean linear intercept (MLI) and lung vasculature were evaluated. **Chapter 6** is a review of the literature describing the role of arginase in various pulmonary diseases. Evidence from animal models as well as from clinical studies is presented. **Chapter 7** is dedicated to unraveling the role of arginase in the pathophysiology of COPD. Using the guinea pig model of COPD described in **Chapter 5**, the effects of pretreatment by inhalation of the specific arginase inhibitor (2S)-amino-boronohexanoic acid (ABH) on LPS-induced pulmonary inflammation and remodelling were investigated. In addition to some of the parameters evaluated in **Chapter 5** (neutrophilia, MUC5A/C expression, hydroxyproline content and pulmonary vascular dimensions), arginase activity and IL-8 levels in the lung as well as right ventricle mass were evaluated in this disease model.

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