

Pathogenesis of combined pulmonary fibrosis and emphysema

Common pathogenetic pathways

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INTRODUCTION

Emphysema is defined on pathology grounds as an abnormal permanent enlargement of air spaces distal to the terminal bronchioles, accompanied by the destruction of alveolar walls and without obvious fibrosis¹. Idiopathic Pulmonary Fibrosis (IPF) has a distinct appearance on HRCT defined as the UIP and possible UIP pattern². Although initially considered as two separate disorders (as it is clearly depicted in the definition of emphysema), emphysema and fibrosis have been found to co-exist. The widespread use of HRCT given its high accuracy for the diagnosis of emphysema and IPF helped in recognizing the co-existence of these two diseases. The association of IPF and emphysema was initially described by Wiggins et al in 1990³. The term Combined Pulmonary Fibrosis and Emphysema (CPFE) was first proposed by Cottin et al⁴, who retrospectively described a homogenous group of 61 patients with both emphysema of the upper zones and diffuse parenchymal lung disease with fibrosis of the lower zones of the lungs on chest computed tomography. The patients were all smokers, male (but one), presenting with dyspnea on exertion, relatively preserved lung volumes and a disproportionate reduction in DLco. Although in the pivotal study by Cottin, patients with connective tissue disease were excluded, the CPFE syndrome has been described in the latter group⁵. Interestingly, in a retrospective study by Tzouveleki et al⁶, a significant increased number of CPFE patients exhibited elevated serum ANA with or without positive p-ANCA titers compared to patients with IPF without emphysema. Furthermore, patients with CPFE and positive autoimmune markers (mainly ANA) exhibited improved survival compared to patients with a negative autoimmune profile. This improvement in survival was also correlated with the presence of a massive infiltration of clusters of CD20+ B cells forming lymphoid follicles⁶.

Whether the co-existence of emphysema and fibrosis is pure coincidence, due to the presence of a common predisposing factor (i.e. smoking) or it is related to shared pathogenetic mechanisms, is not yet clarified. However there are indications supporting the latter theory. Further studies are needed in order to reach a more robust conclusion.

DEFINITION

CPFE is defined by the co-existence of emphysema and pulmonary fibrosis based on HRCT. This definition is broad as it includes patients with any amount of emphysema and any type of fibrosis.

This broad definition has some limitations. The absence of a specific threshold for emphysema leads to the possible inclusion of patients with limited emphysema of no clinical significance. This can underscore the actual impact of emphysema in the natural course of CPFE. Several groups have proposed diagnostic criteria for CPFE that include a threshold of 10% emphysema visible on HRCT.

The inclusion of multiple types of ILD in the definition, leads to a heterogeneous cohort. This makes it impossible to draw comparative conclusions regarding the natural course of CPFE against a specific ILD. For example, it is well established that IPF has a better prognosis compared to iNSIP and also collagen tissue related interstitial lung diseases (CTD-ILD). Hence, the inclusion of iNSIP or CTD-ILD patients in the CPFE population can produce a bias in favor of CPFE when compared to IPF. When studying clinically meaningful endpoints (rate of FVC decline, rate of DLco decline, hospitalizations, acute exacerbations of ILD), the type of fibrotic pattern should be defined. Given the fact that IPF is the commonest ILD, special interest should be given to the syndrome emphysema-IPF, as it can have serious implications in the rate of FVC decline and thus in the evaluation of outcome and impact of therapeutic approaches.

PATHOGENESIS

The relationship between smoking and emphysema is well established. Also, smoking is known to be related with certain interstitial lung diseases as respiratory bronchiolitis interstitial lung disease (RB-ILD), desquamative interstitial pneumonia (DIP), pulmonary Langerhans' cell histiocytosis (PLCH) and IPF^{7,8}. It is noteworthy that in the latest classification of the Idiopathic Interstitial Pneumonias smoking is used as a classification element.⁹ Furthermore, in recent years there is an increasing amount of literature regarding a new form of smoking related interstitial disease linking smoking with fibrosis¹⁰⁻¹⁵. It comprises of distinctly circumscribed fibrotic lesions that radiate from a respiratory bronchiole to the pleura. The interstitial fibrosis is mixed with emphysematous changes. On HRCT these lesions correspond to localized areas of reticulation concentrated about regions of upper and mid

zone emphysema¹⁵. Several names have been proposed for this distinct interstitial lung disease as respiratory bronchiolitis-interstitial lung disease with fibrosis¹⁰, smoking-related interstitial fibrosis (SRIF)^{11,12} airspace enlargement with fibrosis^{13,14} and respiratory bronchiolitis with fibrosis (RBF)¹⁵. Although in the majority of cases RBF doesn't seem to have functional consequences, it provides an additional link between emphysema and fibrosis.

Based on the above, it is not surprising that there is an extremely strong association between smoking and CPFE. Almost all CPFE patients (98%)¹⁶ are current or former smokers. Below we will describe the pathogenesis of CPFE focusing on those aspects that support common pathogenetic routes in favor of a mere co-existence due to a common etiological factor. Smoking is a common predisposing factor for both conditions. However, it seems that in patients with CPFE there are diverse susceptible genetic backgrounds that result in the development of both emphysema and fibrosis.

1. Senescence-Telomeres

A characteristic example of such genetic background is related to accelerated senescence via telomere length shortening. A telomere is a region of repetitive nucleotide sequences at each end of a chromatid. Telomeres shorten with each cell division. Thus, they act as disposable buffers at the ends of chromosomes. When telomeres reach a critical threshold (Hayflick limit), they are sensed as double stranded DNA break and trigger a DNA damage response that provokes cellular senescence and apoptosis¹⁷. Telomerase is a specialized polymerase that renews telomeres. It has two core components: TERT, the telomerase reverse transcriptase, and TERC, the telomerase RNA that provides the template for telomere repeat addition^{18,19}.

Interestingly, telomerase mutations leading to short telomeres have been implicated in the pathogenesis of emphysema²⁰ and IPF²¹. Recently it was shown that telomerase mutations may be associated with a more severe phenotype of emphysema with a frequent occurrence of pneumothorax, especially in women²². Mutations in TERT and TERC are risk factors for pulmonary fibrosis underlying the inheritance in 8–15% of familial cases^{23,24}. Mutations in the essential telomerase genes are also found in 1–3% of sporadic IPF cases^{25,26}. Furthermore, IPF patients have shorter telomeres than age-matched controls, even when no mutation in telomerase is detected^{25,26}. Based on the above it is expected that telomerase mutations have been proposed as a possible common link for the development of CPFE^{27,28}.

Telomerase mutations have important clinical implications. Since they are not restricted to the lungs the term "telomere syndrome" has been proposed²⁸. This contradicts the classic view that IPF has no extra-pulmonary manifestations. When IPF is associated with telomerase mutations, it is in fact part of a telomere-mediated syndrome²⁹. Other manifestations of this syndrome can be present as well.

Indeed, IPF patients are at increased risk for the development of diseases that have been associated with short telomeres as liver cirrhosis²⁵, diabetes^{30,31} and bone marrow failure. Bone marrow failure can manifest as red blood cells macrocytosis, single lineage cytopenias, aplastic anemia, myelodysplastic syndrome, or acute myeloid leukemia^{23,32-34}. The syndrome complex of IPF and bone marrow failure is specific for the presence of germline defects in telomerase³⁵. Thus, in IPF patients, a high degree of suspicion for the presence of hematologic abnormalities is needed.

The presence of hematologic abnormalities can be extremely deleterious regarding outcome, when IPF patients need to receive marrow suppressive drugs in the context of chemotherapy or immunosuppressive drugs after lung transplantation. IPF patients with telomerase mutations that were subjected to lung transplantation, showed an increased rate of hematological complications and renal failure requiring renal replacement therapy³⁶.

2. SFTP

Another susceptible genetic background involves mutations in genes encoding surfactant proteins C and A2 (SFTPC and SFTPA2). Mutations in SFTPC and SFTPA2 have been implicated in the development of pulmonary fibrosis. They exert their fibrogenic effect by increasing the levels of endoplasmic reticulum stress and activating the unfolded protein response (UPR)³⁷⁻³⁹. Cottin et al reported a case of CPFE in a never smoker, 32 year old female with a dominant I73T mutation in the SFTPC gene. Her infant (bearing the same mutation) was also diagnosed with interstitial lung disease⁴⁰. Other genes involved in surfactant metabolism, such as ATP-binding cassette A3 (ABCA3) are also associated with pulmonary fibrosis⁴¹⁻⁴³. Epaud et al reported a case of CPFE in a nonsmoker, 41 year old male with mutations in the ABCA3 gene⁴⁴.

3. MMP-9 and TGF- β

Matrix-metalloproteinase (MMP)-9 has been implicated in the pathogenesis of emphysema and lung fibrosis. Sputum MMP-9 concentrations in COPD patients are directly

associated with the extent of emphysema measured by CT and MMP-9 expression levels are inversely associated with DLco⁴⁵. Regarding fibrosis, it is important that MMP-9 proteolytically cleaves latent TGF- β , leading to its activation⁴⁶. TGF- β is known to play a pivotal role in the development of IPF as it induces fibroblast chemotaxis, proliferation and transdifferentiation into myofibroblasts, and it largely promotes the production and secretion of extracellular matrix compounds, mainly collagen⁴⁷. Furthermore, it seems that TGF- β plays an important regulatory role in the transcription of MMP-9 via the Smad3 signalling pathway. TGF- β ligand activates TGF- β receptor/Smad3 signalling, which in turn collaboratively activates SIRT1 transcription with c-Jun. Subsequently, SIRT1 can deacetylate NF- κ B at lysine 30, as well as histones adjacent to the transcription factors AP-1, NF- κ B, and Pea3 binding sites of the MMP9 promoter. The final result is the suppression of MMP-9 transcription (Figure 1). When Smad3 is missing, this regulatory pathway is neutralized so that MMP9 is epigenetically activated^{48,49}. These developmental epigenetic mechanisms by which Smad3 regulates MMP9 transcription may be important in modulating both emphysema and pulmonary fibrosis. This could explain why both pathologies can appear within the same lung specimen^{49,50}.

4. TNF- α , PDGF- β

Lunndbland et al demonstrated that in transgenic mice, TNF- α overexpression caused pathologic changes consistent with both emphysema and pulmonary fibrosis⁵¹. Hoyle et al studied transgenic mice that expressed the platelet-derived growth factor-B (PDGF- β) gene from the lung-specific surfactant protein C (SPC) promoter. Results of these studies indicated that overexpression of PDGF- β induced distinct abnormalities in the developing and adult lung that led to a complex phenotype encompassing aspects of both emphysema and fibrosis⁵².

5. Neutrophil elastase

In bleomycin treated mice Lucatelli et al demonstrated that emphysematous and fibrotic lesions can be significantly attenuated by using a protease inhibitor active against neutrophil elastase. Also, in a strain of mice that developed both emphysema and fibrosis after chronic cigarette-smoke exposure, the presence of neutrophil elastase in alveolar structures was associated with a positive immunohistochemical reaction for both TGF- β and TGF- α . This led to the suggestion that neutrophil

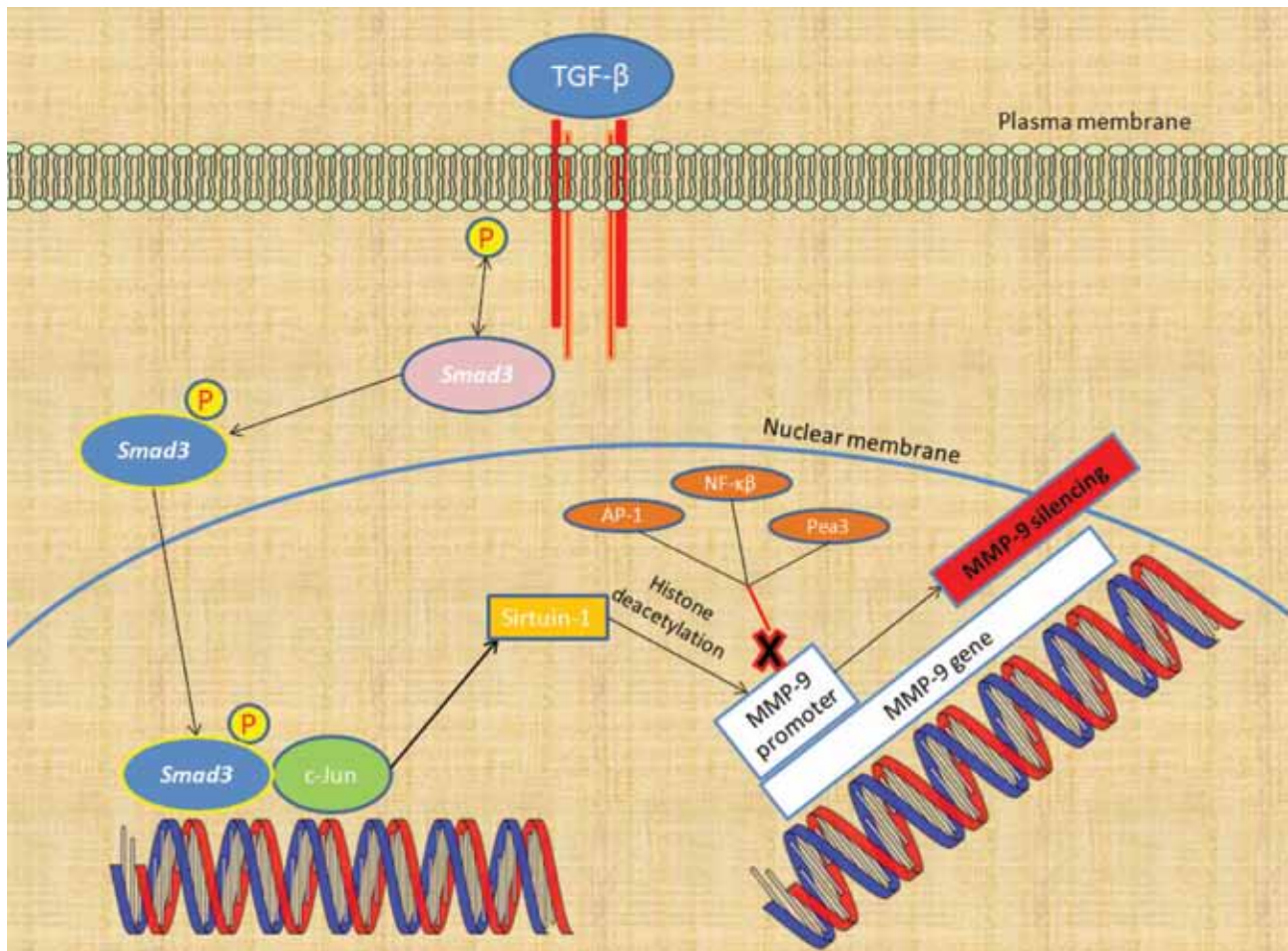


FIGURE 1. Regulation of MMP-9 transcription by TGF- β via Smad3. TGF- β ligand activates TGF- β receptor/Smad3 signalling, which in turn collaboratively activates SIRT1 transcription with c-Jun. Subsequently, sirtuin 1 deacetylates histones adjacent to the binding sites of the MMP9 promoter regarding transcription factors AP-1, NF- κ B, and Pea3. The final result is the suppression of MMP-9 transcription.

elastase could be the missing link between emphysema and fibrosis⁵³. It is possible that neutrophil elastase is involved in the release of the active TGF- β from its initial inactive complex. This complex comprises from TGF- β , latent associated peptide (LAP) and latent TGF- β binding protein (LTBP)⁵⁴.

6. CXCL5 and CXCL8

CXCL5 and CXCL8 (epithelial neutrophil activating peptide 78, and interleukin 8) have been found elevated in patients with IPF^{55,56}. CXCL8 is known to be implicated in the pathogenesis of emphysema⁵⁷. Tasaka et al⁵⁸ examined cytokine levels and differential cell counts in bronchoalveolar lavage (BAL) from patients with IPF. The concentrations of CXCL5 and CXCL8 were significantly

greater in patients with concomitant emphysema. CXCL8 concentrations were significantly correlated with the absolute numbers of neutrophils in BAL fluid and with the extent of emphysema on HRCT and inversely correlated with FVC.

7. Interleukin 13

Overexpression of interleukin 13 (IL-13) in a murine model resulted in the development of lung fibrosis, and airspace enlargement (emphysema)⁵⁹. Recently, it was shown that inhibition of IL-13 via overexpression of IL-13R α 2 resulted in inhibition of bleomycin induced pulmonary fibrosis⁶⁰. This is in line with previous observations where blocking IL-13 with tralokinumab in a humanized mouse model of IPF inhibited lung fibrosis and reduced

epithelial apoptosis in the lung⁶¹.

8. Caveolae

Caveolae are 50 to 100 nm flask-shaped invaginations of the plasma membrane enriched in cholesterol and glycosphingolipids. Caveolin-1, the structural protein component of caveolar membranes, acts as a scaffolding protein to concentrate and functionally regulate signaling molecules. Oxidative stress upregulates Caveolin-1 protein expression. Caveolin-1 plays an important role in the development of emphysema through activation of the ATM-p53-p21 pathway. The development of pulmonary emphysema was significantly inhibited in caveolin-1 null mice compared to wild type mice after 6 months of exposure to cigarette smoking⁶². Regarding IPF, caveolin-1 expression was found reduced in lung tissues and in primary pulmonary fibroblasts from idiopathic pulmonary fibrosis patients, compared with controls⁶³. It is very interesting that restoration of caveolin-1 function in skin fibroblasts from patients with scleroderma reverses their pro-fibrotic phenotype while skin from caveolin-1 null mice exhibits many characteristics found in the skin of scleroderma patients⁶⁴.

9. T-regulatory cells (Treg)

Treg (CD4+CD25+FOXP3+) play a key role in immune regulation. They generally suppress or downregulate

induction and proliferation of effector T cells⁶⁵. In IPF there is a global impairment of Treg⁶⁶. This impairment is not only numerical but also functional. Specifically, Treg from IPF patients exhibited a defective antiproliferative activity and a defective suppression of Th1 and Th2 cytokine secretion. This functional impairment was strongly correlated with a decrease in FVC, DLco and TLC⁶⁶. In lung tissue from emphysema patients⁶⁷ and also in the small airways of COPD patients, an impairment of Treg has also been observed⁶⁸. In COPD patients, Hou et al⁶⁹ dissected Treg into three distinct subpopulations based on the intensity of FOXP3 expression and the secretion of proinflammatory cytokines: denominated resting, active and cytokine secreting Treg. In accordance to the previous studies there was an imbalance between different Treg subsets in peripheral blood and BAL. Specifically, there was a downregulation of suppressive Treg and an increased proportion of the cytokine-secreting non-Treg subpopulation in patients with COPD when compared to smokers with normal pulmonary function tests⁶⁹. The impairment of Treg in emphysema and COPD is correlated with a decrease in FEV1^{68,69}. Thus, Treg impairment provides an intriguing common pathway between IPF and emphysema and a potential therapeutic target⁷⁰.

SYNOPSIS

CPFE vividly illustrates the importance of thinking

TABLE 1. Common pathogenetic mechanisms in emphysema and pulmonary fibrosis

Category	Mechanism	Comments
Susceptible genetic background	Telomerase mutations (TERT, TERC)	8–15% of familial cases and 1–3% of sporadic IPF cases
	Mutations in SFTPC, SFTPA2, ABCA3	Case reports
Cytokines	CXCL5	Increased in BALF from CPFE patients
	CXCL8 (Interleukin 8)	Increased in BALF from CPFE patients
	Interleukin 13	Possible therapeutic target
	TNF-α	Overexpression in a murine model resulted in combined emphysema/lung fibrosis
Growth Factors	TGF-β	Pivotal cytokine in pulmonary fibrosis Regulates MMP-9 transcription via Smad3
	PDGF-β	Overexpression in a murine model resulted in combined emphysema/lung fibrosis
Proteases	MMP-9	Cleavage of latent TGF-β releasing its active form
	Neutrophil elastase	Cleavage of latent TGF-β releasing its active form
Adaptive immunity	Tregs	Impaired (numerically and functionally) in IPF and emphysema patients
Cell anatomy	Caveolae	Oxidative stress acts as a first hit

beyond the obvious. A dogmatic dichotomous categorization between emphysema on one hand and fibrosis on the other has limitations. The recognition and further study of this syndrome can have serious implications. On the level of basic science the exploration of shared pathogenetic mechanisms can offer new therapeutic targets. As described this is an area of intensive ongoing research with very interesting findings (telomere length shortening, mutations in SFTPC and SFTPA2, IL-8, IL-13, TNF- α , MMP-9, Treg, Table 1) that could translate to feasible therapeutic targets. From a clinical trial perspective it is important to clarify if and how the inclusion of CPFE patients in the IPF trials can impact the results. CPFE patients tend to exhibit a delay in the reduction of FVC and DLco. This can underestimate the efficacy of an investigational drug. Further research is needed in order to seek for a better end point for CPFE. Also, it is important to accurately define CPFE as well as the comparator populations in further studies in order to produce comparable results across clinical trials. Finally, from a clinical point early detection of pulmonary hypertension is of utmost importance as it is associated with an ominous prognosis and can have impact on crucial management decisions as early referral for lung transplantation⁷¹⁻⁷³. A high degree of vigilance is required regarding the early diagnosis of lung cancer in CPFE patients as well^{74,75}. Finally, stem cell therapy is a promising approach for COPD⁷⁶ and IPF^{77,78}. Conducting clinical trials of stem cell therapy in CPFE is an intriguing project that could shed further light in the areas of pathogenesis and treatment.

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