by dabigatran in vitro and in vivo [14,15]. Dabigatran is a direct

thrombin inhibitor that reversibly binds to the active site of thrombin preventing the conversion of fibrinogen to fibrin [16]. Recently, we

have demonstrated that binding of dabigatran to thrombin prevents

cleavage of the extracellular N-terminal domain of the PAR-1

receptor [14]. In the absence of dabigatran, thrombin binds to PAR-

1, cleaves the peptide bond between residues Arg-41 and Ser-42,

thereby unmasking a new amino terminus, SFLLRN, which then can

bind to the second extracellular loop of PAR-1 and initiate receptor

signaling [7,17]. Dabigatran-bound thrombin is unable to cleave and

activate PAR-1 [14]. Further, we have shown that dabigatran inhibits

thrombin-induced in vitro differentiation of normal lung fibroblasts

to the myofibroblast phenotype and decreases CTGF, a-SMA, and

collagen type I in scleroderma lung fibroblasts [14]. To test whether

dabigatran attenuates lung injury in a murine model of SSc-ILD, we

used dabigatran etexilate, the oral prodrug of dabigatran. The prodrug

does not have antithrombin activity; however, after oral administration

dabigatran etexilate is rapidly converted by ubiquitous esterases to

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Editorial

A Passion for Scleroderma Research

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My research experience in the field of scleroderma began in year 2000 when I was recruited as a postdoctoral fellow to the laboratory of Dr. Richard M. Silver. Since that time my research interests has focused on understanding the cellular and molecular mechanisms of pulmonary fibrosis in scleroderma patients. The ongoing research in the laboratory can be subdivided in three major projects: The first is thrombin signaling in scleroderma-associated pulmonary fibrosis, and the third project concentrates on the role of genome in scleroderma.

Scleroderma (Systemic Sclerosis, SSc) is an autoimmune rheumatic disease, which affects about 300,000 Americans, primarily females who are 30 to 50 years old at onset. The leading cause of death in scleroderma patients is pulmonary dysfunction as a result of progressive interstitial lung fibrosis. The pathophysiology of pulmonary fibrosis is characterized by differentiation of fibroblasts into activated myofibroblasts, and enhanced synthesis and secretion of extracellular matrix (ECM) molecules by these cells. However, the mechanisms leading to the lung fibroblast activation and excessive deposition of collagen and other ECM proteins remain poorly understood.

In recent years, increasing evidence has accumulated to implicate involvement of the coagulation system in various fibrotic diseases, including SSc-ILD [1]. Activation of coagulation proteases, e.g. thrombin, is one of the earliest events following tissue injury. Thrombin modulates tissue repair responses by altering vascular permeability, stimulating fibroblast and neutrophil migration, and promoting adhesion and spreading of endothelial cells and fibroblasts [2]. It also activates various cell types and induces secretion of several pro-immune, profibrotic, and angiogenic factors [1-3]. Activation of these cells by thrombin is a likely mechanism for the development and progression of pulmonary fibrosis in general, and SSc-ILD in particular.

Our laboratory as well as others has demonstrated dramatically increased levels of thrombin in Broncho alveolar Lavage Fluid (BALF) from scleroderma patients with lung fibrosis and other fibrosing lung diseases [4,5]. It has been shown that BALF from normal subjects has a low level of thrombin activity, while BALF from SSc patients express up to 100-fold higher thrombin activity [4]. Elevated levels of thrombin activity have been also observed in bleomycin-induced pulmonary fibrosis in mice [6]. The majority of the cellular responses to thrombin are mediated via the G protein-coupled protease activated receptor-1(PAR-1) [2,7]. In previous studies we demonstrated that PAR-1 expression is significantly increased in patients with SSc-ILD, notably in lung parenchyma associated with inflammatory and fibroproliferative foci [8]. PAR-1 is co-localized with myofibroblasts in SSc-ILD tissue and appears to decrease during later stages of pulmonary fibrosis when a decreased number of myofibroblasts is observed. Elevated expression of PAR-1 has been also shown in patients with IPF and in a murine model of bleomycin-induced lung fibrosis [1,6].

Previously, we demonstrated that thrombin differentiates normal lung fibroblasts to a myofibroblast phenotype via the PAR-1 receptor and a protein kinase C-dependent pathway [9]. Thrombin is mitogenic for lung fibroblasts [4,8,9] and enhances the proliferative effect of fibrinogen on fibroblasts [10]. Thrombin is also a potent inducer of fibrogenic cytokines, such as transforming growth factor- β (TGF- β) [11] and connective tissue growth factor (CTGF) [12,13]. We recently showed that all this cellular effects of thrombin can be inhibited

the active moiety, dabigatran [16,18]. We observed that dabigatran etexilate significantly reduced thrombin activity and levels of TGF-B1 in BALF, while simultaneously decreasing inflammatory cells and protein concentrations in BALF [15]. Histological lung inflammation and fibrosis were significantly decreased in dabigatran etexilate-treated mice. Additionally, treatment with dabigatran etexilate reduced collagen, CTGF, and a-SMA expression in mice with bleomycininduced lung fibrosis, whereas it had no effect on basal levels of these proteins [15]. This data suggest that dabigatran etexilate may serve as a novel and attractive therapeutic agent for SSc-ILD especially for patients with over expression of thrombin. Fibroblasts isolated from SSc patients, also called activated fibroblasts or myofibroblasts, continue in vitro to generate the excess of ECM proteins and to produce various pro-inflammatory and profibrotic mediators [19]. One of the major cytokines constitutively elevated in scleroderma fibroblasts in vivo and in vitro is CTGF [13, 20,21]. Fonseca et al. [22] recently demonstrated a direct genetic association between CTGF and SSc, identifying a CTGF polymorphism within a novel SP3-dependent transcriptional repressor site that is significantly associated with the susceptibility to SSc [22]. Increased amounts of CTGF are found in scleroderma patients with more extensive skin involvement and severe pulmonary fibrosis [23]. In addition to increased tissue expression, CTGF levels are increased in the sera of patients with SSc compared with healthy controls and patients with other connective tissue diseases; also, patients with diffuse scleroderma exhibit higher levels of CTGF than those with the limited form of the disease [23]. Furthermore, broncho alveolar lavage fluid

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(BALF) from SSc patients with active lung fibrosis was found to contain much higher levels of CTGF as compared with patients without lung involvement [21].

Profibrotic properties of CTGF have been lately well established. It was shown that CTGF promotes deposition of several ECM proteins such as collagen, fibronectin, and tenascin C [20,24,25]. Using CTGFspecific small interfering RNA (siRNA), Xiao et al. [26] found that silencing of CTGF significantly decreases expression of collagen in SSc skin fibroblasts. In addition, CTGF was demonstrated to promote fibroblast proliferation, collagen gel contraction, cell adhesion and migration [13,20,25]. We recently identified a novel CTGF-induced protein, IQ motif containing GTPase activating protein-1 (IQGAP1), which is inherently elevated in lung fibroblasts isolated from SSc patients [27]. We demonstrated that expression of IQGAP1 in SSc lung fibroblasts correlates with the migration rate of cells in a wound healing assay and that IQGAP1 mediates cell migration induced by CTGF in normal lung fibroblasts [27]. The studies to understand the role of IQGAP1 in the pathogenesis of SSc-ILD are currently under progress in our laboratory.

The prognosis for pulmonary outcomes is much poorer and the mortality rate among African Americans with SSc is twice as high when compared with whites [28]. Such racial disparities in health may be explained to some extent by the effects of differences in health behaviors, socioeconomic status, and access to health care and biological factors. To understand the role of genome in scleroderma, one of our projects is focused specifically on hepatocyte growth factor (HGF) receptor, a genetic factor, which we have identified, that impacts upon the molecular mechanisms regulating the ability of HGF to control the expression of ECM proteins, such as collagen and CTGF, in the lung challenged by rheumatic disease.

We recently published the first study to demonstrate an association between deficiency of HGF and cellular mesenchymal-epithelial transition factor (c-MET) signaling and African American race. Our results demonstrated that HGF is reduced in BALF and plasma from African American SSc-ILD patients compared with white SSc-ILD patients. Moreover, in white SSc lung fibroblasts HGF down regulates extracellular matrix proteins, such as collagen and CTGF, whereas in African American SSc fibroblasts HGF is not functional due to a deficiency in phosphorylation of the HGF receptor [29].

The role of HGF as a potent anti-fibrotic factor has been established in several animal models of fibrosis [30,31]. However, the therapeutic potential of HGF for the treatment of fibrosis in humans has yet to be explored. Several recent *in vitro* studies have characterized the antifibrotic effects of HGF on collagen, MMP-1, and CTGF in SSc skin fibroblasts [32,33], whereas our lab has demonstrated the mechanisms by which HGF down regulates collagen and CTGF in human lung fibroblasts [34]. The studies to validate the genetic alteration in the c-MET gene and to determine the correlations between HGF receptor loss or reduced function and the severity of SSc-ILD among African American and white scleroderma patients are under progress.

SSc-ILD is the most serious complication and a leading cause of mortality in scleroderma for which no proven therapy currently exists. Our laboratory, like hundreds other around the world, works hard toward the understanding of etiology and pathogenesis of this disease that will ultimately lead us to the development of novel, effective therapies. It is possible that one of the potential novel therapeutics for SSc-ILD will become dabigatran etexilate which has already demonstrated profound antifibrotic effects *in vitro* and *in vivo*.

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