

**REGULATION OF ALVEOLAR EPITHELIAL CELL SURVIVAL BY
ANGIOTENSIN1-7 AND *MAS***

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ABSTRACT

REGULATION OF ALVEOLAR EPITHELIAL CELL SURVIVAL BY ANGIOTENSIN II AND *MAS*

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Apoptosis of alveolar epithelial cells (AECs) is believed to be a critical event in the pathogenesis of lung fibrosis. It was previously shown that apoptosis of AECs requires autocrine generation of angiotensin (ANG) II. ANG II can also be degraded to ANG1-7 by angiotensin converting enzyme-2 (ACE-2). ACE2 was recently found to be protective but significantly downregulated in experimental lung fibrosis and patients with IPF. In other organ systems, ANG1-7 has been found to inhibit the actions of ANGII through the ANG1-7 receptor *mas*. Therefore, it was hypothesized that ANG1-7 in the lungs might antagonize the actions of ANG II in the regulation of AEC apoptosis. To test this theory, the AEC cell line MLE-12 and primary cultures of human AECs were stimulated by the profibrotic apoptosis inducers ANG II or bleomycin (BLEO). Caspase-3 activation and nuclear fragmentation were used as markers of apoptosis and were measured along with JNK phosphorylation. Exposure to ANG II or BLEO induced caspase-3 activation, nuclear fragmentation, and JNK phosphorylation in cultured AECs. Pretreatment with ANG1-7, at a concentration of 0.1 μ M, prevented JNK phosphorylation and apoptosis. In addition, pretreatment with A779, a specific blocker of the ANG 1-7 receptor *mas*, successfully prevented ANG1-7 induced blockade of JNK phosphorylation and nuclear fragmentation. These data indicate that ANG1-7 prevents AEC apoptosis, and its actions are mediated through the ANG1-7 *mas* receptor.

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To Mom and Dad for their unconditional care, love, and support.

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TABLE OF CONTENTS

LIST OF FIGURES	viii
LIST OF ABBREVIATIONS	x
CHAPTER 1 – GENERAL INTRODUCTION	1
Idiopathic Pulmonary Fibrosis	2
Epidemiology	2
Clinical Features	3
Histological Features	4
Genetic Factors	5
Therapeutic Options.....	6
Pathogenesis of Fibrosis	8
Alveolar Epithelial Cells.....	8
Coagulation and Fibrogenesis.....	10
Profibrotic Cytokines	11
Myofibroblasts	12
Basement Membrane Disruption and ECM Remodeling	15
Apoptosis in Idiopathic Pulmonary Fibrosis.....	19
Apoptosis in General.....	19
<i>Morphological and Biochemical Changes.....</i>	19
<i>Caspases</i>	20
<i>Mechanisms of Apoptosis.....</i>	21
<u>The Extrinsic Pathway</u>	22
<u>The Intrinsic Pathway</u>	23
<u>Caspase Substrates</u>	24
<u>Other Pathways</u>	25
Alveolar Epithelial Cell Apoptosis in IPF	27
<i>Evidence of Apoptosis</i>	27
<i>Induction of Pulmonary Fibrosis</i>	28
<i>Mechanism of Alveolar Epithelial Apoptosis.....</i>	28
<u>p21 and p21 Upregulation.....</u>	28
<u>Fas-FasL Pathway Activation</u>	29
<u>TGF-β Induction of Apoptosis.....</u>	30
<u>Oxidative Stress</u>	30
<u>Alveolar Epithelial-Fibroblast Crosstalk</u>	31
The Renin Angiotensin System	33
General Ras	33
Local Ras	34
Bleomycin-Induced Experimental Pulmonary Fibrosis Model	36
Pulmonary Angiotensin System in IPF	38
General Ras	36
Local Ras	37

CHAPTER 2 – ANGIOTENSIN 1-7 REGULATES ALVEOLAR EPITHELIAL CELL SURVIVAL THROUGH THE RECEPTOR MAS	42
Abstract.....	43
Introduction.....	44
Methods.....	54
Results	56
Discussion.....	66
REFERENCES.....	69

LIST OF FIGURES

Figure 1.1. Pathogenesis of idiopathic pulmonary fibrosis [Figure 7.1 modified from (Selman et al., 2001)	17
Figure 1.2. Induction of apoptosis [Figure 11.9C.1 modified from (Siegel and Lenardo, 2002)]	26
Figure 1.3. The Renin Angiotensin System (RAS)	35
Figure 2.1. Quantitation of ISEL-positive nuclei in alveolar (A) and airway (B) epithelial cells after Bleo, CAPTO, and ZVAD-fmk administration	47
Figure 2.2. Quantitation of collagen by PR staining of alveolar (A) and peribronchial (B) lung parenchyma.)	48
Figure 2.3. Dose-dependent induction of nuclear fragmentation by BLEO in primary rat AECs and blockade by inhibitors of caspases, endonucleases, ANG-converting enzyme (ACE), and ANG-receptor interaction	49
Figure 2.4. Blockade of Bleo-induced apoptosis in primary AECs by selective caspase or ANG receptor blockers	50
Figure 2.5. ANGII induces activation of caspase-9	51
Figure 2.6. Pathways for the formation of ANG 1-7	52
Figure 2.7. Summary of Earlier Published Data.....	53
Figure 2.8. ANG1-7 inhibits ANG II induced JNK phosphorylation through the <i>mas</i> receptor.....	59
Figure 2.9. ANG1-7 inhibits BLEO induced JNK phosphorylation through the <i>mas</i> receptor.....	60
Figure 2.10. ANG1-7 inhibits BLEO induced caspase-3 activation of MLE-12 cells.....	61
Figure 2.11. ANG1-7 inhibits BLEO induced caspase-9 activation of primary AECs through the <i>mas</i> receptor	62
Figure 2.12. ANG1-7 inhibits bleomycin-induced nuclear fragmentation through the <i>mas</i> receptor in MLE-12 cells.....	63
Figure 2.13. ANG1-7 inhibits bleomycin-induced nuclear fragmentation through the <i>mas</i> receptor in primary human AECs	64

Figure 2.14. Summary	65
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LIST OF ABBREVIATIONS

ACE	Angiotensin-Converting Enzyme
ACE-2	Angiotensin Converting Enzyme-2
AEC	alveolar epithelial cell
AGT	Angiotensinogen
Ang 1-7	Angiotensin 1-7
Ang II	Angiotensin II
Ang III	Angiotensin III
Ang IV	Angiotensin IV
AT ₁	Type 1 Angiotensin receptor
AT ₂	Type 2 Angiotensin receptor
ATI	Type I alveolar epithelial cell
ATII	Type II alveolar epithelial cell
ECM	extracellular matrix
EMT	epithelial-mesenchymal transition
HIF-1	Hypoxia Inducible Factor -1
IL	Interleukin
ILD	Interstitial Lung Disease
IPF	Idiopathic Pulmonary Fibrosis
JNK	c-Jun N-terminal kinase
MAPK	mitogen activated protein kinase
NFκB	nuclear factor-kappa B

PI3K	Phosphoinositide 3-kinase
RAS	Renin Angiotensin System
ROS	Reactive Oxygen Species
RT-PCR	Reverse Transcriptase-Polymerase Chain Reaction
siRNA	Small interfering RNA
SNP	Single Nucleotide Polymorphism
SP-C	surfactant protein C
TGF- β 1	Transforming Growth Factor-beta 1
TNF- α	tumor necrosis factor-alpha

Chapter 1

GENERAL INTRODUCTION

IDIOPATHIC PULMONARY FIBROSIS

Idiopathic pulmonary fibrosis (IPF) is a chronic and progressive fibrotic lung disease resulting from injury to the lung and an ensuing fibrotic response that leads to the thickening of the alveolar walls and the obliteration of the alveolar space (Fonseca et al., 2000). The American thoracic Society and the European Respiratory Society recently issued a consensus statement that defined IPF as a distinctive type of chronic fibrosing interstitial pneumonia of unknown etiology limited to the lung and associated with the histology of unusual interstitial pneumonia (American Thoracic Society, European Respiratory Society, 2002). IPF is classified as a specific presentation of idiopathic interstitial pneumonia, which is in turn a type of interstitial lung disease (American Thoracic Society, European Respiratory Society, 2002).

Epidemiology

IPF is the most common of the idiopathic interstitial pneumonias and is associated with the worst prognosis (American Thoracic Society, 2002). Deaths from IPF are estimated to be nearly 14 times greater than those from asbestos (Wells and Mannino, 1996). Differences in disease definition, classifications, diagnostic approach, and epidemiological design of the studies greatly influence the findings. In the United States, a recent study estimated prevalence to range from 14.0 to 42.7 cases per 100,000 population, depending on the criteria used for diagnosis (Raghu et al., 2006). This is higher than the previously reported prevalence of 3 to 6 per 100,000 persons (Cherniack et al., 1991). The median survival rate of IPF is estimated to be from 3 to 5 years (Selman et al., 2001).

Respiratory failure is the most frequent cause of death, and has been reported to account for over 80% of all fatalities (Martinet et al., 2005).

Two thirds of IPF patients are older than 60 years of age with a mean age of 66 at diagnosis (Johnston et al., 1997). The incidence, prevalence, and death rate of IPF increase with age (Coultais et al., 1994). IPF is reported to be more common in males than in females (Hubbard 1996). Some studies also suggest that the risk of developing IPF increases with environmental or occupational exposure to dusts, organic solvents, or urban pollution (Iwai, 1994). Other risk factors associated with pulmonary fibrosis include smoking, gastro-esophageal reflux disease, commonly prescribed drugs, diabetes mellitus, infectious agents, and genetic factors (Zisman et al., 2005).

Clinical Features

The clinical features for IPF are not specific and can also occur in various other pulmonary disorders. The symptoms usually vary with the extent of lung damage, the rate of disease progression, and the development of complications, such as lung infections or cor pulmonale. IPF usually presents with inspiratory crackles, dyspnoea on exertion, and dry, non-productive cough. Common symptoms also include weight loss and fatigue. In more than 80% of patients, bi-basilar, end-expiratory rales are found upon physical examination (American Thoracic Society, 2000). In addition, up to half of all patients develop digital clubbing where ends of the fingers become thick or club-shaped (Johnston et al., 1997). At late stages of the disease, patients also experience shortness of breath at rest as well as cyanosis of the lips and fingers. Pulmonary hypertension may also present late in the course (Panos et al., 1990).

There are currently no laboratory tests specific for the diagnosis of IPF. Laboratory evaluations are done to rule out alternative causes of interstitial lung disease such as sarcoidosis or connective tissue disease-pulmonary fibrosis. Pulmonary function tests show restrictive impairment, reduced diffusing capacity for carbon monoxide, and arterial hypoxemia exaggerated or elicited by exercise (Selman et al., 2001). Greater than 90% of patients have abnormal chest radiographs that show diffuse bilateral interstitial or reticulonodular infiltrates in the basilar and subpleural regions of the lung (Kazerooni et al., 1997; Wells et al., 1993). High-resolution computerized tomography scanning shows variable but limited ground-glass opacity subpleural honeycombing. Extensive honeycombing, septal thickening, and a lack of ground glass opacities are usually indications of a poor prognosis.

The diagnosis of idiopathic pulmonary fibrosis must include: 1) a compatible clinical history; 2) the exclusion of other known causes of interstitial lung disease (such as drug injuries, environmental exposures, or collagen vascular disease); 3) a surgical lung biopsy showing usual interstitial pneumonia histologic pattern (American Thoracic Society, 2000).

Histological Features

The histological features of IPF are described as usually interstitial pneumonia (UIP), which can be seen in other diseases as well. The histological hallmark of usual interstitial pneumonia is heterogeneous appearance of normal-appearing lung alternating with areas of peripheral fibrosis, interstitial inflammation, and honeycomb changes, on low magnification (American Thoracic Society, 2000). The inflammatory component is

typically light with mostly lymphocytes and plasma cells. At higher-power magnification, the border between the fibrotic and normal lung, called the fibroblastic foci, can be seen with accumulations of dense, relatively acellular, collagen bundles. Fibroblastic foci are the regions of fibroblastic proliferation and represent the primary site of ongoing injury. Fibroblastic foci are rich in extracellular matrix and mesenchymal cells with cell phenotypes ranging from proliferating fibroblasts to fully differentiated smooth muscle cells. The most abundant cell type is the myofibroblast, which contribute to active contraction, distorted architecture and excess collagen deposition (Kuhn & McDonald 1991). Although not pathognomonic, the number of fibroblastic foci may be an important prognostic factor and correlate with a worse prognosis (Nicholson et al., 2002, King et al., 2001).

Genetic factors

Evidence suggests that genetic factors may play an important role in the pathogenesis of IPF. It has been estimated that up to 4% of IPF is familial and appears to be inherited as an autosomal dominant trait (Lawson & Lloyd, 2006). The familial form of IPF has been reported in monozygotic twins raised in different environments, in genetically related members of several families, in consecutive generations in the same families, and in family members separated at an early age (Steele et al., 2005). Genetic studies found two mutations of the SP-C gene to be associated with familial lung fibrosis (Thomas et al., 2002). These mutations lead to misfolding of the protein resulting in abnormal surfactant production and injury in type II alveolar epithelial cells (Thomas et al., 2002). Angiotensin II is a growth factor that plays a key role in the physiopathology of IPF. A nucleotide

substitution of an adenine instead of a guanine (G-6A) in the proximal promoter region of the angiotensin II precursor, angiotensinogen, has been associated with an increased gene transcription rate. A recent case-control study found that the distribution of G-6A genotypes and alleles did not significantly differ between cases and controls (Molina et al., 2008). The G-6A polymorphism of the angiotensinogen gene was found to be associated with idiopathic pulmonary fibrosis progression but not with disease predisposition (Molina et al., 2008). The G-6A polymorphism of the angiotensinogen gene could have a predictive significance in idiopathic pulmonary fibrosis patients.

Therapeutic options

The two main types of therapeutic agents used to treat IPF are the anti-inflammatory medications and the anti-fibrotic agents. Most therapies are, however, largely ineffective with limited evidence to suggest that any one treatment improves survival or disease progression. The conventional management of IPF with anti-inflammatory medications is based on the concept that pulmonary fibrosis is an inflammatory disorder and the suppression of inflammation prevents progression to fibrosis. These therapies continue to be used despite the lack of evidence of inflammation in the pathogenesis of IPF (Raghu et al., 2004). Clinical studies indicate that response to steroids is poor in IPF and the use of other immunosuppressive or aggressive cytotoxic agents only offer a marginal benefit at best (King et al., 2001). Recently, more anti-fibrotic agents are being used in the treatment of IPF based on the concept that the disease is a fibrotic condition with a lack of significant inflammatory component (Selman et al., 2004). A meta-analysis of the combined results of multiple studies showed that interferon- γ 1b (IFN- γ 1b) therapy may be efficacious in

improving survival and is associated with reduced mortality in patients with IPF (Bajwa et al., 2005). However, this benefit of IFN- γ 1b therapy was not seen in patients with more severe disease (Martinez et al., 2005).

Currently, the only option for patients with end stage pulmonary fibrosis and those who fail to respond to medical treatment is lung transplantation. Patients are referred for transplantation when progressive symptoms have a vital capacity less than 60-70% and a corrected diffusion capacity below 50-60% of predicted (The American Society for transplant physicians, 1998). The five-year survival rate for patients following transplant is estimate to be 50-60% (American Thoracic Society, 2000). A recent study estimated that lung transplantation reduces the risk of death by 75% (Thabut et al., 2003).

To develop new therapies, recent studies have focused on targeting angiotensin II (ANG II) and its role in the pathogenesis of pulmonary fibrosis. These studies showed that inhibition of the RAS using angiotensin converting enzyme inhibitors (ACEi's), angiotensin receptor blockers or receptor deletion was able to prevent experimental lung fibrosis (Wang et al., 2000b, Marshall et al., 2004, Uhal et al., 2007a, Molteni et al., 2000, Molina-Molina et al., 2006) and improve lung function in IPF patients (Woo et al., 2003).

PATHOGENESIS OF FIBROGENESIS

Until recently, pulmonary fibrosis has been considered to be an inflammatory disorder where lower respiratory tract inflammation leads to derangements of the alveoli resulting in scarring of the lung parenchyma (Gallin et al., 1992). Inflammation results in the loss of functional alveolar-capillary unit, the accumulation of collagen, and the formation of the honeycomb lung. Recent studies, along with the limited success of anti-inflammatory treatments, suggest the existence of alternative hypothesis regarding the pathogenesis of IPF. The current hypothesis is that IPF results from alveolar epithelial injury and abnormal wound repair (Selman et al., 2001). This theory is supported by evidence from human lung biopsies where nascent fibrotic foci were colocalized with unrepaired or abnormal epithelia (Uhal et al., 1998).

Alveolar Epithelial Cells

The alveolus is the basic functional unit of the lung where gas exchange occurs between the air and the capillaries. The site of gas exchange is composed of the capillary endothelium and alveolar epithelium separated by a thin interstitium composed of connective tissue, ECM and a few fibroblasts (Fonseca et al., 2000). The alveolar epithelium is composed of two types of alveolar epithelial cells (AECs) that are morphologically and functionally distinct. The first is the thin and elongated type I (AT I) cells that are terminally differentiated and cover 95% of the alveolar surface to facilitate gas exchange. AT I cells are metabolically active and express cell surface receptors for a variety of substances, including extracellular matrix (ECM) proteins, growth factors, and cytokines. The second are type II (AT II) alveolar epithelial that reside in alveolar corners

and are cuboidal in shape with rounded nuclei. AT II cells secrete surfactant, facilitate trans-epithelial movement of water, function as antigen presenting cells and serve as the progenitor of both types of epithelial cells (Uhal, 1997). AT II cells are important in the maintenance and repair of the alveolar epithelium by replacing lost cells and restoring normal tissue architecture and lung function. The regeneration of a continuous epithelium is vital in maintaining barrier function and in limiting airway hyper-activity. This process is most critical in response to lung injury and other insults. Epithelial injury and blunted epithelial repair are sufficient to promote pulmonary fibrotic processes (Adamson et al., 1988).

One important step in the healing process is the rapid re-epithelialization of the denuded area through epithelial cell migration, proliferation and differentiation. The inhibition of alveolar epithelial repair in animal models increases the severity of subsequent fibrosis (Haschek and Witschi, 1979), suggesting that the efficient repopulation of the denuded alveolar basement membrane acts to suppress subsequent fibroblast proliferation and extracellular matrix accumulation. IPF alveolar epithelium show significant loss of AT I cells, hyperplasia of AT II cells and altered expression of adhesion molecules and MHC antigens (Kasper, 1996). AT II cells in IPF have decreased capacity to restore damaged AT I cells, resulting in epithelial cuboidalization and the presence of transitional reactive phenotypes (Kasper, 1996), abnormalities in pulmonary surfactant (McCormack et al., 1991), and alveolar collapse (Burkhardt, 1989).

The epithelial cells overlying fibroblastic foci are found to be hyper- and dysplastic, with abnormal morphology and gene expression patterns (Kasper et al., 1996). These cells secrete a variety of profibrotic cytokines and participate in a bidirectional communication

network with neighboring fibroblasts where each cell type influences the proliferation/survival of the other. AEC apoptosis was also detected adjacent to myofibroblasts-containing fibroblastic foci, the presumed primary sites of epithelial injury in IPF. Ongoing apoptosis is believed to be a key component in the progression of IPF and appears to be essential for the development of transforming growth factor-beta (TGF- β) induced lung fibrosis (Lee et al., 2004; Uhal et al., 1998).

Coagulation and fibrinolysis

The disruption of blood vessels and extravasation of blood into the wound after tissue injury reestablishes hemostasis while also providing a provisional extracellular matrix in which the repair process can begin. AT II cells and macrophages convert the leaked in fibrinogen into fibrin, which then becomes scaffold for fibroblast migration (Gross et al., 1991). The efficient and orderly resolution of the fibrin matrix will reconstitute a normal alveolar space. Continued deposition of fibroblasts and dysregulated coagulation, however, will lead to destruction of the alveolar space and contribute to pulmonary fibrosis. Whether fibrin is deposited or resorbed in the alveolar space is critical in the development of fibrosis and is determined by the balance of procoagulatory, fibrinolytic, and antifibrinolytic activities in the lungs. Fibrin removal is slowed in IPF lungs as evidenced by increased procoagulant and anti-fibrinolytic activities found in bronchoalveolar lavage fluid from IPF patients (Gunter et al., 2000).

During wound repair, the fibrinolytic process is required to cleave a path for epithelial cell migration. The plasminogen activation system mediates this process and is crucial in tissue remodeling. Plasminogen, when activated to plasmin by tissue-type

plasminogen activator (tPA) or urokinase-type plasminogen activator (uPA), is the primary fibrinolytic enzyme responsible for degrading fibrin clots and promoting wound reepithelialization. The overexpression of uPA has been shown to decrease fibrosis in a murine model, indicating a protective role of uPA in the development of pulmonary fibrosis (Sisson et al., 2002). Plasminogen activator inhibitors (PAIs) negatively regulate plasminogen. Animal models have shown that the overexpression of PAIs inhibits plasmin activity and promotes the formation of fibrosis while the lack of PAIs increase plasmin activity and prevents fibrosis (Eitzman et al., 1996). Alveolar epithelial cells synthesize uPA and its receptor uPAR as well as PAI-1 (Hasegawa et al., 1997; Simon et al., 1992). PAI-1 is strongly induced by TGF- β and may play a role in TGF- β induced fibrosis.

In normal repair, epithelial cells are able to dissolve the fibrin barrier and migrate throughout the denuded wound surface. The proximity of uPA and its receptor greatly increases the rate of plasminogen activation on the alveolar epithelium and the clearance of fibrin from the alveolar spaces (Hattori et al., 1989). Alveolar epithelial cells in the IPF lung, however, seem to play a role in increasing procoagulant and antifibrinolytic activities. BAL fluid from IPF patients was found to contain significantly more PAIs but the same amount of uPA as compared to normal individuals (Kotani et al., 1995). This suggests that the IPF alveolar microenvironment prefers a pro-coagulant and anti-fibrinolytic state that promotes ECM accumulation and inhibits alveolar reepithelialization.

Profibrotic Cytokines

Alveolar epithelial cells in IPF also express several cytokines and growth factors that promote fibroblast migration and proliferation as well as extracellular matrix

accumulation. Damaged AECs produce platelet-derived growth factor (PDGF) (Antoniades et al., 1990), tumor necrosis factor alpha (TNF- α) (Kapanci et al., 1995), endothelin-1 (Giaid et al., 1993), and are the primary producers of TGF- β (Khalil et al., 1996). TGF- β has profound effects on epithelial cells and fibroblasts and is a central regulator of pulmonary fibrosis. TGF- β promotes epithelial cell apoptosis (Hagimoto et al., 2002), epithelial-to-mesenchymal transition (EMT) (Kim et al., 2006), epithelial cell migration (Yu et al., 2008), collagen synthesis, fibroblast proliferation and transformation into myofibroblasts (Scotton and Chambers, 2007). Studies with animal models found that overexpression of TGF- β leads to extensive and progressive fibrosis with limited inflammation (Sime et al., 1997). PDGF released by epithelial cells is a potent mitogen and chemoattractant for fibroblasts. PDGF mRNA and protein were found to be upregulated in the epithelial cells of IPF patients (Antoniades et al., 1990). Hyperplastic AT II cells in pulmonary fibrosis synthesize TNF- α , which promotes DNA synthesis and proliferation of fibroblasts (Battegay et al., 1995). Endothelin-1 also plays a role in stimulating fibroblast proliferation and transdifferentiation into myofibroblasts (Shahar et al., 1999). Together, these cytokines released by AECs after injury promote proliferation, migration, and matrix deposition by fibroblasts.

Myofibroblasts

Fibroblasts are highly versatile and are able to interconvert between several distinct but related cell types. This phenotypic plasticity of fibroblasts is crucial to tissue repair after injury. After lung injury, fibroblasts are responsible for secreting ECM proteins to provide a tissue scaffold for epithelial cell migration during repair. Additionally, the dissolution of this scaffold along with apoptosis of fibroblasts and myofibroblasts are

critical for restoration of normal tissue structure (Desmouliere et al., 1995; Lorena et al, 2002).

In IPF, AECs lose their ability to provide normal and homeostatic fibroblast-suppressive functions and instead initiate the pathologic process by producing most of the factors that induce the fibroblast phenotypic changes. In the fibroblastic foci of the injured lung, fibroblasts continually modify their interactions with the microenvironment. They first become migratory, then proliferative, and finally become profibrotic and secrete abundant ECM (Kuhn et al., 1989). Activation of fibroblasts by the secreted factors produce myofibroblasts that express contractile proteins and have ultrastructural features that are intermediate between fibroblasts and smooth muscle cells (Gabbiani et al., 1971). The contractile function of myofibroblasts is important in the re-epithelialization process by bringing wound margins closer together. Myofibroblasts also have high synthetic capacity for ECM proteins (Ignatz and Massague, 1986), growth factors/cytokine (Finlay et al., 2000), growth factor receptors (Thannickal et al., 1998), integrins (Heino et al. 1989), and oxidants (Thannickal et al 1989).

In addition to sending activation signals to mesenchymal cells, AECs themselves can also undergo phenotypic transition to fully differentiated mesenchymal cells such as fibroblasts and myofibroblasts (Zavadil et al., 2005). This process is called epithelial to mesenchymal transition (EMT). Epithelial cells undergoing EMT are able to lose polarity, disassemble cell adhesion systems, produce cell-motility machinery, and move from one location to another (Zavadil et al., 2005). EMT plays a pivotal role in cellular transdifferentiation during embryonic morphogenesis and tumor progression. In the adult, injury to epithelial cells induces EMT and contributes to fibrosis in a number of organs

(Iwano et al., 2002; Saika et al., 2004). EMT has been investigated as a mechanism underlying fibrosis in renal and lens epithelium. Injured renal tubular epithelial cells during kidney fibrosis migrate through damaged basement membranes into the interstitium and fully transdifferentiate into fibroblasts and myofibroblasts (Fan et al., 1999; Kalluri et al., 2003). About 36% of new fibroblasts come from EMT of local epithelium in kidney fibrosis (Iwano et al., 2002).

The stimulatory input of epidermal growth factor (EGF), hepatocyte growth factor (HGF), fibroblast growth factors (FGF), and especially TGF- β factors stimulate epithelial cells to lose polarity, express basement membrane-degrading matrix metalloproteinases (MMPs), undergo cytoskeletal rearrangements, and express machinery necessary for motility, leading to migration and complete transition to a mesenchymal phenotype (Grunert et al., 2003; Kalluri et al., 2003; Savagner et al., 2001). It was recently demonstrated *in vitro* that TGF- β 1 induces EMT in AECs (Willis et al., 2005).

In addition to the highly synthetic phenotype of fibroblasts in IPF, there is also an imbalance in the production of MMPs and tissue inhibitors of metalloproteinases (TIMPs) that control MMP activity (Ramos et al., 2001; Selman et al., 2000). MMPs are a family of highly regulated zinc-dependent peptidases and have been implicated in the remodeling of ECM and cell migration (Nagase and Woessner, 1999). TIMP-2 is expressed by myofibroblasts within the fibrotic foci and may induce mesenchymal cell proliferation. This in part explains the survival of mesenchymal cells in the fibroblast foci in IPF, in contrast to the expected cell death as observed in normal wound healing (Desmouliere et al., 1995). In addition, TIMP-2 has been found to contribute to the irreversible structural remodeling in IPF (Fukuda et al., 1998).

Myofibroblasts have also been shown to secrete angiotensin peptides that may induce apoptosis of adjacent alveolar epithelial cells (Uhal et al., 1998). The signaling crosstalk between AECs and fibroblasts is critical in determining whether fibrosis occurs after lung injury. The various roles and phenotypes of fibroblasts and myofibroblasts make them key effector cells in the pathogenesis of fibrosis.

Basement Membrane Disruption and ECM Remodeling

The basement membrane plays an important role in maintaining the integrity and differentiation of the alveolar epithelium. The basement membrane is a complex structure composed of type IV collagen, laminin, entactin, fibronectin, and heparin sulfate-chondroitin proteoglycans (Yurchenco and Schittny, 1990). The disruption of the basement membrane contributes to the pathogenesis of fibrosis. MMP proteins produced by the myofibroblasts in IPF have been shown to degrade different components of the basement membrane, type IV collagen in particular (Segura et al., 2000). In addition, MMP proteins in IPF have been found to coincide in areas with denuded alveolar basement membrane (Hayashi et al., 1996; Selman et al 2000). These findings suggest that myofibroblasts play a role in the degradation of the basement membrane to facilitate their migration into the alveolar spaces. This disrupted basement membrane may also contribute to the failure of an orderly repair of damaged AT I cells.

Fibroblasts and myofibroblasts also play a central role in the synthesis, deposition, and remodeling of the ECM. Increase in the deposition of ECM, including fibrillar collagens, fibronectin, elastic fibers, and proteoglycans, results in the aberrant tissue remodeling observed in IPF (Selman et al., 1996). This remodeling eventually leads to

extensive structural disorganization in the lung microenvironments and progressive loss of the alveolar-capillary units.

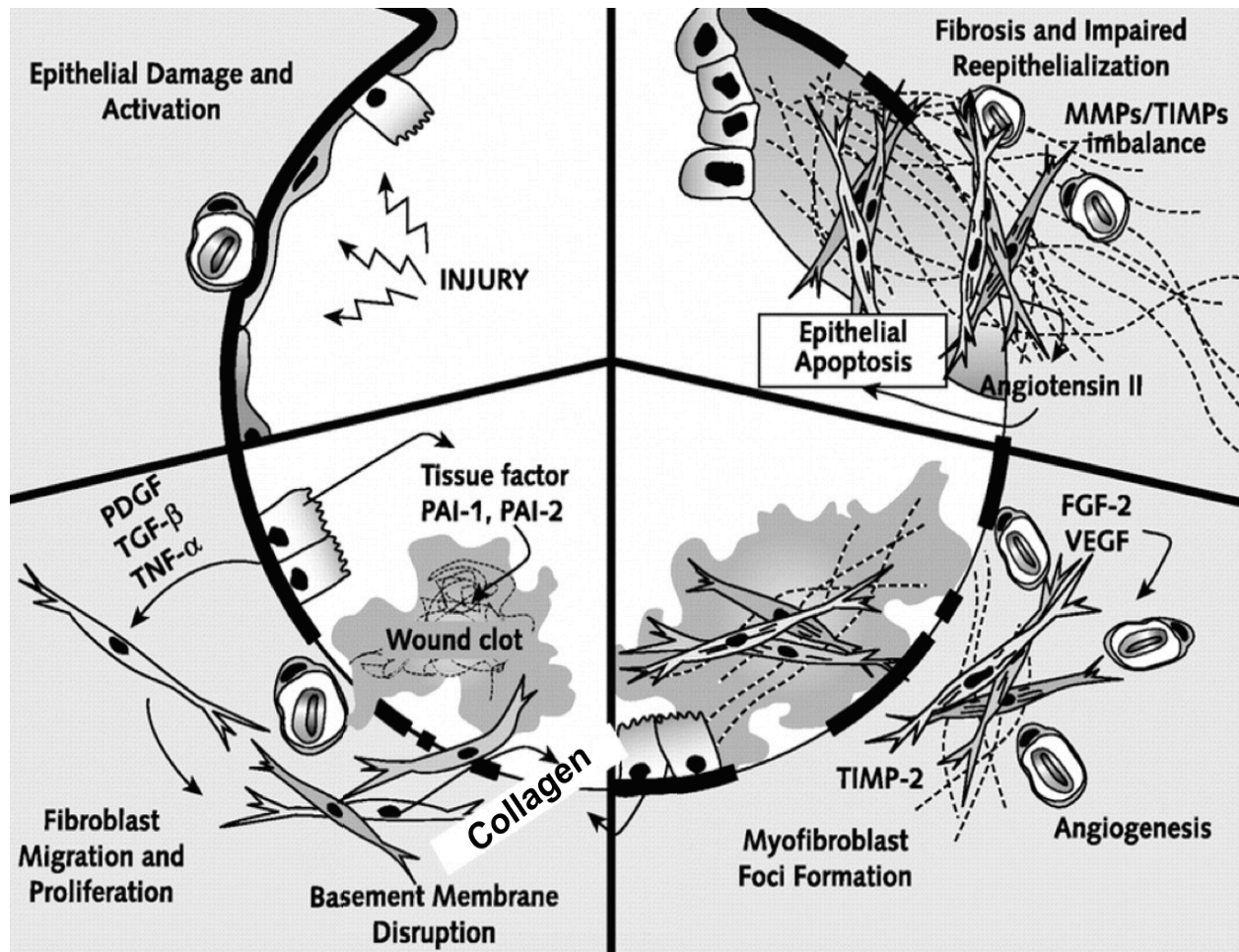


Figure 1.1.

Pathogenesis of idiopathic pulmonary fibrosis [Figure 7.1 modified from (Selman et al., 2001)]. Alveolar epithelial cell damage (top left) induces an antifibrinolytic environment in the alveolar spaces, which then enhances wound clot formation. Alveolar epithelial cells also secrete growth factors that induce migration and proliferation of fibroblasts and differentiation into myofibroblasts (bottom left). Myofibroblasts secrete extracellular matrix proteins, mainly collagens. Neovascularization is induced by angiogenic factors TIMP-2, FGF-2, and VEGF (bottom right). An imbalance between

interstitial collagenases and tissue inhibitors of matrix metalloproteinases (MMPs) provokes the progressive deposit of extracellular matrix (top right). Myofibroblasts also produce angiotensin II, which provokes alveolar epithelial cell death, further impairing re-epithelialisation. PDGF: platelet-derived growth factor; TGF: transforming growth factor; TNF: tumour necrosis factor; PAI: plasminogen activator inhibitor; TIMP: tissue inhibitors of metalloproteinases; VEGF: vascular endothelial growth factor; FGF: fibroblast growth factor.

APOPTOSIS IN IDIOPATHIC PULMONARY FIBROSIS

Apoptosis is programmed cell death that is a critical physiological process in the development and maintenance of tissue homeostasis. Apoptosis is important in almost all cell types to provide a balance between cellular proliferation and turnover. In IPF, apoptosis affects the development of fibrosis through both increased apoptosis of epithelial cells resulting in inefficient reepithelialization and resistance to apoptosis of fibroblasts resulting in increased fibrosis (Antoniou et al., 2007).

Apoptosis in General

Morphological and Biochemical Changes

The cell undergoes a highly regulated series of characteristic morphological and biochemical changes during apoptosis that distinguish it from normal and necrotic cells. The first morphological changes during apoptosis are cytoplasmic shrinking, loss of cell-cell contacts and active membrane blebbing (Allen et al., 1997). In addition, the cytoplasm and nuclear chromatin condense and fragment into membrane-bound vesicles. Phosphatidylserine (PS) externalizes at the intact cell surface and promotes signal recognition by phagocytic cells. Endogenous DNase degrades chromosomal DNA and cleaves the internucleosomal regions into double stranded DNA fragments. Another marker of apoptosis is the activation of cytosolic caspases. Caspases are cysteine proteases that cleave and degrade a specific subset of cellular proteins in cells undergoing apoptosis. The entire process of apoptosis occurs in a predictable sequence that concludes with the engulfment of apoptotic bodies by other cells.

Caspases

Caspases are a family of aspartate-specific cysteine proteases that are present in all animal cells as inactive zymogens. In apoptosis, caspases function as both effectors in cell disassembly and as initiators in initiating this disassembly in response to proapoptotic signals. Initiator caspases are the first to be activated in a death pathway. They then activate effector caspases through a caspase cascade. Proteolytic processing at conserved aspartic acid (Asp) residues triggers caspases to assume active states. During activation, the zymogen pro-proteins are cleaved at a site known as the linker region to generate the large and small subunits of the active enzymes. One small and one large subunit make up one catalytic unit of the active caspase enzyme. Each catalytic unit contains one active site and a dimer of identical catalytic units make up one active caspase enzyme. The N-terminal prodomain regulates activation of the proenzyme and is highly variable in sequence and length. Caspases are unusually highly specific and recognize the Asp residues they cleave within tetrapeptide motifs (Nicholson and Thornberry, 1997). This strict specificity ensures that caspases are signaling proteases intended for specific protein cleavage and not for protein degradation.

Caspase proteins collaborate in proteolytic cascades where caspases activate themselves and each other. Caspase enzymes cleave their substrates at Asp residues, and are also activated by proteolytic processing at Asp residues. Activated upstream initiator caspases then activate downstream effector caspases. The N-terminal domains of proform initiator caspases are large and function as protein interaction modules that allow them to interact with various proteins that trigger caspase activation. The proforms of downstream

effector caspases contain only short N-terminal prodomains that appear to serve no function. The effector caspases depend mostly on initiator caspases for their proteolytic processing and activation. Consequently, the sequence of the cleavage sites of zymogen effector caspases generally matches the preferred tetrapeptide specificities of the initiator caspases. Additionally, cleavage sites of proteins that were identified as caspase substrates that undergo processing during apoptosis also match with the preferred specificities of the effector caspases (Thornberry et al., 1997). Most caspases in mammals and higher eukaryotes are directly involved in cell death except for a few, such as caspase-1, 4, and 5 that are involved in processing of pro-inflammatory cytokines in humans. The tetrapeptide specificities of these cytokine-processing proteases do not match the cleavage sites of most of the proteins known to undergo cleavage during apoptosis, but instead they match with the sequences of the cleavage sites within pro-cytokines (Salvesen and Dixit, 1997).

Initiator caspases can be activated by a variety of mechanisms to initiate the apoptotic cascade. The underlying biochemical mechanisms of initiator caspase activation are all remarkably similar and can be explained by the induced proximity model (Salvesen and Dixit, 1999). The zymogen forms of unprocessed caspases are not completely inactive but rather possess weak protease activities that, in some cases are at approximately 1% of the level of the fully active enzymes. Protein interactions can bring the zymogens close together and allow them to trans-process each other to produce fully active proteases.

Mechanisms of Apoptosis

Apoptosis can be induced by various physiological stimuli such as deformation of the cell membrane, growth factor withdrawal, inflammatory cytokines, osmotic shock or

UV irradiation, as well as a number of molecules and agents. When stimulated, the cells undergo the activation of pro-apoptotic signaling pathways that transduce a signal to the interior of the cell and activate the apoptotic machinery.

The mechanism of apoptosis begins with the initiation phase, during which various apoptotic stimuli lead to activation of caspases, followed by the execution phase where the activated caspases induce cell death. The caspase cascade can be activated by many pathways, two of which have been studied in detail. Both of these pathways lead to the activation of the signaling cascade that results in the cleavage of inactive pro-caspase molecules into active caspase proteins.

The Extrinsic Pathway/Receptor-mediated Pathway

The first is the extrinsic or death receptor pathway that involves the activation of death receptors present in the cell membrane, such as Fas and tumor necrosis factor (TNF) receptors I and II (Figure 1.2). Death receptors are protein interaction modules that consist of a compact bundle of six α -helices (Huang et al., 1996). Death receptors bind to each other with specificity directed by differences in their surface residues. Fas receptor can be activated by either the Fas ligand (FasL) on the surface of cytotoxic lymphocytes, or by a soluble form of FasL (sFasL) that can be cleaved from cell membranes by MMP-7 and MMP-3 (Matsuno et al., 2001; Powell et. al 1999). Upon ligation to FasL, the Fas receptor forms microaggregates at the cell surface. This allows the adaptor molecule, Fas-associated protein with death domain (FADD), to be recruited to the cytosolic tail of the Fas receptor. FADD contains a death effector domain (DED) at its amino-terminus and a Death Domain (DD) at its carboxy-terminus. Caspase-8 contains two DEDs at its amino-terminus. FADD

binds Fas receptor and at least three molecules of procaspase-8 through homotypic interactions. Within this death inducing signaling complex (DISC), procaspase-8 is able to undergo activation by autocatalytic cleavage via the induced proximity mechanism. During activation, the N-terminal prodomain of procaspase-8 is cleaved off and the active protease is released into the cytosol. The active caspase-8 then cleaves and activates downstream caspases-3, 6, and 7 as well as the pro-death protein Bid. The Bid cleavage fragment then targets the mitochondria and leads to a mitochondrial amplification of the caspase pathway.

The Intrinsic Pathway/Mitochondrial Pathway

The intrinsic pathway responds to ionizing radiations, chemotherapeutic drugs, mitochondrial damage, and environmental cues. Death triggers increase mitochondrial permeability, leading to the release of cytochrome *c* and the subsequent recruitment and activation of caspase-9 (Zou et al., 1999). The release of cytochrome *c* triggers the formation of the apoptosome complex. The apoptotic protease-activating factor 1 (Apaf1) is the main component of the apoptosome. Apf-1 is a multidomain protein that consists of three functional regions: an N-terminal caspase-recruitment domain (CARD); a nucleotide-binding and oligomerization domain (NOD); and a string of WD40 repeats in the C-terminal of the protein. In the absence of an apoptotic signal, Apaf-1 exists in a monomeric form. Cytochrome *c* binding allows the compact Apaf-1 molecule to stretch out and polymerize upon binding ATP (Acehan et al., 2002). Studies of the apoptosome through electron cryomicroscopy reveal it as a seven-spoked wheel with the caspase-9 recruitment domain, CARD of Apaf-1, in the central hub (Renatus et al., 2001). Caspase-9 is activated through dimerization of monomers within the apoptosome. In contrast to the activation of

caspase-8, the N-terminal prodomains of caspase-9 is not cleaved and the caspase-9 enzyme must remain bound to Apaf-1 to be fully active (Rodriguez et al., 1999). Activated caspase-9 is then released from the complex to cleave and activate downstream caspases-3, 6, and 7 (Budihardjo et al., 1999). Members of the B-cell lymphoma (Bcl)-2 family proteins, including the anti-apoptotic Bcl-2/Bcl-xL and the pro-apoptotic Bax/Bid, regulate the release of cytochrome *c* from the mitochondria and act to suppress or promote apoptosis. Activated Bax/Bid migrate from the cytosol to the mitochondria to initiate release of cytochrome *c* from the mitochondria and promote apoptosis. Bcl-2/Bcl-xL are primarily localized in the outer member of the mitochondrial and are able to inhibit the release of cytochrome *c* to suppress apoptosis. The balance of anti-apoptotic proteins and pro-apoptotic proteins determines the release of cytochrome *c* and activation of downstream caspases and therefore the cell's susceptibility to apoptotic stimuli. In addition, both Bcl-2/Bcl-xL and Bax/Bid are transcriptional targets of the tumor suppressor protein p53, which induces cell cycle arrest and apoptosis in response to DNA damage (Earnshaw et al., 1999)

Caspase Substrates

Activation of caspases leads to downstream cleavage of key structural components of the cytoskeleton and nucleus as well as proteins involved in the apoptosis signaling pathway. Pro-and anti-apoptotic regulatory proteins are among the first to be cleaved by activated caspases. Caspase-3 cleavage of Bcl-2/Bcl-xL inactivates the anti-apoptotic properties of these proteins by producing a pro-apoptotic product that increases mitochondrial permeability and release of cytochrome *c* (Li et al., 1998). Caspase-8 can

also induce cytochrome *c* release by cleaving pro-apoptotic Bid (Lou et.al, 1994). These cleavage events allow the extrinsic pathway to communicate with and utilize the mitochondrial pathway to amplify apoptotic signals.

Other Pathways

The MAPK pathway allows signals to be transmitted from the cell membrane to the nucleus in response to various stimuli. The MAPK pathway also induces the phosphorylation of intracellular substrates such as protein kinases and transcription factors to regulate cellular functions such as cell growth, differentiation, and apoptosis (Kyriakis and Avruch, 1996). The MAPK family of kinases include: the extracellular signal-regulated kinases (ERKs), ERK-1 (p44 MAPK) and ERK-2 (p42 MAPK); stress-activated protein kinases (SAPKs), also referred to as c-Jun, NH₂-terminal kinases (JNKs), which include p54 SAPK (SAPK α /, β , JNK2) and p45 SAPK (SAPK γ , JNK1); and the p38 kinases (α , β , γ , and δ) (Widmann et al., 1999). Activation of the SAPK/JNKs and p38 kinases play a role in apoptosis and activation of the ERK1/ERK2 pathways are involved in cell survival (Earnshaw et al., 1999).

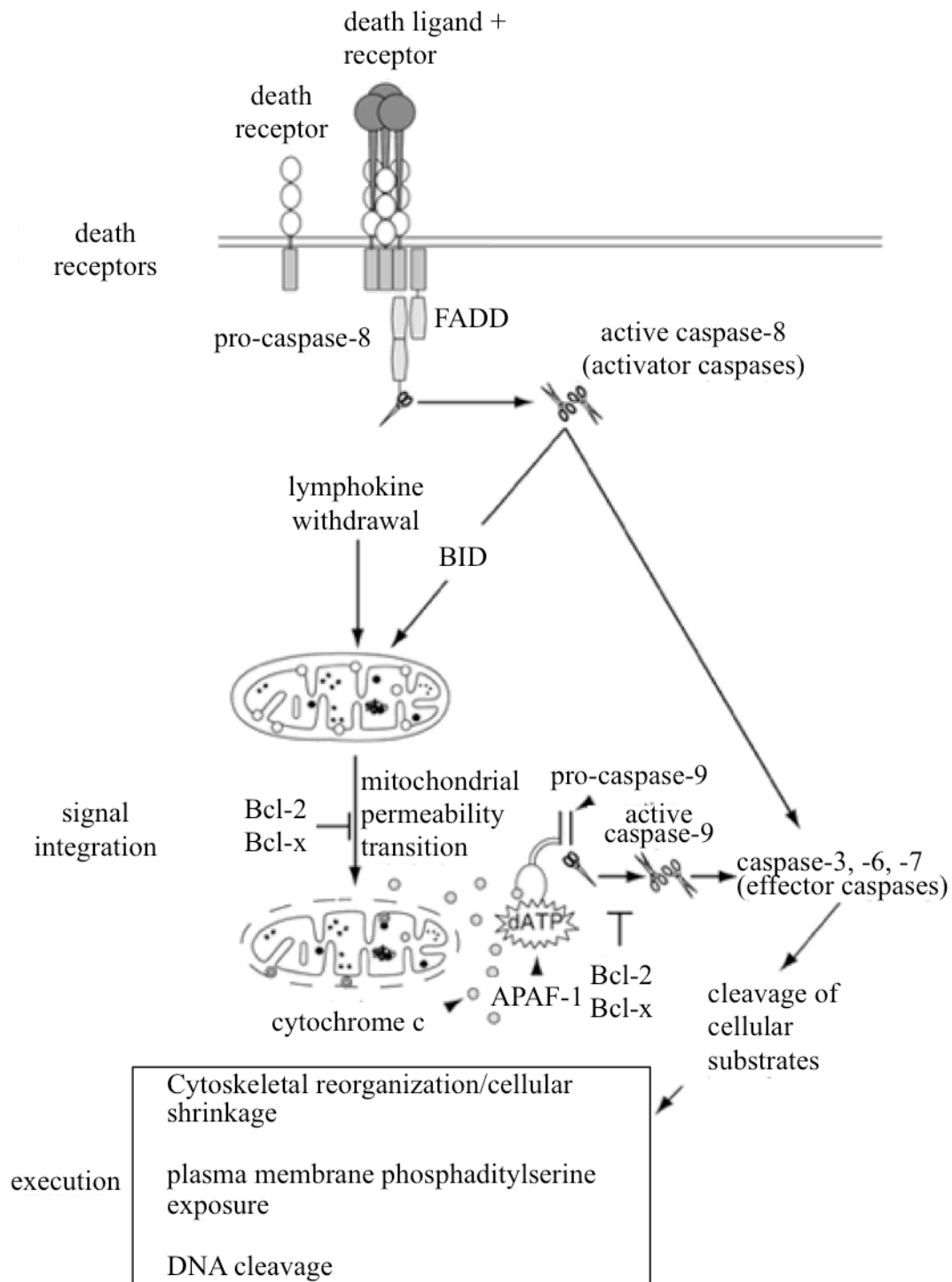


Figure 1.2.

Induction of apoptosis [Figure 11.9C.1 modified from (Siegel and Lenardo, 2002)]

Alveolar Epithelial Cell Apoptosis in IPF

Alveolar epithelial cell injury and death have consistently been found in human IPF. The survival and recovery of epithelial cells is key to normal repair of damaged epithelia. Fibrosis occurs through the pathological repair process of tissue remodeling. Often, the degree of remodeling is closely associated with the patient's prognosis. Hence, a greater understanding of epithelial cell apoptosis may lead to the development of effective strategies for treatment.

Evidence of Apoptosis

Alveolar epithelial cell injury is a characteristic feature of IPF. In the normal lung, ATI cells makeup 40% of the AEC population and cover 95% of the alveolar surface (Mason and Shannon, 1997). The number of AT1 cells are markedly decreased in the IPF lung after injury and cell death. Recently, increased apoptosis of ATII cells has been found in areas of IPF that do not have established apoptosis (Barbas-Filho et al., 2001), and in epithelial cells that overlay myofibroblasts (Uhal et al., 1998). Additionally, increased expression of proapoptotic proteins and decreased expression of anti-apoptotic proteins has also been found in the IPF lung (Plataki et al., 2005). Experiments using caspase inhibitors also demonstrate the involvement of apoptosis in fibrosis. The tripeptide broad-spectrum caspase inhibitor, benzyloxycarbonyl-Val-Ala-Asp fluoromethylketone (Z-VAD.fmk), inhibits the intracellular activation of caspase-like proteases *in vivo*, and attenuates bleomycin-induced pulmonary fibrosis in mice (Kuwano et al., 2001; Wang et al., 2000). Thus, there is growing evidence that subpopulations of AECs in IPF are undergoing

apoptosis in association with the dysfunctional repair of the damaged alveolar epithelium.

Induction of Pulmonary Fibrosis

Studies of animal models have provided further understanding of whether excessive AEC apoptosis is sufficient to lead to fibrogenesis. The induction of AEC apoptosis using aerosolized anti-Fas antibody was found to associate with the development of pulmonary fibrosis in mice (Hagimoto et al., 1997). Targeted transgenic overexpression of bioactive TGF- β 1 to the murine lung produces a transient wave of epithelial apoptosis that is followed by mononuclear-rich inflammation, tissue fibrosis, myofibroblast hyperplasia, and honeycombing (Lee et al., 2004). Furthermore, a null mutation of early growth response gene (EGF)-1 or caspase inhibition can rescue the fibrotic phenotype (Lee et al., 2004).

Mechanism of Alveolar Epithelial Apoptosis

p53 and p21 upregulation

Lung tissues from patients with IPF reveal upregulation of p53 and p21 in lung epithelial cells (Kuwano et al., 1996). In normal cells, p53 responds to DNA damage and mediates cell cycle arrest, DNA repair, and apoptosis. p53 expression is upregulated in response to a variety of stresses. ATII cell apoptosis is associated with upregulation of p53 and p21 expression in diffuse alveolar damage (Guinee et al., 1996). ATII cells of patients with IPF also express high levels of p53 in association with DNA strand breaks (Kuwano et al., 1996; Plataki et al., 2005;). p53 and p21 are also overexpressed in AECs treated with bleomycin (Mishra et al., 2000; Kuwano et al., 2000). p21 is a critical downstream effector in the p53 pathway of growth control in mammalian cells and is induced after exposure to

DNA-damaging agents (el-Deiry et al., 1993). p21 enhances cell survival by promoting DNA repair or by modifying cell death caused by exposure to hyperoxia (O'Reilly et al., 2001). Transfer of the p21 gene through adenovirus to epithelial cells attenuates bleomycin-induced pulmonary fibrosis in mice (Inoshima et al., 2004). p21 also regulates activation of caspase-3. The formation of a procaspase-3-p21 complex is essential for cell survival (Suzuki et al., 1998; Suzuki et al., 1999), suggesting that p21 may act as a key regulator of DNA replication and repair after lung damage.

Fas-FasL Pathway Activation

Fas is expressed on the luminal surface of a subset of AT II cells (Fine et al., 1997). In patients with acute lung injury, alveolar epithelial damage is in part associated with the local upregulation of the Fas-FasL pathway and subsequent activation of the apoptotic cascade in epithelial cells (Albertine et al., 2002). In lung tissues of IPF patients, FasL mRNA and protein expression are upregulated in infiltrating inflammatory cells and Fas protein is overexpressed in the epithelial cells (Kuwano et al., 1999). BAL fluid obtained from patients with IPF or ARDS could induce apoptosis on small airway epithelial cells through the Fas-FasL pathway (Matute-Bello et al., 1999; Hagimoto et al., 2002).

Bleomycin is used to induce pulmonary fibrosis in animal models of lung injury and fibrosis. FasL mRNA in this model is upregulated in infiltrating lymphocytes. In addition, excessive apoptosis is detected in the bronchiolar and alveolar epithelial cells where Fas protein expression is upregulated (Hagimoto et al., 1997). Neutralization of FasL prevents the development of fibrosis in this model (Kuwano et al., 1999). These findings indicate that the Fas-mediated apoptotic pathway is essential in the bleomycin-induced pulmonary

fibrosis model. Furthermore, the inhibition of caspases may provide a new approach to treating pulmonary fibrosis.

TGF- β Induction of Apoptosis

TGF- β is a strong chemotactic attractant for monocytes and macrophages as well as a potent promoter of ECM production. There is a consistent increase in TGF- β production in epithelial cells and macrophages in lung tissues of IPF patients (Khalil et al., 1991) and in bleomycin-induced pulmonary fibrosis in rodents (Raghow et al., 1989). Gene transfer of a TGF antagonist reduces the fibrotic response to bleomycin (Kolb et al., 2001). TGF- β 1 directly induces epithelial cell apoptosis via Fas and caspase-3 activation, and by enhancer the FasL-Fas interaction (Hagimoto et al., 2002). The Bax-mediated, Bid-activated pathway is involved in the pathogenesis of pulmonary fibrosis. TGF- β 1 significantly stimulates Bax and Bid expression and causes the release of MMP-12 and tissue inhibitor of metalloprotease-1 in mice (Kang et al., 2007). Bid-deficient mice are protected from developing fibrosis after TGF- β 1 activation, indicating that Bid is required for AEC apoptosis and bleomycin-induced fibrosis in mice (Budinger, 2006). In an experiment using mice, TGF- β 1 overexpression in lung epithelial cells induced apoptosis and fibrosis. This induction was attenuated when a caspase inhibitor was administered from day 0 but not from day 5 after TGF- β 1 overexpression (Lee et al., 2004), indicating that TGF- β 1-induced epithelial cell apoptosis is a critical early event in pulmonary fibrosis.

Oxidative Stress

Oxidative stress has been proposed to significantly contribute to epithelial cell damage in IPF (Kuwano et al., 2003). A number of studies demonstrate the increase of oxidative stress in IPF. Patients with IPF exhibit an increase in the spontaneous production of oxidants in the alveolar epithelial lining fluid (Saleh et al., 1997) as well as a significant reduction in antioxidant capacity in the plasma and BAL fluid (Rahman et al., 1999). Apoptosis plays a key role in hyperoxic lung injury (Albertine et al., 2002). Hyperoxia can affect both ATI and endothelial cells as well as induce DNA damage in ATII cells (Roper et al., 2004). Hyperoxia amplifies ventilator-induced cytokine production, neutrophil influx, and apoptosis through activation of the JNK and ERK pathway (Ward et al., 1983). It was traditionally thought that alveolar inflammatory cells produce the reactive oxygen species (ROS) that induce AEC injury/apoptosis in IPF (Cantin et al., 1987; Stausz et al., 1990). However, recent studies suggest that structural cells of the lung, particularly activated myofibroblasts, produce sufficient amounts of extracellular ROS to induce injury/apoptosis of adjacent epithelial cells. Bleomycin induces AEC apoptosis and fibrosis through an increase of ROS (Wallach-Dayana et al., 2006).

Alveolar Epithelium-Fibroblast Crosstalk

Severe injury to the lung epithelial cells followed by insufficient repair disturbs normal epithelial-fibroblast interactions. The lack of reepithelialization after damage allows fibroblasts to proliferate and eventually resulting in pulmonary fibrosis. Epithelial cells control fibroblasts through the release of cytokines. At the same time, fibroblasts can also affect alveolar epithelial cells by releasing factors that disrupt their normal repair.

Abnormal fibroblasts isolated from fibrotic human lungs produce factors that can induce apoptosis and necrosis of AECs *in vitro* (Uhal et al., 1995).

THE RENIN ANGIOTENSIN SYSTEM

General RAS

The traditional view of the renin angiotensin system (RAS) characterizes it as an endocrine system with important roles in the regulation of fluid homeostasis, electrolyte metabolism and blood pressure. Angiotensinogen (AGT), an α -glycoprotein, is released from the liver and cleaved by renin in the circulation to form the decapeptide, Angiotensinogen I (ANG I). Renin is an aspartyl protease secreted by the granular cells in the juxtaglomerular apparatus of the kidney in response to decreased renal perfusion and plasma sodium concentration. ANG I is activated by angiotensin converting enzyme (ACE) to form the octapeptide angiotensin II (ANG II). ACE, a membrane-bound metalloproteinase, is primarily expressed in high concentrations on the surface of endothelial cells in the pulmonary capillaries. Alternative pathways involving enzymes other than renin and ACE have also been shown to generate ANG I and ANG II. For example, ANG II may be generated by chymase released from mast cells in the vascular tissue under certain pathological conditions. ANG II is considered the main effector peptide of the RAS and exerts its actions through specific cell surface angiotensin receptors.

The main receptors of ANG II are AT₁ and AT₂, both belong to a superfamily of seven transmembran G-protein couple receptors. The AT₁ receptor mediates most of the well-known actions of ANG II, such as vasoconstriction, sodium retention, aldosterone release, and cell growth and proliferation, etc. Actions mediated by the AT₂ receptor are less well characterized but appear to counteract AT₁ effects by promoting vasodilation, cell differentiation, and inhibition of cell growth and apoptosis (de Gasparo et al., 2000).

In addition to ANG I and ANG II, several functional peptides can be generated through alternative cleavage pathways. These include angiotensin III (ANG III), angiotensin IV (ANG IV), and angiotensin 1-7 (ANG 1-7). ANG III is generated from ANG II by aminopeptidase A and plays a similar role as ANG II. Cleavage of ANG III by aminopeptidase M generates ANG IV, which is involved in the vascular inflammatory response (Ruiz-Ortega, 2007). ANG 1-7 heptapeptide is generated through cleavage of ANG II by angiotensin-converting enzyme 2 (ACE2). ANG 1-7 was found to have actions opposing those of ANG II and could act as a part of a counterbalancing mechanism in controlling the actions of ANG II in specific tissues (Ferrario et al., 2006).

Local RAS

Recent evidence of RAS components expression in most organs and tissues suggests the existence of a “local” RAS system that responds to physiological or pathophysiological stimuli in an autocrine, paracrine and/or intracrine manner. This local system contains all the components necessary to produce ANG II and other angiotensin peptides along with their respective receptors, as well as renin/prorenin receptors. Most, if not all of the renin found in local RAS is derived from renal renin. Local RAS systems may operate independently of or interact with “circulating RAS” to exert novel and tissue specific actions. These include cellular actions such as regulation of cell growth, differentiation, proliferation and apoptosis, reactive oxygen species (ROS) generation, hormonal secretion, tissue inflammation, and fibrosis (Leung et al., 2007; Montani & Van Vliet 2004).

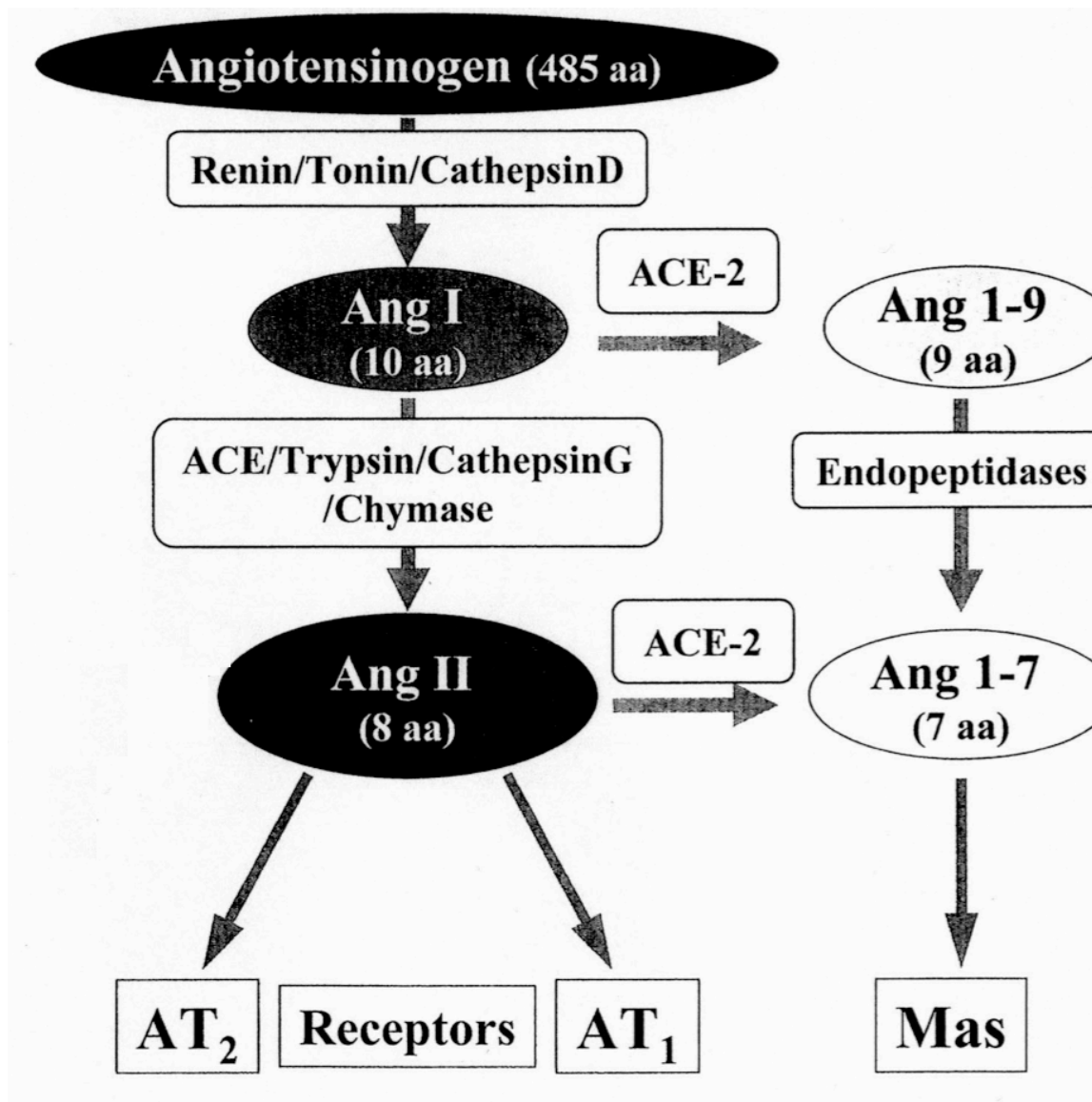


Figure 1.3.

The Renin Angiotensin System (RAS). The diagram illustrates the enzymatic cleavage of angiotensinogen protein into angiotensin I (ANG I), angiotensin II (ANG II), angiotensin 1-9 (ANG 1-9), angiotensin 1-7 (ANG 1-7). Also shown are the known receptors for angiotensin peptides: AT₁ (angiotensin receptor 1), AT₂ (angiotensin receptor 2), Mas. ACE: angiotensin converting enzyme; ACE-2: angiotensin converting enzyme-2; aa: amino acid.

BLEOMYCIN-INDUCED EXPERIMENTAL PULMONARY FIBROSIS MODEL

The bleomycin-induced experimental pulmonary fibrosis model is one of many animal models of pulmonary fibrosis that have been developed to address the pathogenesis of pulmonary fibrosis. Bleomycin is an antineoplastic antibiotic that was isolated from a strain of *Streptomyces verticillus* (Umezawa et al., 1966). Bleomycin is often used to treat squamous cell carcinomas and various lymphomas (Ichikawa et al., 1967, 1969; Yogoda et al., 1972; Blum et al., 1973). Instead of being just a single peptide, bleomycin consists of a family of complex glycopeptides with different amine groups (Umezawa et al., 1966, 1967; Umezawa 1973, 1974). The mechanisms of the antineoplastic actions of bleomycin are cell cycle-dependent and complex. Bleomycin can intercalate between DNA base pairs, leading to DNA unwinding and impaired protein synthesis. Bleomycin also reduces molecular oxygen to superoxide and hydroxyl radicals, producing radical oxygen species that then attack DNA and cause strand cleavage (Sausville et al., 1978).

Treatment with bleomycin has very little side effects on bone marrow (Kimura et al., 1972; Boggs et al., 1974) and immunocompetence (Dlugi et al., 1974; Lehane et al., 1975). Despite these advantages, bleomycin has severe toxic side effects on the lung and skin organ systems (Thrall and Scalise, 1995). After parenteral administration, bleomycin concentrations are highest in the lungs and skin, likely due to the lack of a hydrolase, which inactivates bleomycin (Umezawa et al., 1972).

The bleomycin-induced pulmonary fibrosis animal model is currently the most commonly used model in studying the pathogenesis of fibrotic lung disease. Pulmonary fibrosis induced by bleomycin is dependent on dosage and administration route, most common of which are parenteral and intratracheal. Various animals, such as mice and rats,

have been used in models of bleomycin-induced pulmonary fibrosis (Thrall and Scalise). The development of lung injury in these models is similar regardless of species and route of administration. The lung injury has three stages (Thrall and Scalise, 1995): (1) the acute inflammatory stage where the inflammatory mediator systems are activated and pulmonary edema appears; (2) The subacute stage where collagen is synthesized and the level of net lung collagen elevates; (3) the chronic stage that is dominated by the metabolism of connective tissue towards reepithelialization. The bleomycin animal models are valuable tools in studying the pathogenesis of fibrotic lung disease.

PULMONARY ANGIOTENSIN SYSTEM IN IDIOPATHIC PULMONARY FIBROSIS

Recent evidence suggests the existence of a local intrinsic RAS system in the distal lung parenchymal that plays a central role in the signaling of apoptosis in alveolar epithelial cells (Wang et al., 1999). ANG II in the lung, derived from its precursor AGT, drives fibrosis through upregulation of collagen gene expression in lung fibroblasts (Marshall et al., 2004; Marshall et al., 2000), induction of apoptosis in alveolar epithelial cells (Li et al., 2003), and other profibrotic actions (Marshall, 2003).

AGT in the local system is synthesized within the lung tissue itself by alveolar epithelial cells and myofibroblasts. Primary cultures of human or rat AECs can synthesize and secrete AGT, which is converted to ANG II when undergoing apoptosis induced by Fas ligand (Wang et al., 1999), TNF- α (Wang et al., 2000), amiodarone (Bargout et al., 2000) and bleomycin (Li et al., 2003). Myofibroblasts isolated from lungs of IPF patients were shown to synthesize AGT mRNA and protein constitutively (Wang et al., 1999). Intratracheal administration of antisense oligonucleotides against AGT mRNA prevented bleomycin-induced lung fibrosis in rats without affecting circulating levels of ANGEL protein, suggesting that local pulmonary synthesis of ANG II from the precursor AGT within the lung is required for lung fibrogenesis (Li et al., 2007).

The local angiotensin systems often utilize components of endocrine RAS system in its pathways. For example, synthesis of local ANG II in the heart and vessel walls rely on the uptake of circulating renin through either diffusion into the interstitial space or through binding to prorenin receptors (Re, 2004). Renin mRNA has not been detected in the lungs. Other proteases in the lung, such as cathepsin D (Cat D), are capable of cleaving AGT to

ANG I. Cat D expression is upregulated in the fibrotic lungs of both rat and human. Cat D is also overexpressed in the lungs of patients with pulmonary fibrosis (Kasper et al., 1996), and is induced in lung cells during apoptosis (Kasper et al., 1999). It is possible that production of ANG I from AGT in the lungs is mediated by Cat D.

The decapeptide ANG I is inactive until it is cleaved into ANG II by the dipeptidyl carboxypeptidase ACE. ANG II is thought to be responsible for most of the physiological and pathophysiological effects of the RAS. ACE in the pulmonary vascular endothelium is primarily responsible for the conversion of ANG I to ANG II in the circulation (Oparil et al., 1971). ACE2, the first known homologue of ACE, was recently discovered to be another regulator of the RAS system (Donoghue et al., 2000). ACE2 was first described for its ability to cleave ANG I into ANG 1-9, a peptide with no known function (Donoghue et al., 2000). It was later found to also degrade ANG II to the biologically active peptide, ANG 1-7 (Vickers et al., 2002). *In vitro* studies demonstrate that the catalytic efficiency of ACE2 for ANG II is 400 times greater than for ANG I (Vickers et al., 2002), suggesting that ACE2 is primarily involved in the conversion of ANG II to ANG 1-7. ACE2 expression has been identified in the heart, kidney, testis, gastrointestinal tract, brain, and lung (Hammer et al., 2002). Despite its similarities to ACE, the *in vitro* enzymatic activities of ACE2 is unaffected by ACE inhibitors (Donoghue et al., 2000; Tipnis et al., 2000).

Both receptors of ANG II, AT1 and AT2, are found to be expressed in the lung. In the human lung, AT1 receptor mRNA and proteins were found on vascular smooth muscle cells, macrophages and in the stroma underlying the airway epithelium (Bullock et al., 2001). AT2 receptor RNA and protein were highly expressed on the bronchial epithelial

cell brush border, endothelial cells, and underlying mucous glands of the epithelium (Bullock et al., 2001). A study of the rat lung also found AT1 to be expressed on alveolar macrophages, alveolar type II cells, vascular smooth muscle cells, endothelial cells and fibroblasts of the rat lung (Otsuka et al., 2004). The AT1 expression in the lung was significantly upregulated after induction of fibrosis (Otsuka et al., 2004), suggesting that ANG II promotes lung fibrosis via the AT1 receptor.

Although AGT in the lung can be synthesized by myofibroblasts, they have a limited ability to convert the AGT to ANG II (Wang et al., 1999). Alveolar epithelial cells, on the other hand, are capable of proteolytically processing AGT to ANG II as well as undergoing apoptosis in response to the ANG II produced (Wang et al., 1999). Thus, it is theorized that the death of AECs adjacent to underlying myofibroblasts in the fibrotic lung is due to production of AGT by the myofibroblasts and its subsequent conversion to ANG II by the epithelial cells.

The local RAS system can also induce apoptosis of AECs without myofibroblasts. Purified ANG II was shown to induce apoptosis of AECs in primary culture through the AT1 receptor (Papp et al., 2002). The induction of AEC apoptosis by Fas ligand (Wang et al., 1999), TNF- α (Wang et al., 2000), amiodarone (Bargout et al., 2000), and bleomycin (Li et al., 2003) were blocked by AGT antisense mRNA oligonucleotides, ACE inhibitors, and ANG II receptor antagonists, indicating that AEC apoptosis as induced by these agents require angiotensin synthesis de novo. The inhibition of bleomycin-induced AEC apoptosis with either an ACE inhibitor or ANG receptor antagonist was observed both *in vitro* and *in vivo*, indicating that ANG II may be required for apoptosis in both cultured cells and the intact lung.

Angiotensin system antagonists have been shown to attenuate lung fibrosis in experimental models. Applications of ACE inhibitors, which presumably prevent ANG II production, have been shown to block pulmonary fibrosis in animal models induced by various agents. Treatment with the ACE inhibitor captopril, for example, effectively inhibited monocrotaline (Molteni et al., 1985) as well as γ irradiation-induced lung fibrosis in rats (Ward et al., 1990). Captopril was also shown to inhibit the proliferation of human lung fibroblasts *in vitro* (Nguyen et al., 1994). The specific AT1 receptor antagonist L158809 and the nonthiol ACE inhibitor enalapril were also shown to have similar effects on radiation-induced pulmonary fibrosis in rats (Molteni et al., 2000). Captopril was also found to prevent collagen deposition on bleomycin-treated rats (Wang et al., 2000). Recently, it was shown that captopril and AT1 selective antagonist losartan block the amiodarone induced pulmonary fibrosis in rats (Uhal et al., 2002). Together, these results suggest that ANG II plays an important role in lung fibrosis through the AT1-receptor.

The ACE inhibitor captopril as well as caspase inhibitors have been found to block pulmonary fibrosis through the inhibition of apoptosis. Caspases play key roles in apoptosis signaling. Application of the broad-spectrum caspase inhibitor ZVADfmk, or captopril both attenuated epithelial apoptosis and collagen deposition in intratracheally administered bleomycin-induced pulmonary fibrosis in rats (Wang et al., 2000). Delivery of the same caspase inhibitor through inhalation also inhibited bleomycin-induced in mice (Kuwano et al., 2001). These results indicate that both ANG II and epithelial apoptosis are required for lung fibrogenesis.

Chapter 2

ANGIOTENSIN 1-7 REGULATES ALVEOLAR EPITHELIAL CELL SURVIVAL THROUGH THE RECEPTOR MAS

ABSTRACT

Apoptosis of alveolar epithelial cells (AECs) is believed to be a critical event in the pathogenesis of lung fibrosis. It was previously shown that apoptosis of AECs requires autocrine generation of angiotensin (ANG) II. ANG II can also be degraded to ANG1-7 by angiotensin converting enzyme-2 (ACE-2). ACE2 was recently found to be protective but significantly downregulated in experimental lung fibrosis and patients with IPF. In other organ systems, ANG1-7 has been found to inhibit the actions of ANGII through the ANG1-7 receptor *mas*. Therefore, it was hypothesized that ANG1-7 in the lungs might antagonize the actions of ANG II in the regulation of AEC apoptosis. To test this theory, the AEC cell line MLE-12 and primary cultures of human AECs were stimulated by the profibrotic apoptosis inducers ANG II or bleomycin (BLEO). Caspase-3 activation and nuclear fragmentation were used as markers of apoptosis and were measured along with JNK phosphorylation. Exposure to ANG II or BLEO induced caspase-3 activation, nuclear fragmentation, and JNK phosphorylation in cultured AECs. Pretreatment with ANG1-7, at a concentration of 0.1 μ M, prevented JNK phosphorylation and apoptosis. In addition, pretreatment with A779, a specific blocker of the ANG 1-7 receptor *mas*, successfully prevented ANG1-7 induced blockade of JNK phosphorylation and nuclear fragmentation. These data indicate that ANG1-7 prevents AEC apoptosis, and its actions are mediated through the ANG1-7 *mas* receptor.

INTRODUCTION

Idiopathic pulmonary fibrosis is a pathological condition resulting from lung injury and an ensuing fibrotic response that leads to thickening of the alveolar walls and the obliteration of the alveolar space (Fonseca et al., 2000). The etiology of the disease is unknown. Current evolving hypothesis about pathogenesis is that IPF results from the epithelial injury and a failure of reepithelialization (Selman et al., 2001). Alveolar epithelial cell apoptosis is a critical event in the pathogenesis of pulmonary fibrosis and the regulation of AEC apoptosis is of great interest and potential clinical importance. The bleomycin animal models are valuable tools in studying the pathogenesis of fibrotic lung disease. Recent studies in our lab and other labs using the bleomycin-induced models in rats and mice suggested a role of epithelial apoptosis as the profibrotic event in fibrosis. First, apoptosis of AECs was found in both patients with IPF (Uhal, et al., 198) and animal models (Hagimoto et al., 1997). Second, induction of apoptosis in the epithelium is sufficient to produce a fibrotic response (Hagimoto et al., 1997). And lastly, several labs have shown that the blockade of apoptosis could prevent the fibrotic response (Wang et al., 2000; Kuwano et al., 2001).

Recent work from this lab demonstrated that the exposure of cultured AECs to Fas ligand (Wang et al, 1999), tumor necrosis factor- α (Wang et al., 2000) or bleomycin (Li et al., 2003) induce expression of angiotensinogen mRNA and protein and its cleavage to the peptide Angiotensin II. They also showed that bleomycin induces fibrosis through induction of alveolar epithelial cell apoptosis (Wang et al., 2000). They found that the ACE inhibitor captopril and the broad-spectrum caspase inhibitor ZVAD-fmk were able to reduce bleomycin-induced apoptosis (Figure 2.1). Moreover, captopril and ZVAD-fmk also

prevented the bleomycin-induced accumulation of collagen (Figure 2.2). Together, these results indicate that the induction of apoptosis is required for bleomycin-induced fibrosis and that ANG II and caspases play an important role in this pathway.

In another study, our lab demonstrated that bleomycin-induced apoptosis of alveolar epithelial cells requires ANG II synthesis *de novo*. Li et al. showed that bleomycin-induced nuclei fragmentation was blocked by ZVAD-fmk, captopril and the nonselective ANG receptor antagonist saralasin (Figure 2.3). In addition, they also demonstrated that bleomycin-induced apoptosis of primary AECs was blocked by the caspase-3-selective blocker DEVD-fmk and the AT₁ receptor selective blocker losartan (Figure 2.4), suggesting that AT₁ receptor mediates bleomycin-induced apoptosis as it does ANGII-induced AEC apoptosis. Together, these data indicates that the production of ANGII is a necessary event in bleomycin-induced AEC apoptosis.

Bleomycin induces alveolar cell apoptosis through the JNK-dependent activation of the mitochondrial death pathway (Lee et al., 2005). Bleomycin-induced Bax activation was inhibited by the expression of a dominant negative JNK in MLE-12 cells (Lee et al., 2005). Additionally, recent evidence show that both bleomycin and ANGII induces caspase 9 activation, suggesting that ANG II, just like bleomycin, also induces apoptosis through the mitochondrial death pathway requiring the activation of phosphorylation of JNK (Figure 2.5).

Angiotensin converting enzyme 2(ACE-2) degrades ANG II into the heptapeptide ANG 1-7 (Figure 2.6). In other organ systems, ANG1-7 was found to prevent some of the actions of ANGII through the ANG1-7 receptor *mas* (Gava et al., 2009; Lara et al., 2010; Tallant et al, 2005). Precisely which lung cell types express active *mas* signaling is

currently unknown. This study reports evidence that ANG1-7 in the lung acts through the ANG1-7 receptor *mas* to inhibit ANGII signaling and AEC apoptosis.

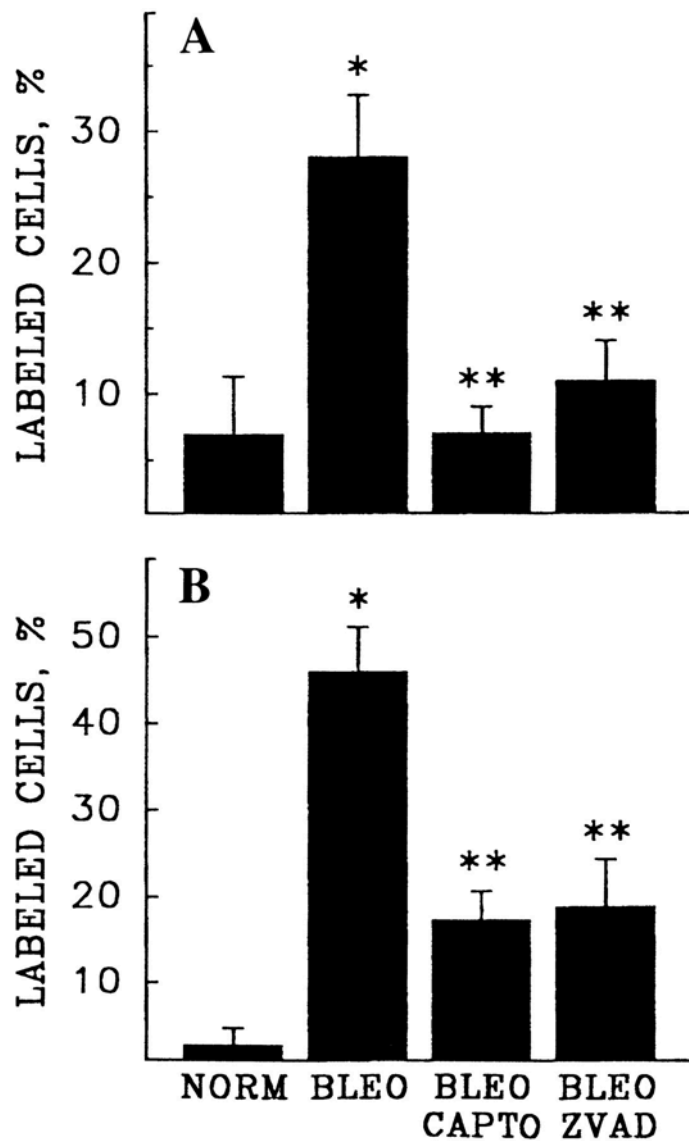


Figure 2.1.

Quantitation of ISEL-positive nuclei in alveolar (A) and airway (B) epithelial cells after Bleo, CAPTO, and ZVAD-fmk administration. NORM, animals exposed to vehicle only. ISEL-positive nuclei were scored as a percentage of the total epithelial nuclei within the alveolar and airway epithelial cell populations. See methods for details. * $P < 0.01$ vs. NORM. ** $P < 0.01$ vs. Bleo (both by ANOVA and Student-Newman-Keuls test). (Adapted from Wang et al., 2000)

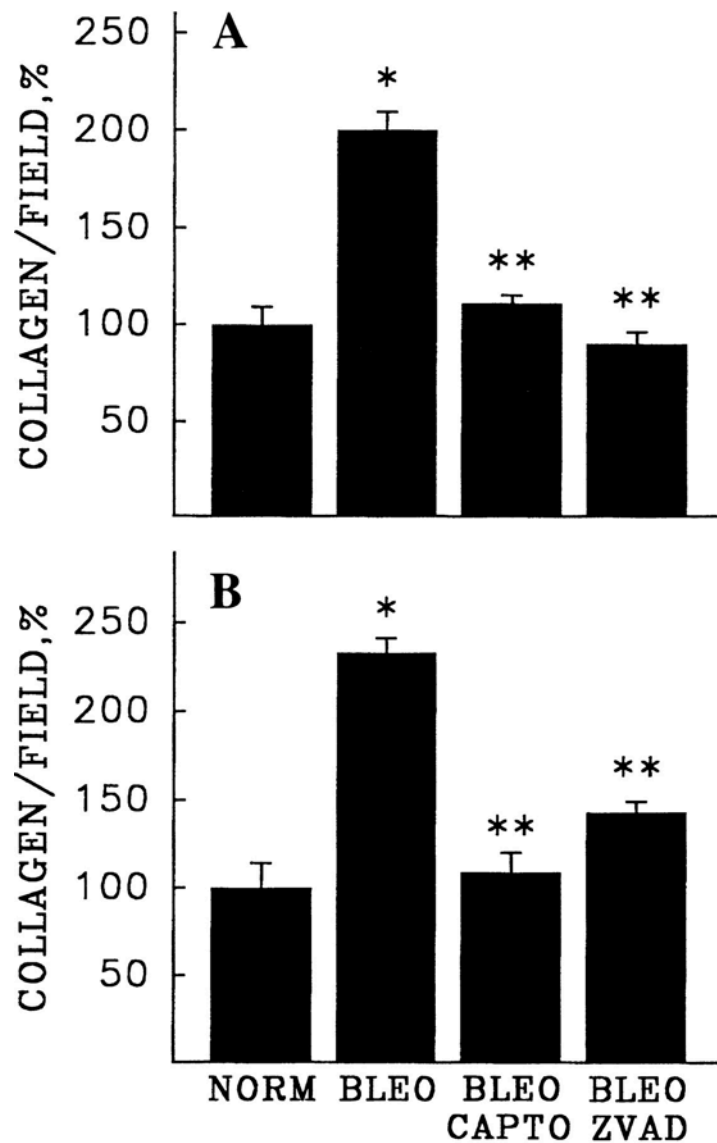


Figure 2.2.

Quantitation of collagen by PR staining of alveolar (A) and peribronchial (B) lung parenchyma. Collagen was quantitated from polarized light images as the total number of white pixels per unit area within the alveolar and peribronchial lung parenchyma. See methods for details. * $P < 0.01$ vs. NORM. ** $P < 0.01$ vs. Bleo (both by ANOVA and Student-Newman-Keuls test). (Adapted from Wang et al., 2000)

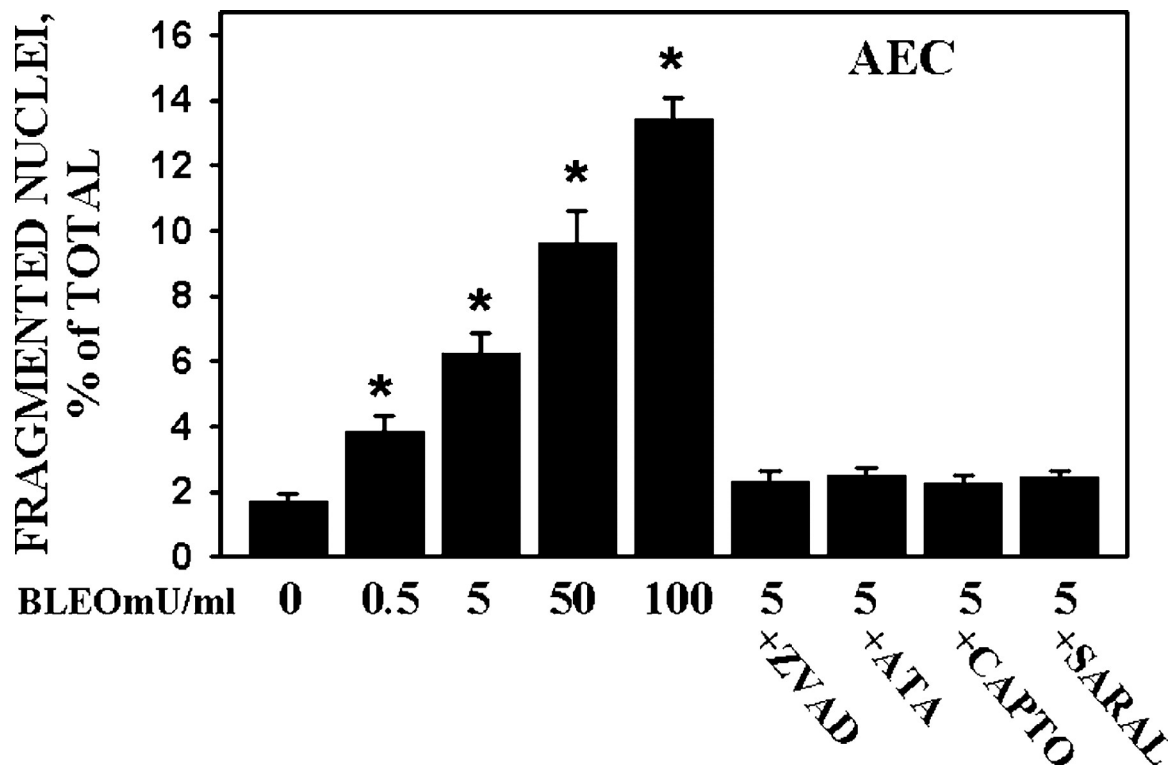


Figure 2.3.

Dose-dependent induction of nuclear fragmentation by BLEO in primary rat AECs and blockade by inhibitors of caspases, endonucleases, ANG-converting enzyme (ACE), and ANG-receptor interaction. Rat AECs were isolated and challenged with the indicated concentrations of Bleo on day 2 of primary culture (see materials and methods). Putative inhibitors were added 30 min before addition of Bleo; nuclear fragmentation was scored as described in Fig. 1 B and materials and methods. ZVAD, N-benzylcarboxy-Val-Ala-Asp-[O-Me]-CH₂F (60 μM); ATA, aurintricarboxylic acid (10 μM); Capto, captopril (500 ng/ml); Saral, saralasin (50 μg/ml). Bars are the means ± SE of at least 4 observations; *P < 0.05 vs. control (0.0 Bleo). (Adapted from Li et al., 2002)

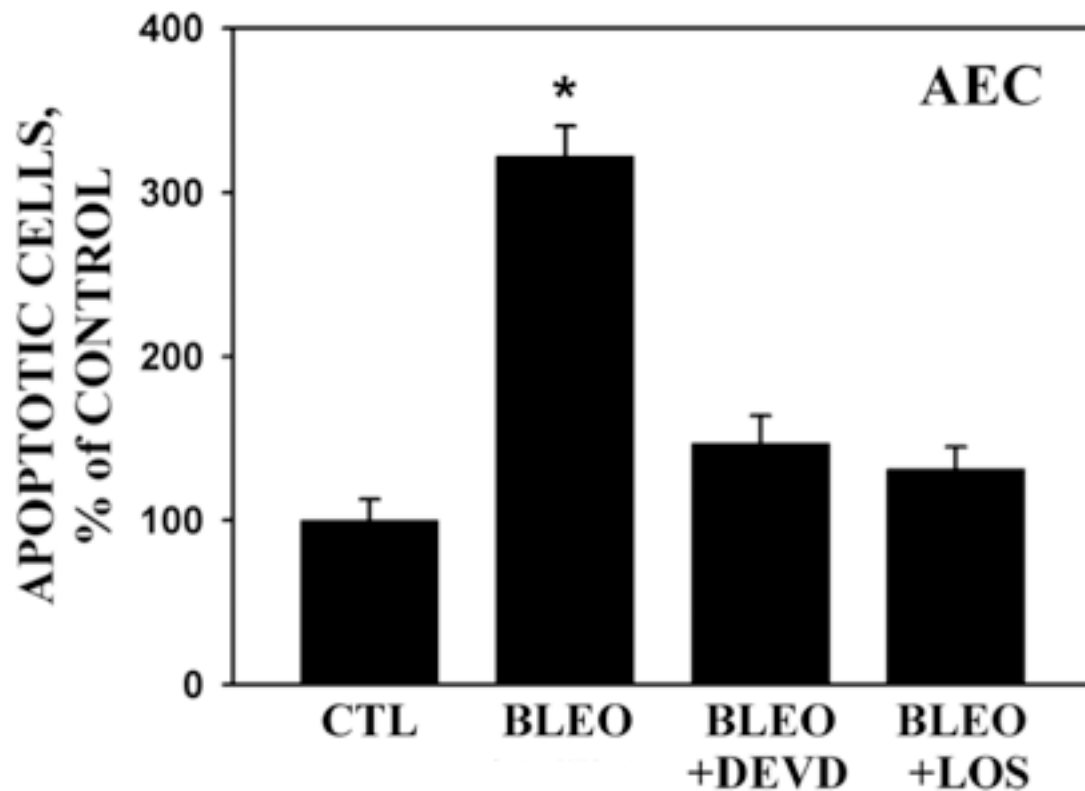


Figure 2.4.

Blockade of Bleo-induced apoptosis in primary AECs by selective caspase or ANG receptor blockers. Rat AECs were isolated and challenged with 25 mU/ml Bleo alone or in the presence of the caspase-3-selective inhibitor Asp-Glu-Val-Asp-[O-Me]-CH₂F (DEVD-fmk, 60 μ M) or the ANG receptor AT₁-selective antagonist losartan (Los, 10⁻⁶ M). Control cultures (Ctl) received blocker vehicles only. Nuclear fragmentation was scored as described in Fig. 1 and materials and methods. Bars are means $\pm$ SE of at least 4 observations; *P < 0.05 vs. control. (Adapted from Li et al., 2002)

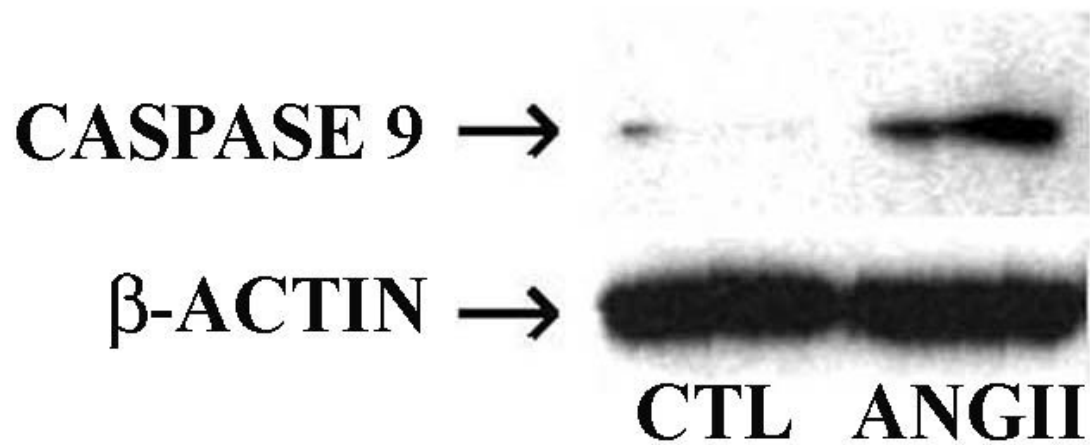


Figure 2.5.

ANGII induces activation of caspase-9. Primary cultures of rat type II alveolar epithelial cells were cultured in serum-free medium for 6 hours with or without ANGII. Cells were harvested with NP40 buffer for Western blotting. (Xiaopeng Li, 2006)

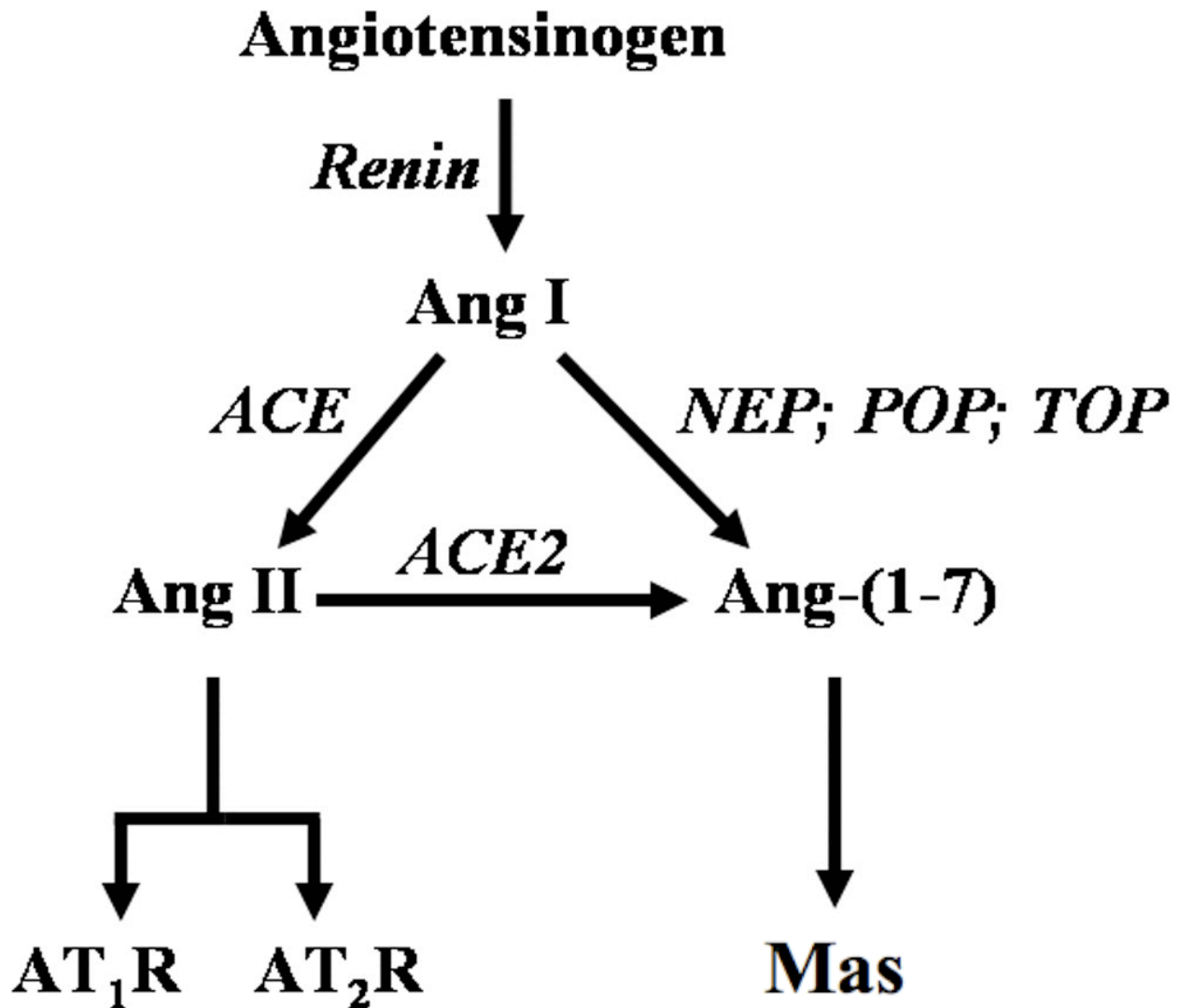


Figure 2.6.

Pathways for the formation of ANG 1-7. Renin cleaves angiotensinogen into the 10-aa peptide ANG I. ANG I is then converted to angiotensin II by ACE or to ANG 1-7 by neprilysin (NEP), prolyl-endopeptidase (POP) or thimet oligopeptidase (TOP). ANG 1-7 can also be generated from ANG II through ACE2. ANG II can activate either the AT₁ subtype of angiotensin receptor or the AT₂ ANG II subtype receptor. ANG 1-7 activates the Mas receptor.

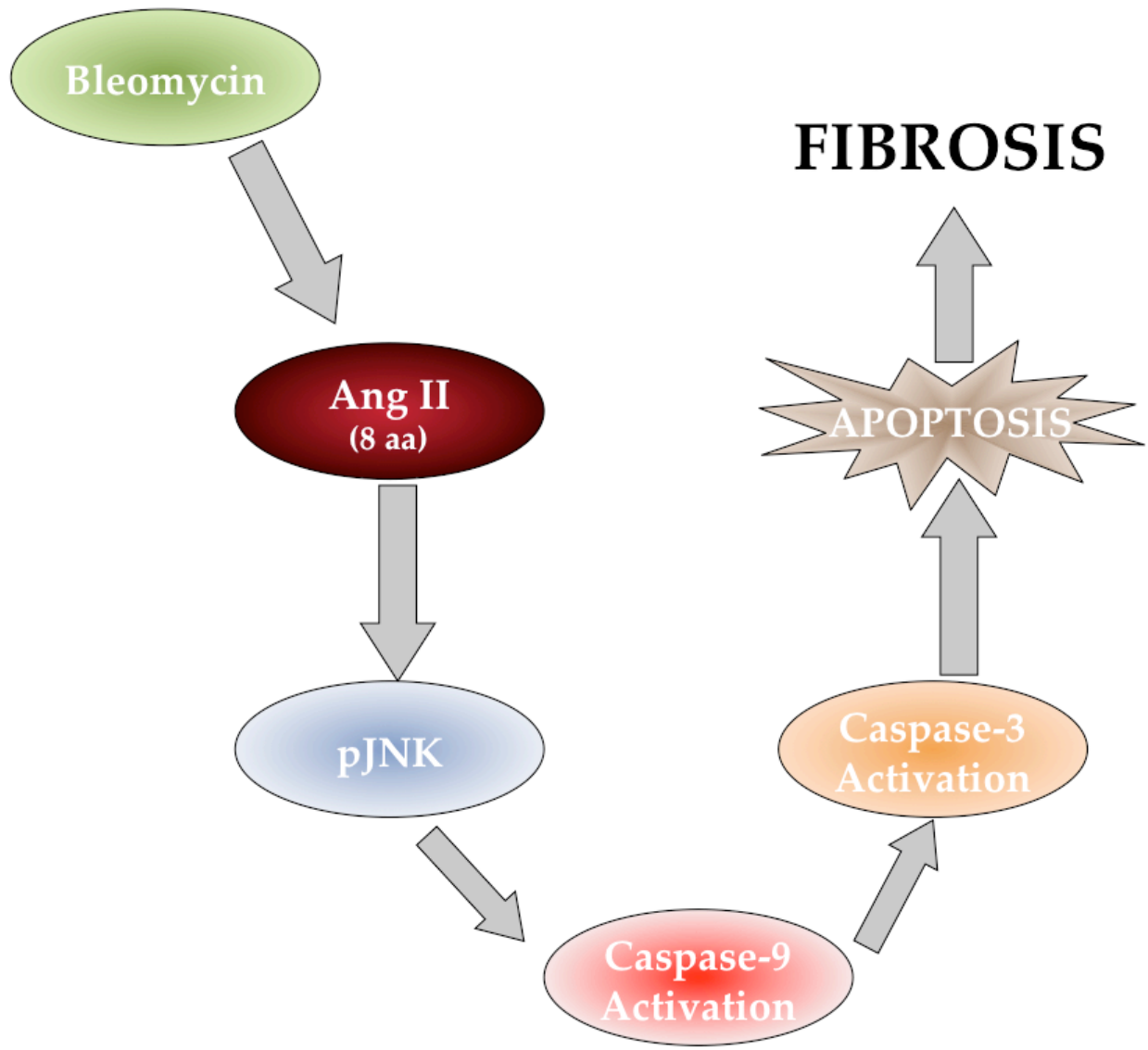


Figure 2.7.

Summary of Earlier Published Data. For interpretation of the references to color in this and all other figures, the reader is referred to the electronic version of this thesis.

METHODS

Cell culture

The mouse lung epithelial MLE-12 cell line was a gift from the laboratory of Dr. Jeffrey Whitsett, University of Cincinnati, OH, and was grown in complete HITES medium for MLE12 supplemented with 2% fetal bovine serum according to ATCC's guidelines. All cells were grown in 12-, 24-, or 6-well chambers and were analyzed at subconfluent densities of 50-80%. 12 hours before treatment, cells were switched from growth medium containing fetal bovine serum to serum-free medium. All subsequent incubations with bleomycin and/or other test agents were performed in serum free medium. In all experiments, cells were treated with inhibitors or antagonists 30 minutes prior to treatment with bleomycin or ANGII. For exposure of extended periods, cells were exposed to test agents as described and after one hour, culture media were replaced with new media containing fresh A779 and ANG1-7 but not bleomycin. Due to the low biological half-lives of the A779 and ANG1-7 peptides, they were replaced every 2 hours until cells were harvested. Primary human alveolar epithelial cells isolated from human lung tissue were obtained from ScienCell Research Laboratories. They were cultured in Alveolar Epithelial Cell Medium according to ScienCell Research Laboratories guidelines.

Detection of Apoptosis

Apoptotic cells were detected by nuclear fragmentation with propidium iodide (PI) after enzymatic digestion of ethanol-fixed cells with DNase-free RNase in PBS containing 5µg/ml PI. During fixation with 70% ethanol, detached cells were retained by centrifugation of the 24-well or 12-well culture plates. Cells with discrete nuclear

fragmentation containing condensed chromatin were scored as apoptotic. Apoptotic cells were scored over four separate microscopic fields from each of four culture wells per treatment group. The active forms of caspase-3 and caspase-9 were detected by Western blotting using antibodies specific for the active cleaved forms.

Western Blotting

For caspase-3 and caspase-9 detections, cells were harvested and lysed in an NP40-based lysis buffer containing protease inhibitors. For detection of phosphor-proteins, cells were harvested and lysed with a modified lysis buffer containing 50mM HEPES, 150mM NaCl, 10% glycerol, 1% Triton X-100, 1uM EGTA, 1.5mM MgCl₂, phosphatase inhibitor cocktail (Phospho Stop, Roche, Nutley, NJ) and the protease inhibitor cocktail (Complete Mini, Roche, Nutley, NJ). Proteins were separated on polyacrylamide gels and transferred to PVDF membranes. Immunoreactive bands were visualized by ECL detection systems (Thermo Scientific, Rockford, IL).

RESULTS

Earlier results from this lab, as summarized in figure 2.7, proposes that pulmonary fibrosis induced by bleomycin begins with an increased production of ANG II. ANG II then induces JNK phosphorylation and downstream activation of caspase 9 and 3. The signaling eventually leads to apoptosis of the alveolar epithelial cells followed by fibrosis. In this study, we wanted to investigate how Angiotensin 1-7 fits in to this pathway and what role it plays in the regulation of AEC apoptosis.

To begin examining the hypothesis that ANG1-7 acts to oppose the actions of ANG II, we assessed the effect of ANG II on JNK phosphorylation in the presence and absence of purified ANG1-7 in serum-free media. JNK phosphorylation has previously been shown to be required for apoptosis in alveolar epithelial cells (Lee et al., 2005). Figure 2.8 shows that 5 minutes of exposure to ANG II (10^{-7} M) also induces JNK phosphorylation. 30 minute pre-incubation with ANG1-7 (10^{-7} M) prevented the ANG II-induced increase in JNK phosphorylation, indicating that ANG1-7 is able to block the actions of ANG II at equimolar concentrations. Moreover, incubation with the specific *mas* receptor blocker A779 blocked the effects of ANG1-7. This indicates that ANG1-7 is able to signal through its receptor *mas* to prevent ANGII induced JNK phosphorylation.

In a previous study, it was shown that bleomycin-induced apoptosis of AECs requires autocrine synthesis of ANGII and its subsequent binding to ANG receptor AT₁ (Li et al., 2003). Given that ANG1-7 inhibited ANG II induced JNK phosphorylation (Figure 2.8), we theorized that it may also block the JNK phosphorylation that is required in bleomycin-stimulated JNK phosphorylation. A 30-minute pretreatment with ANG1-7 (10^{-7} M) prevented the increase in p-JNK observed 5 minutes after exposure of MLE-12 cells to

purified bleomycin (10^{-7} M) (Figure 2.8). We then tested bleomycin-induced JNK phosphorylation again with a 30-minute pretreatment of the *mas* antagonist A-779 (D-Ala⁷-Ang1-7, 10^{-7} M). The A779 pretreatment prevented the pJNK inhibition by ANG1-7 (Figure 2.9), suggesting that ANG1-7 prevents bleomycin-induced JNK phosphorylation through the receptor *mas*.

Since ANG1-7 prevented both ANG II- and bleomycin-induced JNK phosphorylation, and JNK phosphorylation is a required event in bleomycin-induced apoptosis, we then examined if ANG1-7 might also prevent AEC apoptosis. In Figure 2.10, preincubation with ANG1-7 blocked the bleomycin-induced activation of caspase-3 in MLE-12 cells. Figure 2.11 shows that ANG1-7 could also prevent bleomycin-induced caspase-9 activation in primary cultures of human alveolar epithelial cells and that the *mas* antagonist A779 prevents the action of ANG1-7, confirming that bleomycin induces apoptosis through the mitochondrial pathway as previously established. Furthermore, Figure 2.12 shows that ANG1-7 prevented the bleomycin-induced increase in nuclear fragmentation detected in MLE-12 cells after 12 hours exposure to bleomycin ($p < 0.01$). Preincubation with the *mas* antagonist A779 prevented the effect exerted by ANG1-7. It was previously demonstrated that bleomycin-induced AEC apoptosis involves the release of cytochrome c, indicating that apoptosis is occurring versus necrosis. In Figure 2.13A, MLE-12 cells were treated with 25mU/ml bleomycin for 20 hours and stained with active caspase-3 antibodies. The results further confirm the activities of apoptosis. Figure 2.13 also shows that ANG1-7 could also prevent bleomycin-induced nuclear fragmentation in primary cultures of human alveolar epithelial cells ($p < 0.01$). As seen in MLE-12 cells, the blockade of nuclear fragmentation by ANG1-7 was prevented by A779. Both nuclear fragmentation and

caspase-3 activation are markers of apoptosis. Figure 2.14 provides a brief summary of the results of this study and related earlier works, along with topics for future research.

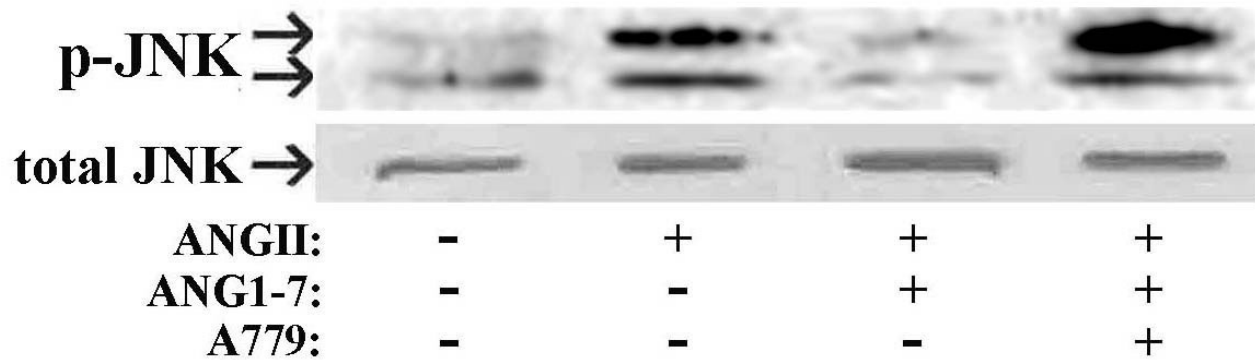


Figure 2.8.

ANG1-7 inhibits ANG II induced JNK phosphorylation through the *mas* receptor. MLE-12 cells in serum free conditions were preincubated with specific *mas* antagonist A779 10^{-7} M for 30 minutes followed by incubation in 10^{-7} M ANG1-7 for an additional 30 minutes. Afterwards, cells were treated with 5mU/ml ANG II for 10 minutes and harvested for Western blotting.

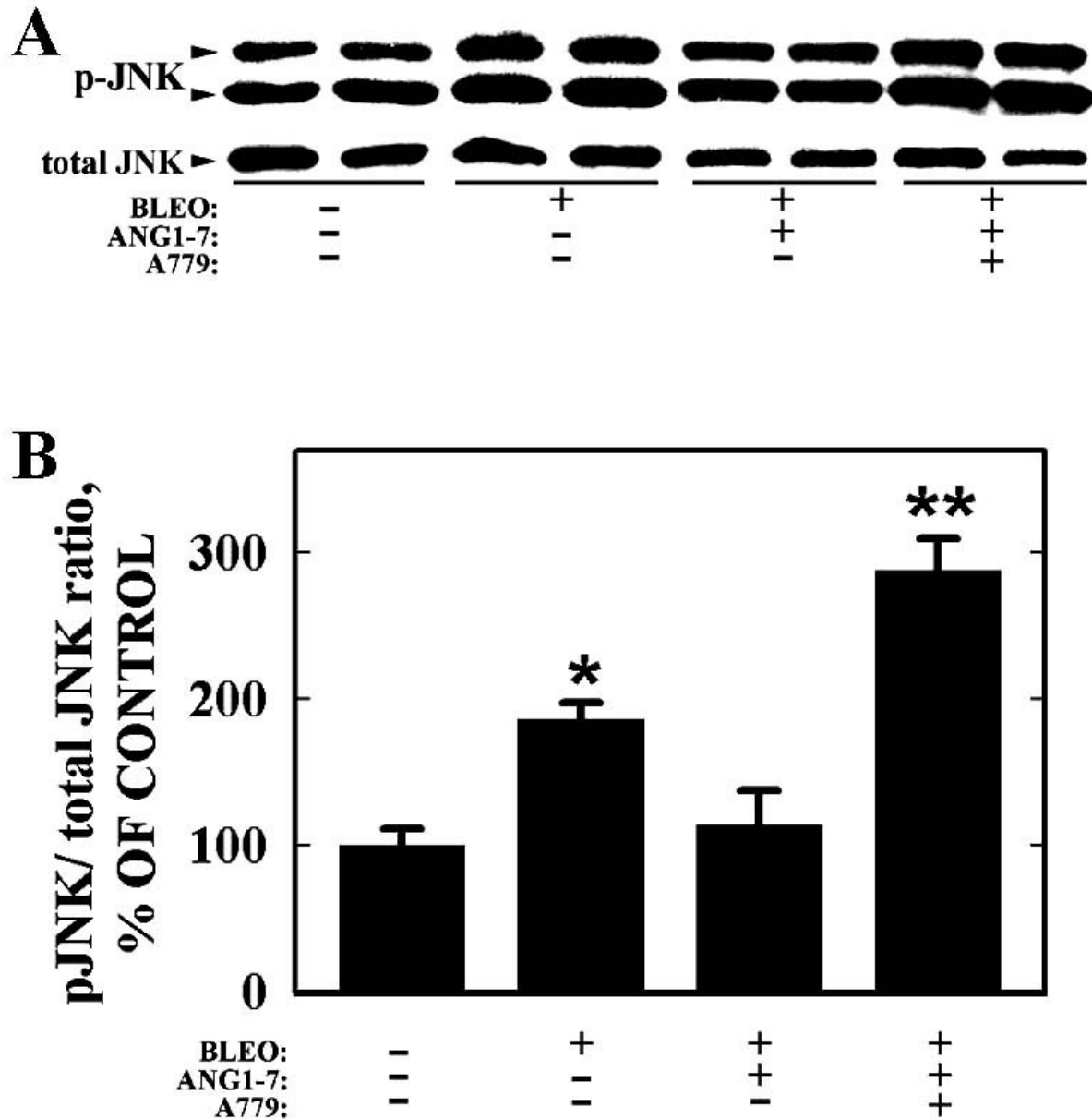


Figure 2.9.

ANG1-7 inhibits BLEO induced JNK phosphorylation through the *mas* receptor. A: MLE-12 cells in serum free conditions were preincubated with specific *mas* antagonist A779 10^{-7} M for 30 minutes followed by incubation in 10^{-7} M ANG1-7 for an additional 30 minutes. Afterwards, cells were treated with 5mU/ml bleomycin (BLEO) for 5 minutes and harvested for Western blotting. B: Ratio of the densitometric quantification of the Western blot performed in Panel A and the Western blot of total JNK of the same conditions (data not shown). Bars are the means. *: $p < 0.05$ vs. CTRL and **: $p < 0.01$ vs CTRL by ANOVA and Student-Newman-Keul's post-hoc analysis.

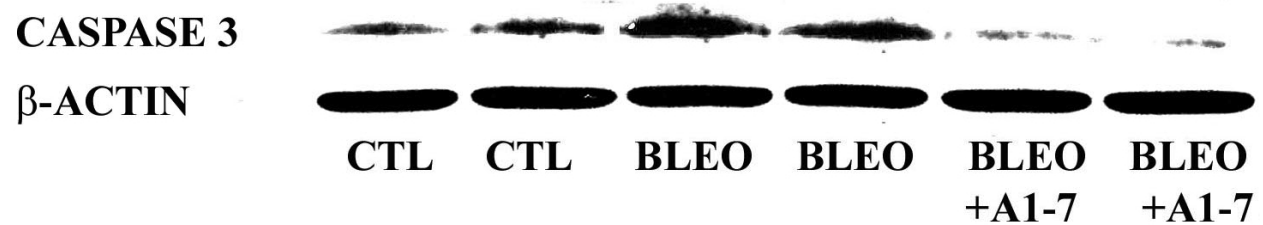


Figure 2.10.

ANG1-7 inhibits BLEO induced caspase-3 activation of MLE-12 cells. MLE-12 cells in serum free conditions were preincubated in 10^{-7} M ANG1-7 for 30 minutes. Afterwards, cells were treated with 5mU/ml bleomycin (BLEO) for 2 hours and harvested for Western blotting.

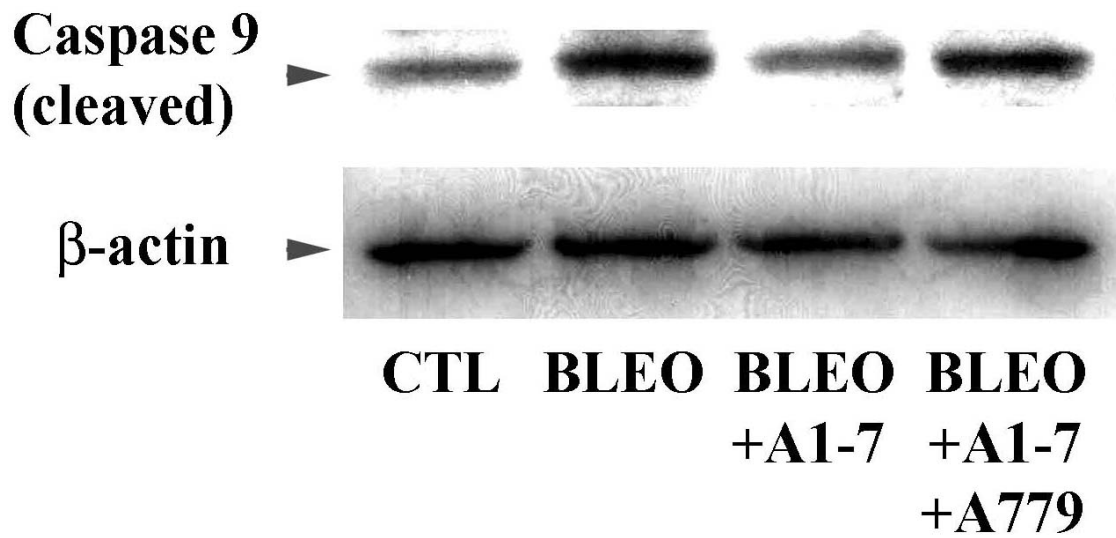


Figure 2.11.

ANG1-7 inhibits BLEO induced caspase-9 activation of primary AECs through the *mas* receptor. Primary cultures of human alveolar epithelial cells were preincubated with the specific *mas* antagonist A779 (10^{-7} M) for 30 minutes followed by ANG1-7 (10^{-7} M) for an additional 30 minutes. Thereafter the cells were exposed to bleomycin (BLEO, 5mU/ml) for an additional 6 hours.

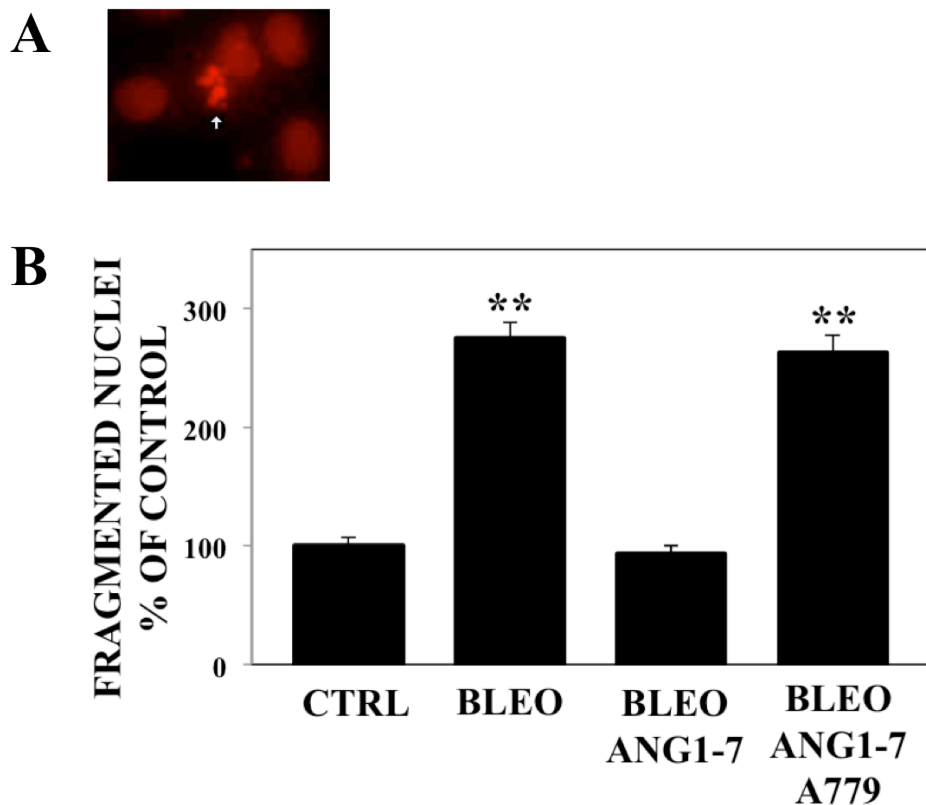
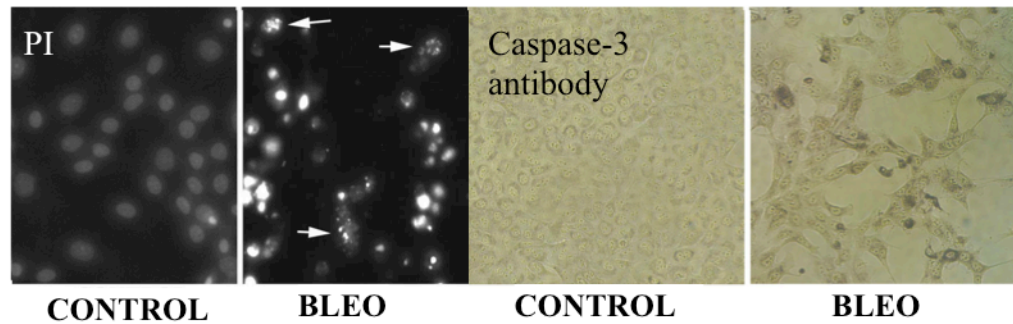
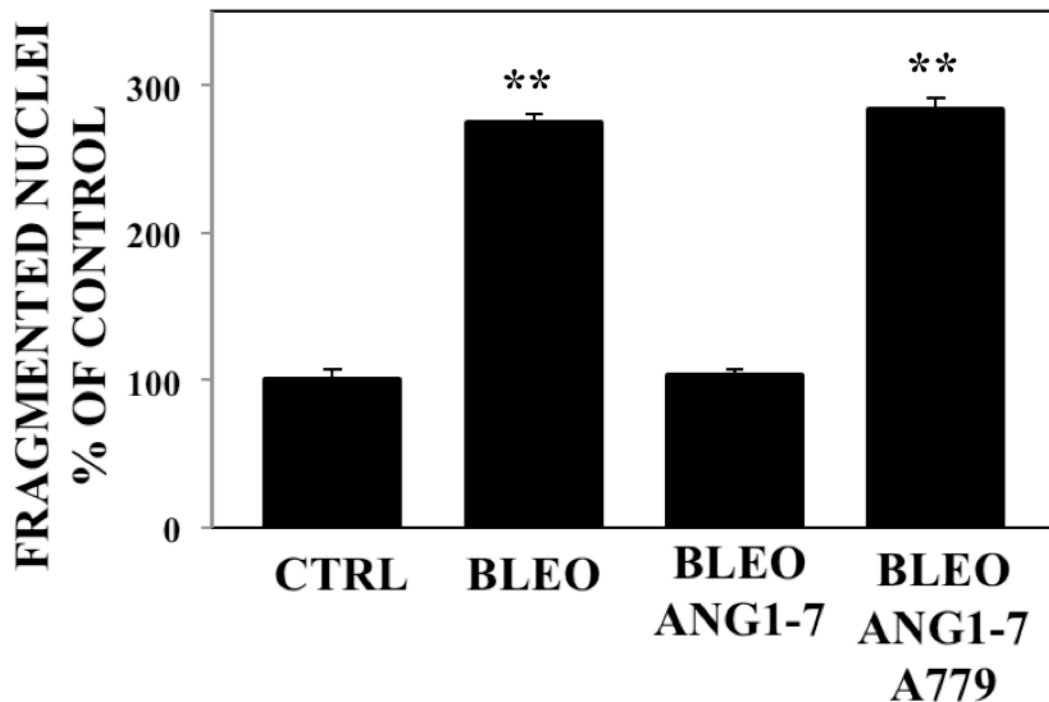


Figure 2.12.

ANG1-7 inhibits bleomycin-induced nuclear fragmentation through the *mas* receptor in MLE-12 cells. A: Primary cultures of AECs fixed in 70% ethanol and stained with propidium iodide. Arrow points to cells that exhibit chromatin condensation and nuclear fragmentation. B: MLE-12 cells in serum free conditions were preincubated with specific *mas* antagonist A779 10^{-7} M for 30 minutes followed by incubation in 10^{-7} M ANG1-7 for an additional 30 minutes. Afterwards, cells were treated with 5mU/ml bleomycin (BLEO) for 12 hours. A779 and ANG1-7 were replaced every 2 hours. Cells were then harvested for detection of nuclear fragmentation. Absolute value for basal (CTRL) nuclear fragmentation was $3.30 \pm 0.25\%$. Bars represent the means $\pm$ S.E.M. of four separate cell cultures. **: $p < 0.001$ vs. CTRL by ANOVA and Student-Newman-Keul's post-hoc analysis.

A**B****Figure 2.13.**

ANG1-7 inhibits bleomycin-induced nuclear fragmentation through the *mas* receptor in primary human AECs. A. Cells treated with 25 mU/ml bleomycin for 20 hrs stained with PI or with active caspase-3 antibody. Arrows indicate fragmented nuclei with condensed chromatin (bright fluorescence). B. Primary human alveolar epithelial cells in serum free conditions were preincubated with specific *mas* antagonist A779 10^{-7} M for 30 minutes followed by incubation in 10^{-7} M ANG1-7 for an additional 30 minutes. Afterwards, cells were treated 10^{-7} with 5mU/ml bleomycin (BLEO) for 12 hours. A779 and ANG1-7 were replaced every 2 hours. Cells were then harvested for detection of nuclear fragmentation. Absolute value for basal (CTRL) nuclear fragmentation was $3.11 \pm 0.21\%$. Bars represent the means $\pm$ S.E.M. of four separate cell cultures. **: $p < 0.001$ vs. CTRL by ANOVA and Student-Newman-Keul's post-hoc analysis.

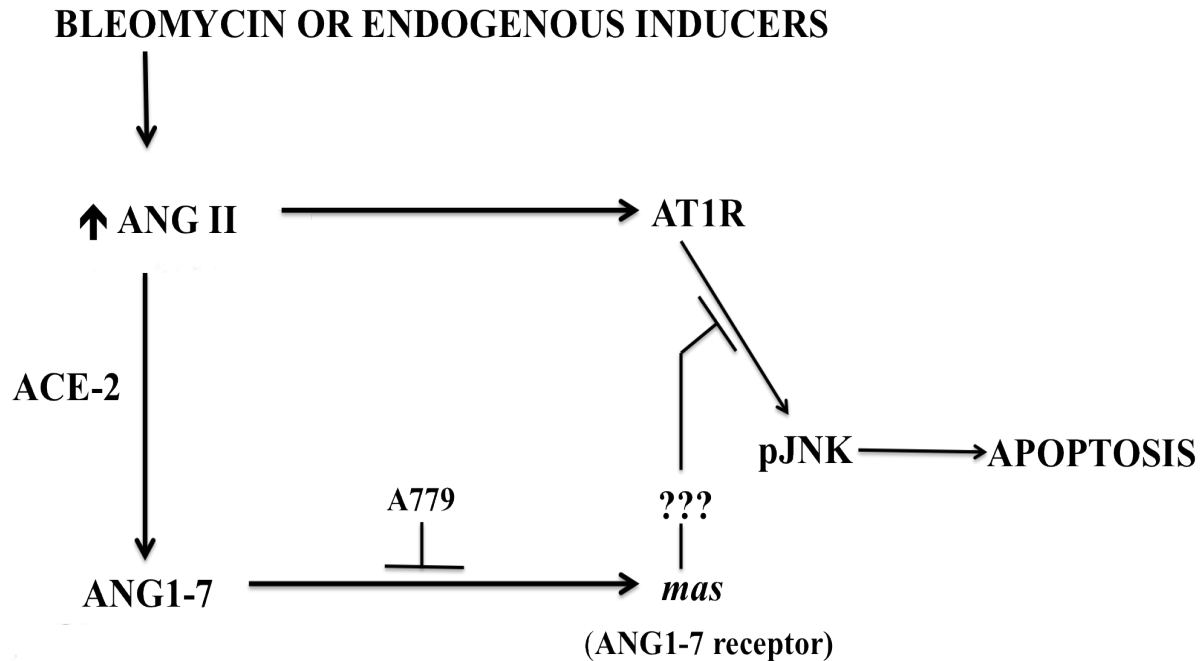


Figure 2.14.

Summary. The autocrine production of ANGII is required for AEC apoptosis and can be stimulated by bleomycin or the endogenous proapoptotic inducers Fas ligand or TNF- α . Apoptosis signaling proceeds through the ANG receptor AT₁ and JNK phosphorylation (pJNK). ANG 1-7 inhibits JNK phosphorylation and apoptosis through binding to the specific ANG 1-7 receptor *mas*. The mechanisms of how activation of *mas* leads to inhibition of JNK phosphorylation are currently unknown and may be topics for future inquiry.

DISCUSSION

Recent studies have established a new regulatory axis in the renin-angiotensin system (RAS). Much attention has been paid to studying the role of the ACE-2/ANG1-7/*mas* axis in organ injury and fibrogenesis. The heptapeptide ANG1-7, has been shown to oppose the vasoconstrictive, proliferative, and profibrotic actions of ANG II in a number of tissues (Ferrario et al., 2005; Ferrario 2010). ANG1-7 can be generated through cleavage by a variety of enzymes including neprilysin, prolyl-endopeptidase, and thimet oligopeptidase (Gallager and Tallant, 2004). The primary pathway of ANG 1-7 synthesis is through ACE2 degradation of ANG II, in which ACE2 removes the ANG II C-terminal phenylalanine to produce ANG1-7 (Li et al., 2008). Figure 2.6 illustrates the cleavage pathway of AGT into ANG 1-7. Studies of the adult mouse lung indicate that the type II pneumocytes are the only cells to express ACE2 in the lungs (Wiener et al., 2007).

The heptapeptide ANG1-7 can bind to the receptor *mas*, a seven trans-membrane protein with domains containing sequences characteristic of G protein-coupled receptors encoded by the oncogene of the same name (Young et al., 1988). Some suggested that ANG1-7 may also bind to the AT₁ receptor (Zisman, 2005). Binding data, however, reveal low affinity of ANG1-7 for AT₁ or AT₂ receptors (Fontes et al., 1994). Currently there is very little data regarding the functions of ANG1-7 in apoptosis and its role in the regulation of organ systems. In the liver, ACE-2 and ANG1-7 may be protective against the development of liver fibrosis (Warner et al., 2007), the molecular and cellular mechanisms of which are still unclear.

Evidence from numerous cell systems demonstrates the ability of ANG1-7 to inhibit the effects of ANG II. For example, in the heart, ANG 1-7 prevents the development of

cardiac fibrosis induced by ANG II infusion (Grobe et al., 2007), and attenuates the ANG II-stimulated production of endothelin-1 and leukaemia inhibitory factor in cardiac fibroblasts (Iwata et al., 2005). In the rat proximal tubular cells, ANG 1-7 inhibited ANG II-induced phosphorylation of MAPKs including p38, ERK1/2, and JNK while also partially blocking activation of TGF- β 1 promoted by ANG II (Su et al., 2006). Also in the kidney, ANG1-7 was found to reverse ANG II-induced stimulated Na⁺-ATPase (Lara et al., 2010). In all of these systems, ANG 1-7 blockade of ANG II actions was inhibited by a knockdown of *mas* or the application of the specific *mas* receptor blocker A779.

Consistent with these results from other experimental systems, Figure 2.8 demonstrates the ability of ANG 1-7 to prevent JNK phosphorylation stimulated by ANG II in alveolar epithelial cells. In addition, Figure 2.12 indicates that ANG 1-7 also prevents apoptosis induced by bleomycin. The finding that bleomycin activates caspase-9 (Figure 2.11) confirms that bleomycin also induces apoptosis through the mitochondrial pathway and that this induction is prevented by A1-7, whose effects, in turn, were blocked by A779. Together, our data reinforces the previously established concept that autocrine production of ANG II and its binding to the AT₁ receptor is required in bleomycin-induced apoptosis of alveolar epithelial cells. As seen in Figure 2.12 and 2.13, the *mas* antagonist A779 was able to block the actions of ANG1-7 in both the MLE cell line and primary human AECs. This suggests that in alveolar epithelial cells, the ability of ANG1-7 to inhibit ANG II actions is mediated through the *mas* receptor, just as in other cell types studied thus far.

It is still unclear as to how intracellular signaling through the *mas* receptor can counteract the effects of ANG II. Evidence indicate that factors such as Akt phosphorylation, protein kinase C (PKC) activation and MAPK inhibition may play a role

in this pathway. In adult ventricular myocytes and endothelial cells, ANG 1-7 causes NO synthase phosphorylation and release through the phosphatidylinositol 3-kinase (PI3K)-protein kinase B (Akt)-dependent pathway (Sampaio et al., 2007; Dias-Peixoto et al., 2008). This was effectively prevented by the *mas* antagonist A779 or knockout of the *mas* gene. Moreover, it was found in rat extracardiac tissues that ANG 1-7 induces the *in vivo* phosphorylation of Akt by phosphorylating at threonine 308 and serine 473 (Muñoz et al., 2010). Studies with isolated perfused rat atria also showed ANG 1-7 to be exerting its effects through the *mas*/PI3K/Akt pathway (Shah et al., 2010). In addition, they also found ANG 1-7 to activate the Na⁺/H⁺ exchanger-1 and CaMKII through the *mas* receptor. However, little is known about how these mechanisms interact to antagonize the signaling of ANG II in the cell. It was previously shown that PKC expression is required for ANG II-stimulated AEC apoptosis in primary cultures (Papp et al., 2002). Recent evidence reveals that in renal tubule cells, ANG1-7 signals via the *mas* receptor to inhibit ANG II-stimulated Na⁺-ATPase activity by reversing PKC activation through a Gs/PKA pathway (Lara et al., 2010). Future studies focusing on the cAMP/PKA pathways in AECs might further elucidate the mechanism by which ANG1-7 inhibits ANG II signaling.

In summary, this study demonstrated that ANG1-7 acts through the receptor *mas* to block phosphorylation of JNK stimulated by either ANG II or bleomycin. ANG 1-7 treatment also inhibited bleomycin-induced apoptosis of alveolar epithelial cells, in agreement with the previously established requirement of JNK phosphorylation in AEC apoptosis. Together, the data demonstrates that ANG 1-7, produced from ANG II by ACE2, acts through the *mas* receptor to play a key role in the regulation of alveolar epithelial cell survival by preventing both JNK phosphorylation and subsequent apoptosis.

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