Role of integrin-mediated TGF β activation in the pathogenesis of pulmonary fibrosis

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Abstract

IPF (idiopathic pulmonary fibrosis) is a chronic progressive disease of unknown aetiology without effective treatment. IPF is characterized by excessive collagen deposition within the lung. Recent evidence suggests that the lung epithelium plays a key role in driving the fibrotic response. The current paradigm suggests that, after epithelial injury, there is impaired epithelial proliferation and enhanced epithelial apoptosis. This in turn promotes lung fibrosis through impaired basement membrane repair and increased epithelial-mesenchymal transition. Furthermore, fibroblasts are recruited to the wounded area and adopt a myofibroblast phenotype, with the up-regulation of matrix-synthesizing genes and down-regulation of matrix-degradation genes. There is compelling evidence that the cytokine TGF β (transforming growth factor β) plays a central role in this process. In normal lung, TGF β is maintained in an inactive state that is tightly regulated temporally and spatially. One of the major TGF β -activation pathways involves integrins, and the role of the $\alpha v\beta 6$ integrin has been particularly well described in the pathogenesis of IPF. Owing to the pleiotropic nature of TGF β , strategies that inhibit activation of TGF β in a cell- or disease-specific manner are attractive for the treatment of chronic fibrotic lung conditions. Therefore the molecular pathways that lead to integrin-mediated TGF β activation must be precisely defined to identify and fully exploit novel therapeutic targets that might ultimately improve the prognosis for patients with IPF.

Introduction

The prognosis of IPF (idiopathic pulmonary fibrosis) is poorer than some cancers, with 5-year survival rates of 43% [1] and a median survival of 2.4 years. Furthermore, the incidence of IPF, currently approx. 4500 new cases annually in the U.K., is rising [1]. The pathogenesis of IPF is incompletely understood, but the lung epithelium is thought to play a key role in orchestrating the fibrotic response [2]. The current paradigm suggests that, after lung injury, there is epithelial damage with subsequent destruction of the alveolar-capillary basement membrane (Figure 1). This permits fibrogenic cell infiltration of the alveolar interstitium, and excess abnormal matrix synthesis characteristic of pulmonary fibrosis. Failure of epithelial repair, through reduced epithelial proliferation and increased apoptosis [3], promotes fibrosis. Furthermore, it has been estimated that up to 30% of fibroblasts in fibrotic lung disease may be derived from epithelial cells [4], hence the epithelium may contribute to fibrosis through EMT (epithelial-mesenchymal transition). The epithelium may also promote fibroblast proliferation, collagen synthesis and myofibroblast transdifferentiation via the paracrine effects of growth factors such as TGF β (transforming growth factor β)

[5]. Indeed, TGF β may mediate many profibrotic effects in the alveolar epithelium following lung injury.

Given the lack of effective treatment for IPF, there is considerable work focused on identifying novel therapeutic targets through understanding the pathogenesis of IPF. The present review focuses on the pathways involved in integrinmediated TGF β activation in the pathogenesis of pulmonary fibrosis.

TGF β is a central mediator in the development of pulmonary fibrosis

TGF β is a member of the TGF β superfamily, a highly conserved group of cytokines including bone morphogenic proteins, activins and inhibins. There are three mammalian isoforms (TGF- β 1, - β 2 and - β 3), encoded by separate genes with distinct and related functions. The effects of TGF β 1 have been best characterized in pulmonary fibrosis, and from this point onwards, references to TGF β will relate to TGF β 1 unless specified otherwise.

TGF β is a pleiotropic cytokine that is ubiquitously expressed by all cells and tissues within the body. Studies of knockout mice have highlighted the crucial developmental functions of TGF β , which also plays a central role in a diverse range of processes, including wound healing, immunity and carcinogenesis. Its pleiotropic effects are notable as it can either inhibit or stimulate the same cellular processes in a cell-type- or tissue-specific manner.

TGF β has profound effects on epithelial cells and fibroblasts, which are central to the pathogenesis of pulmonary fibrosis. TGF β promotes epithelial cell apoptosis [6], EMT

Key words: epithelium, idiopathic pulmonary fibrosis, integrin, transforming growth factor- β (TGF β).

Abbreviations used: COPD, chronic obstructive pulmonary disease; EMT, epithelialmesenchymal transition; IPF, idiopathic pulmonary fibrosis; *itgb6*, $\beta6$ integrin subunit gene; LAP, latency-associated peptide; LPA, lysophosphatidic acid; LPAR2; LPA receptor 2; LTBP, latent transforming growth factor- β -binding protein; MMP, matrix metalloproteinase; PAR-1, proteaseactivated receptor-1; TGF β , transforming growth factor β .

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Figure 1 | Pathology of IPF

(a) Histological section of lung from patient with IPF stained with Masson's trichrome. Shows increased matrix (M) with bronchiolization of type 1 cells (B) and traction of alveoli. Alveoli contain proteinacious exudate and inflammatory cells including activated macrophages (AM). (b) Cartoon depicting histological features of IPF. Alveolar basement membrane is disrupted with type 2 cell hypertrophy and bronchiolization of type 1 cells. There is exudation of intravascular fluid with inflammatory and apoptotic cells within the alveoli. Beneath the epithelial cell layer there is provisional matrix deposition containing increased matrix, fibroblasts and myofibroblasts and platelets.



[7], epithelial cell migration [8], collagen synthesis, fibroblast proliferation and transdifferentiation into myofibroblasts [5]. Furthermore, the role of TGF β is well described in IPF. TGF β is increased in tissue samples from both animal models of IPF [9] and IPF patients [10]. Overexpression of an adenovirus encoding active TGF β leads to persistent pulmonary fibrosis [11], and inhibiting TGF β with soluble TGF β receptor [12], or a TGF β receptor 1 (ALK5) inhibitor [13], ameliorates pulmonary fibrosis. Furthermore, mice null for the TGF β signalling molecule Smad3 are protected from pulmonary fibrosis [14].

The crucial role of TGF β in the pathogenesis of IPF makes it an attractive therapeutic target. Unfortunately, its pleiotropic nature and roles in normal tissue homoeostasis may make global inhibition of TGF β problematic in the treatment of chronic diseases such as IPF. Pre-clinical and clinical studies to date have not found convincing evidence of toxicity associated with TGF β blockade, although there are concerns that long-term toxicity may be a serious potential pitfall of global TGF β inhibition [15,16]. Thus identifying and targeting tissue- or disease-specific mechanisms of TGF β activation are attractive alternatives for the treatment of IPF.

TGF β must be activated in order to have a biological effect

TGF β is secreted by most cells in association with the LTBPs (latent TGF β -binding proteins) as the large latent complex, which is sequestered in the matrix. TGF β itself is synthesized as the small latent complex consisting of active TGF β non-covalently associated with the LAP (latency-

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associated peptide). The tissue specificity of TGF β may be partially determined by the LTBPs, which bind TGF β in an isoform-specific fashion [17]. Furthermore, TGF β associated with the latent complex is quiescent while stored in the matrix. Therefore, for TGF β to exert any biological effects, it must be activated by dissociating from, or altering its interaction with, the LAP. Several processes can cause this, including physical processes such as temperature extremes, low pH and oxidation. TGF β can also be activated by a number of proteases, including plasmin, tryptase, thrombin, elastase, MMP (matrix metalloproteinase)-2 and MMP-9 [18– 21]. However, the importance of these mechanisms *in vivo* has yet to be defined. The best characterized mechanisms of *in vivo* TGF β activation are mediated by interactions with thrombospondin or integrins.

Integrin-mediated activation of TGF β

Integrins are heterodimeric transmembrane proteins consisting of α and β subunits. They are capable of binding to extracellular matrix proteins as well as a range of other molecules, including cell-surface ligands, transmembrane proteins, soluble proteases and growth factors [22].

The mammalian genome encodes 18 α subunits and eight β subunits that heterodimerize to form 24 $\alpha\beta$ integrin combinations. Eight integrins, including all five αv containing integrins, bind ligands through an RGD (Arg-Gly-Asp) sequence. An RGD sequence is found in the LAP of TGF β 1 and TGF β 3, which facilitates the activation of TGF β by at least four αv -containing integrins ($\alpha v\beta$ 3, $\alpha v\beta$ 5, $\alpha v\beta$ 6 and $\alpha v\beta$ 8) *in vitro*. The LAP of TGF β 2 does not contain an

Figure 2 | Two known mechanisms of integrin-mediated ${\rm TGF}\beta$ activation

(a) Latent TGF β binds the $\alpha \nu \beta 6$ and $\alpha \nu \beta 8$ integrins at the interface of the α and β subunit and the RGD motif of the TGF β LAP. The LAP is tethered to the matrix, or cell surface, through LTBP. (b) Activation of TGF β by the $\alpha \nu \beta 8$ integrin does not require association of the cytoplasmic domain and the cytoskeleton. The $\alpha \nu \beta 8$ integrin presents TGF β to MMP-14 which proteolytically digests the LAP and permits the active TGF β to diffuse freely. However, $\alpha \nu \beta 6$ integrin-mediated TGF β activation leads to structural changes within the latent TGF β complex mediated via traction induced on the tethered $\alpha \nu \beta 6$ integrin–LAP complex via the cytoskeleton. There is no release of free TGF β from the complex and therefore there is an absolute requirement for cell-cell contact between the $\alpha \nu \beta 6$ integrin-expressing cell and the TGF β -responsive cell.



RGD motif and no integrin-mediated activation of TGF β 2 has been described.

In vivo, activation of TGF β by integrins appears to play a major role during development and in various disease models. Integrins can activate TGF β via two main mechanisms (Figure 2): a protease-dependent mechanism ($\alpha\nu\beta$ 8) and a mechanism involving cell traction ($\alpha\nu\beta$ 3, $\alpha\nu\beta$ 5 and $\alpha\nu\beta$ 6).

Studies using genetically modified mice have demonstrated that integrins are important in the non-proteolytic activation of TGF β in the lung [23] and on dendritic cells [24]. The replacement of aspartic acid in the RGD motif with glutamic acid renders it unable to bind integrins. Mice bearing this mutation have a phenotype almost identical with TGF β 1-null mice [25], suggesting that integrins contribute significantly to the developmental effects of TGF β activation *in vivo*. Recent data using $\alpha v \beta 6$ (*itgb* $6^{-/-}$, where *itgb*6 is $\beta 6$ integrin subunit gene) and $\alpha v \beta 8$ (*itgb* $8^{-/-}$) -null mice have suggested that the phenotype associated with TGF β 1- and TGF β 3-null mice is due primarily to loss of these two integrins [26].

$\alpha v \beta 6$ integrin-mediated TGF β activation

The $\alpha\nu\beta6$ integrin was the first integrin to be implicated in the activation of TGF β [23]. Expression of $\alpha\nu\beta6$ is significantly increased in injured epithelia, but overexpression is not sufficient to promote fibrosis [27]. The $\alpha\nu\beta6$ integrin itself must be activated before TGF β activation can occur, and the

evidence suggests that this occurs in response to cytoskeletal changes.

The cytoplasmic tail of the β 6 subunit binds to the actin cytoskeleton, and mutation of this domain or treatment of cells with cytochalasin D, an inhibitor of actin polymerization, abolishes $\alpha\nu\beta6$ integrin-mediated TGF β activation [23]. Additionally, in cells lacking LTBP-1, $\alpha\nu\beta6$ cannot activate TGF β , therefore $\alpha\nu\beta6$ integrin-mediated TGF β activation is critically dependent on the association of latent TGF β with LTBP-1 within the large latent complex [28]. TGF β -activating activity can be restored by inducing the expression of a short fusion protein with regions mimicking the LAP-binding and extracellular matrix-binding domains of LTBP-1 [28]. Overall, these studies suggest that $\alpha\nu\beta6$ -mediated TGF β activation is dependent on tethering the $\alpha\nu\beta6$ integrin-TGF β complex to the cell surface.

The $\alpha v \beta 6$ integrin is constitutively bound to TGF β , suggesting that the system is primed to detect injurious stimuli. A pathway through which injurious signals are transmitted to the $\alpha v \beta 6$ integrin has been described [29,30]. Thrombin and LPA (lysophosphatidic acid) are G-proteincoupled receptor agonists released from platelets following injury. Both of these substances have been shown to promote $\alpha v \beta 6$ integrin-mediated TGF β activation and are implicated in the development of IPF [31,32]. Thrombin and LPA induce $\alpha v \beta 6$ integrin-mediated TGF β activation via PAR1 (protease-activated receptor-1) and LPAR2 (LPA receptor 2) respectively, which in turn induce cytoskeletal changes via the G-protein $G_{\alpha q}$, RhoA and Rho kinase [30]. Therefore inflammatory mediators released in response to tissue injury activate TGF β , by binding to epithelial cell-surface receptors and inducing cytoskeletal contraction, resulting in a conformational change in the $\alpha v\beta 6$ integrin– latent TGF β complex.

In vivo data have demonstrated the importance of $\alpha\nu\beta6$ integrin-mediated TGF β activation in the pathogenesis of IPF. Mice that do not express the $\beta6$ subunit ($itgb6^{-/-}$ mice) cannot form the $\alpha\nu\beta6$ integrin. These mice develop mild inflammation in the lungs and skin, but are protected from bleomycin-, LPS (lipopolysaccharide)- and ventilatorassociated lung injury, as well as bleomycin-induced pulmonary fibrosis and renal fibrosis [23,29,33]. Additionally, an anti- $\alpha\nu\beta6$ monoclonal antibody has been shown to prevent pulmonary fibrosis, renal fibrosis and acute lung injury *in vivo* [33–36].

Expression of *itgb6* is increased at the mRNA level in bleomycin-induced lung fibrosis [30] and at the protein level in IPF patients [30,35]. This suggests that, although increased $\alpha\nu\beta6$ integrin expression is insufficient to increase TGF β activation, abnormal regulation of the $\alpha\nu\beta6$ subunit may participate in the pathogenesis of IPF. Indeed, TGF β itself up-regulates *itgb6* expression [37], possibly mediated via the *Ets1* transcription factor [38]. This process can be inhibited by blocking TGF β or the $\alpha\nu\beta6$ integrin, suggesting a selfamplifying paracrine loop [39]. Thus it is possible that lung injury promotes $\alpha\nu\beta6$ integrin-mediated TGF β activation, which in turn amplifies the signal through increasing *itgb6*

Figure 3 | Possible pathogenic mechanism of IPF

Initial alveolar injury leads to release of platelet-derived mediators, including thrombin and LPA. These activate their respective G-protein-coupled receptors, PAR1 and LPAR2, and induce a signalling cascade involving $G_{\alpha q}$, RhoA and Rho kinase (ROCK), leading to actin polymerization and cellular traction on the cytoplasmic domain of the $\alpha v \beta 6$ integrin, permitting activation of matrix- or cell-associated TGF β (the initiating pathway). Active TGF β acts in a paracrine fashion on neighbouring cells, leading to receptor Smad phosphorylation and translocation and increased synthesis of the *itgb6* gene in epithelial cells (the amplification pathway). Owing to the disruption of the basement membrane, cell-cell contact is possible between epithelial cells and fibroblasts enabling epithelial cell induction of fibroblast- and myofibroblast-derived matrix proteins (the fibroqenic pathway).



gene expression (Figure 3). Understanding this process could identify new therapeutic targets for IPF.

$\alpha v \beta 8$ integrin-mediated TGF β activation

The $\alpha\nu\beta$ 8 integrin, like $\alpha\nu\beta$ 6, can bind and activate TGF β . However, the mechanism of TGF β activation is distinct from that of $\alpha\nu\beta$ 6. Whereas $\alpha\nu\beta$ 6-mediated TGF β activation relies upon actin polymerization and retains active TGF β at the cell surface, $\alpha\nu\beta$ 8 integrin-mediated TGF β activation does not require the cytoplasmic domain of the β subunit, and is therefore not influenced by cytoskeletal contraction.

It seems that $\alpha \nu \beta 8$ activates TGF β by acting as a cellsurface shuttle, presenting TGF β to a cell-surface protease. The $\alpha \nu \beta 8$ integrin co-localizes at the cell surface with MMP-14, also known as MT1-MMP (membrane-type 1 matrix metalloproteinase). Consequently, latent TGF β is presented to MMP-14 by the $\alpha \nu \beta 8$ integrin, which results in the proteolytic cleavage of the LAP and the release of active TGF β [40].

The $\alpha v \beta 8$ integrin is crucial for vascular and brain development and for suppression of the adaptive immune system [24,41]. In the lung, $\alpha v \beta 8$ integrin-mediated TGF β activation has been shown to delay epithelial wound closure and inhibit bronchial epithelial proliferation *in vitro* [42]. The importance of $\alpha v \beta 8$ -mediated TGF β activation *in vivo* is supported by the development of autoimmune colitis in mice with conditional loss of the $\alpha v\beta 8$ integrin in dendritic cells [24].

Although there is currently no evidence for the involvement of the $\alpha\nu\beta8$ integrin in the pathogenesis of pulmonary fibrosis, it may play an important role in the airway fibrosis seen in COPD (chronic obstructive pulmonary disease) [39]. Squamous metaplasia, part of the pathological process associated with COPD, is driven by $\alpha\nu\beta6$ integrin-mediated TGF β activation. This activates a paracrine loop involving IL (interleukin)-1 β -induced $\alpha\nu\beta8$ integrin expression on fibroblasts, resulting in TGF β activation and subsequent $\alpha\nu\beta6$ integrin up-regulation and increased squamous metaplasia [39]. Similar pathways could be active in the pathogenesis of IPF, which requires further study.

$\alpha v\beta 3$ and $\alpha v\beta 5$ integrin-mediated TGF β activation

Both the $\alpha v\beta 3$ and the $\alpha v\beta 5$ integrins are up-regulated in the dermal epithelium in systemic sclerosis. Both can activate TGF β in scleroderma fibroblasts and this is associated with the transition of fibroblasts into fibrogenic myofibroblasts [43-45]. Activation of TGF β by the $\alpha v \beta 5$ integrin is not inhibited by protease inhibitors [43], and cellular traction has been proposed to induce TGF β activation via $\alpha v \beta 3$ and $\alpha v \beta 5$ integrins. Myofibroblasts can liberate and activate TGF β from extracellular stores by transmitting a contractile force via the $\alpha v\beta 5$ integrin to latent TGF β [46]. Functionblocking antibodies against $\alpha v\beta 5$ inhibited this contractionmediated TGF β activation, which also occurred to a lesser extent with $\beta 1$ and $\alpha v \beta 3$ integrins [46]. This integrinmediated TGF β activation was limited to culture substrates with stiffness comparable with fibrotic tissue, suggesting that this mechanism maintains $TGF\beta$ activation after fibrosis has begun. This is the first direct evidence that mechanical stress can activate an extracellular matrix-bound cytokine [46].

However, there is currently no direct evidence for $\alpha\nu\beta\beta$ and $\alpha\nu\beta5$ integrin-mediated TGF β activation *in vivo*. In mice null for the $\beta\beta$ integrin subunit [47], the $\beta5$ subunit [48] or both [49], there is no apparent loss of TGF β activity. Furthermore, recent data have demonstrated that many of the developmental and immune effects of TGF β are mediated via the $\alpha\nu\beta6$ and $\alpha\nu\beta8$ integrins [26]. The role of TGF β activation by the $\alpha\nu\beta3$ and $\alpha\nu\beta5$ integrins thus needs to be clarified in disease models.

Conclusions

TGF β is a key cytokine in the pathogenesis of IPF. It is stored as an inactive molecule in the extracellular matrix and is primed for activation following injury. Our understanding of the mechanisms of integrin-mediated TGF β activation has increased dramatically in recent years. The $\alpha v\beta 3$, $\alpha v\beta 5$, $\alpha v\beta 6$ and $\alpha v\beta 8$ integrins have all been shown to activate TGF β *in vitro* and could theoretically influence the pathogenesis of IPF. However, the $\alpha v\beta 6$ integrin has been studied the most extensively, and is the most highly implicated in the pathogenesis of IPF.

As integrins can activate TGF β , and thus promote IPF, they are attractive therapeutic targets, not least because they are spatially and temporally restricted. The $\alpha\nu\beta6$ integrin is particularly attractive in this regard, because it is restricted to epithelial tissues, and inhibition of $\alpha\nu\beta6$ integrins would minimize disruption to the crucial homoeostatic functions of TGF β . Because IPF is a disease characterized by the presence of activated fibroblasts and myofibroblasts, which do not express the $\alpha\nu\beta6$ integrin, it has been suggested that therapies targeting multiple integrins simultaneously may be superior [50]. Additionally, given that proteases can act with integrins in TGF β activation, protease inhibitors themselves are worthy of further investigation. Ultimately, given the complex nature and the unpredictable course of IPF, combinations of these therapeutics could be necessary.

Finally, the molecular pathways leading to integrinmediated TGF β activation and the self-amplifying pathways that increase integrin gene expression are still incompletely understood. Dissecting these molecular pathways, and identifying where dysregulation occurs during the pathogenesis of IPF should help us to better understand the disease as well as identify novel therapeutic strategies for the treatment of IPF, which may hopefully improve the prognosis of this devastating disease.

Acknowledgements

We thank Dr Mark Travis (University of Manchester, Manchester, U.K.) and Dr Dean Sheppard (University of California San Francisco, San Francisco, CA, U.S.A.) for their helpful comments during the preparation of the paper.

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Received 28 April 2009 doi:10.1042/BST0370849