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MTOR SIGNALLING IS AN ESSENTIAL PATHWAY FOR TGF- β 1 INDUCED α SMA AND COLLAGEN GENE EXPRESSION

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Emerging evidence suggests a role for PI3K/mTOR signalling in the pathology of organ fibrosis. The aim of this study was to delineate PI3K/mTOR signalling in response to TGF- β_1 stimulation of primary human lung fibroblasts (HLFs), and to investigate the role of this pathway in TGF- β_1 mediated myofibroblast differentiation and collagen synthesis.

A time-course of SMAD 2/3 and Akt (Ser473) phosphorylation, the major downstream effector of the PI3K/mTOR pathway, was constructed to assess TGF- β_1 induced signalling kinetics in HLFs. TGF- β_1 (1 ng/ml) induced rapid phosphorylation of SMAD2/3, peaking at 1 h, followed by Akt phosphorylation which peaked 12 h after initial stimulation. Maximal expression of *ACTA2* and *COL1A1* was observed 36 h after TGF- β_1 stimulation, correlating with the delayed time-course of Akt phosphorylation.

To investigate the role of the PI3K/mTOR pathway in TGF- β_1 induced myofibroblast differentiation and collagen gene expression, HLFs were treated with pharmacological titrations of potent pathway inhibitors. Maximal Akt signalling and expression of *ACTA2* and *COL1A1* were significantly inhibited by a dual PI3K/mTOR inhibitor, while SMAD phosphorylation was unaffected. Treatment with an ATP competitive mTOR inhibitor also resulted in significantly reduced Akt phosphorylation and expression of *ACTA2* and *COL1A1*, in response to TGF- β_1 . In contrast, treatment of HLFs with either an allosteric or ATP competitive Akt inhibitor showed no inhibitory effect on TGF- β_1 induced gene expression.

These data suggest PI3 kinase/mTOR signalling is an important component in TGF- β_1 induced α SMA and collagen gene expression and that an mTOR dependent, Akt independent pathway mediates this functional response in primary HLFs.

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VASCULAR ENDOTHELIAL GROWTH FACTOR (VEGF) EXPRESSION IN THE IPF LUNG – A ROLE FOR ANTI-ANGIOGENIC ISOFORMS?

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Introduction VEGF has been implicated in the pathogenesis of IPF. Differential splicing of the VEGF gene produces an alternative family of isoforms (VEGF $_{xxx}$ b) that have anti-angiogenic properties, in contrast to conventional isoforms (VEGF $_{xxx}$). Currently available literature on the role of VEGF in IPF has not differentiated between these families of isoforms and thus a degree of literature re-appraisal is required.

Hypotheses

- The balance of VEGF_{xxx}:VEGF_{xxx}b isoforms may be important in IPF pathogenesis
- VEGF_{xxx}b isoforms may abrogate the development of IPF



Abstract S137 Figure 1 The effect of VEGF₁₆₅b on the development of murine BLM-induced pulmonary fibrosis

Bleomycin was given to WT type mice (n = 6 per group) by oro-pharyngeal aspiration (Day 0). rhVEGF $_{165}$ b was administered by IP injection (1 μ g per mouse, bi-weekly) from days 10 to 21. Fibrosis was scored and examined histologically by Masson's trichrome staining. The development of BLM-induced fibrosis was attenuated in mice receiving rhVEGF $_{165}$ b

Methods Human lung sections and BALF were used to quantify isoform expression in the IPF lung and were compared to controls (ELISA and IHC). Explanted 'normal' (NF) and 'fibrotic' (FF) fibroblasts were grown in culture with subsequent total RNA and cell lysate extraction (qPCR and WB). Wild-type mice were administered bleomycin (BLM) then received bi-weekly therapeutic intraperitoneal (IP) injections of rhVEGF₁₆₅b (from day 10). Fibrosis was assessed histologically (Masson's Trichrome and Lung fibrosis score).

Results In the IPF lung, the alveolar epithelium was the most prominent site for total VEGF (PanVEGF isoforms) but also for VEGF $_{165}$ b (n = 10). Addiitonal staining was noted in fibroblasts and lung inflammatory cells. Alveolar and fibrotic cells in the least fibrotic areas of the IPF lung expressed significantly less VEGF $_{165}$ b than severely fibrotic areas (p < 0.001, n = 10). Examination of IPF BALF by ELISA revealed that total VEGF expression was significantly lower compared to control (IPF: 18.04 pg/ml +/- 6.13 n = 15, CTRL 85.72 pg/ml +/- 17.08 n = 13), whilst VEGF $_{165}$ b could not be detected in identical samples.

Explanted NF and FF express comparable quantities of VEGF $_{xxx}$ and VEGF $_{xxx}$ b isoforms at the mRNA and protein level. Rh VEGF $_{165}$ increases the mRNA expression of fibronectin (p < 0.001, n = 4) an effect not seen following the administration of rhVEGF $_{165}$ b.

Administration of rhVEGF $_{165}$ b to mice attenuated the development of BLM-induced pulmonary fibrosis (Masson's Trichrome (Figure 1) and lung fibrosis score (mean score: BLM alone 41.20 vs VEGF $_{165}$ b 30.67, p < 0.01, n = 6 per group)). Conclusion Differential expression of VEGF $_{xxx}$ and VEGF $_{xxx}$ b isoforms occurs in the IPF lung. *In vitro*, recombinant proteins appear to have differential effects on ECM synthesis and *in vivo* attenuate the formation of pulmonary fibrosis. A mouse overexpressing VEGF $_{165}$ b in the lung has been developed to study this concept in greater detail.

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NANODIAMOND DELIVERY OF VASCULAR ENDOTHELIAL GROWTH FACTOR PROMOTES FETAL LUNG DEVELOPMENT IN A RAT MODEL OF CONGENITAL DIAPHRAGMATIC HERNIA

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