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Increased survivin expression contributes to apoptosisresistance in IPF fibroblasts

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Abstract

Fibroblasts perform critical functions during the normal host response to tissue injury, but the inappropriate accumulation and persistent activation of these cells results in the development of tissue fibrosis. The mechanisms accounting for the aberrant accumulation of fibroblasts during fibrotic repair are poorly understood, although evidence supports a role for fibroblast resistance to apoptosis as a contributing factor. We have shown that TGF-B1 and endothelin-1 (ET-1), soluble mediators implicated in fibrogenesis, promote fibroblast resistance to apoptosis. Moreover, we recently found that ET-1 induced apoptosis resistance in normal lung fibroblasts through the upregulation of survivin, a member of the Inhibitor of Apoptosis (IAP) protein family. In the current study, we sought to determine the role of survivin in the apoptosis resistance of primary fibroblasts isolated from the lungs of patients with Idiopathic Pulmonary Fibrosis (IPF), a fibrotic lung disease of unclear etiology for which there is no definitive therapy. First, we examined survivin expression in lung tissue from patients with IPF and found that there is robust expression in the fibroblasts residing within fibroblastic foci (the "active" lesions in IPF which correlate with mortality). Next, we show that survivin expression is increased in fibroblasts isolated from IPF lung tissue compared to cells from normal lung tissue. Consistent with a role in fibrogenesis, we demonstrate that TGF- β 1 increases survivin expression in normal lung fibroblasts. Finally, we show that inhibition of survivin enhances susceptibility of a subset of IPF fibroblasts to apoptosis. Collectively, these findings suggest that increased survivin expression represents one mechanism contributing an apoptosis-resistant phenotype in IPF fibroblasts.

Keywords

Myofibroblast; Idiopathic Pulmonary Fibrosis; Inhibitor of Apoptosis; Lung Fibrosis; Fas

1. INTRODUCTION

Idiopathic Pulmonary Fibrosis (IPF) is a chronic lung disease characterized by the progressive development of scar tissue which can ultimately lead to respiratory failure and death. The incidence of IPF is increasing and, with an average survival of 2 - 3 years

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following the diagnosis, the overall prognosis is poor [1–3]. Although several randomized clinical trials have evaluated a number of therapeutic targets in IPF over the last decade, there continues to be a lack of clearly effective pharmacologic therapy [4]. Despite the generally poor prognosis, a well-recognized clinical feature of IPF is the markedly heterogeneous clinical course, with some patients maintaining stable lung function for years while others experience a rapid deterioration culminating in respiratory failure and death within months of the diagnosis [5]. It is critical to better understand the mechanisms of biologic heterogeneity that underlie the variable clinical course of IPF in order to identify novel targets for intervention that can be applied in a personalized manner [6].

Alveolar scarring in IPF is hypothesized to result from dysfunctional wound repair which is characterized by the accumulation of mesenchymal cells (including fibroblasts and their differentiated phenotype, myofibroblasts) within pathologic lesions called fibroblastic foci [7,8]. During normal reparative responses following injury, mesenchymal cells perform critical functions which provide healing wounds with tensile strength and facilitate the homeostatic restoration of epithelial integrity, including the synthesis, secretion, organization and contraction of new extracellular matrix (ECM), [1,9–11]. Resolution of the normal repair process requires clearance of fibroblasts via apoptosis [12,13]. However, in the dysfunctional repair that is evident in IPF and other diseases characterized by fibrosis, the persistence of reparative fibroblasts leads to excessive deposition and organization of ECM [14]. Several recent studies demonstrate a paucity of apoptosis within the mesenchymal cells that comprise the fibroblastic foci in IPF, highlighting one mechanism that contributes to the persistence of these cells [15–18]. Accumulating studies also show that fibrotic lung fibroblasts are resistant to a variety of apoptotic stimuli, further supporting a role for impaired fibroblast apoptosis in the pathobiology of IPF [18-25]. However, the mechanisms regulating resistance to apoptosis in lung fibroblasts have not been fully elucidated [14].

Recent studies suggest that Inhibitor of Apoptosis (IAP) family proteins can regulate susceptibility to apoptosis in normal lung fibroblasts [18,20,24]. The IAP genes encode at least eight different proteins with diverse functions, including the direct and indirect regulation of apoptosis. The protein family is characterized by the presence of at least one baculoviral IAP repeat (BIR) domain which is critical for the ability of these proteins to inhibit caspase activation in vitro [26,27]. The smallest member of the IAP family, survivin (also known as BIRC5) is a 16.3 kD protein with one BIR domain that is predominantly expressed as a homodimer. Physiologically, survivin serves a critical function in cell-cycle progression and is essential during embryonic development. In adult cells, however, it is typically expressed only in actively dividing cells [28]. As with other IAP family members, survivin inhibits activation of caspase 3, 7 and 9 in-vitro. Moreover, a number of studies demonstrate regulation of cell susceptibility to apoptosis by survivin, although the precise in vivo mechanisms remain unclear [27]. In accord with its crucial function in cell division and apoptosis regulation, inappropriate survivin expression is associated with cancers involving multiple organs including lung, colon, breast, brain and skin [27,29,30]. We have recently shown that induction of survivin by the soluble fibrogenic mediator endothelin-1 (ET-1) reduces the apoptotic susceptibility of normal fetal lung fibroblasts and that the anti-fibrotic mediator prostaglandin E2 enhances fibroblast susceptibility to apoptosis while suppressing survivin expression [20,24]. The goal of this study was to examine the expression of survivin in IPF lung fibroblasts and to determine if survivin regulates IPF fibroblast susceptibility to Fas-mediated apoptosis.

2. MATERIALS AND METHODS

2.1. Cells and Cell Culture

Normal human adult lung fibroblasts (CCL-210) were obtained from ATCC (Manassas, VA). Cells between passages 8 and 16 were cultured as previously described and growth arrested for 24 hours in serum-free media prior to experiments [20]. Patient-derived primary human lung fibroblasts were cultured from the lungs of patients with IPF or from normal lung tissue from patients undergoing thoracic surgery for non-fibrotic disease as we have described previously [31]. Written informed consent was obtained from all subjects in accordance with the University of Michigan Institutional Review Board.

2.2. Reagents

Porcine TGF-β1 was from R&D Systems (Minneapolis, MN). The activating anti-Fas antibody (clone CH11) was from Millipore (Billerica, MA). The mouse monoclonal antibody to survivin used for Western blotting was from Abnova. Antibodies to glyceraldehyde-3-phosphate dehydrogenase (GAPDH) were from Cell Signaling (Danvers, MA). The Cell Death Detection ELISA Kit detecting histone-associated DNA fragments was from Roche Applied Science (Indianapolis, IN). Horseradish peroxidase conjugated secondary antibodies were from Pierce (Rockford, IL). The survivin inhibitor CAY10625, which disrupts the interaction between survivin and Smac/Diablo (second mitochondrial activator of caspases/direct IAP binding protein with low isoelectric point), thereby allowing Smac/Diablo to carry out its pro-apoptotic function, was from Cayman Chemical Company (Ann Arbor, MI) [32]. YM155, a functionally distinct inhibitor which inhibits survivin transcription [33], was from Selleck Chemicals (Houston, TX).

2.3. Immunohistochemistry

4 µm sections on polylysiene coated slides (VWR, Leicestershire, UK) were de-waxed and rehydrated by immersion in xylene followed by decreasing concentrations of ethanol. Sections were washed with tris-buffered saline (TBS, pH 7.6), and endogenous peroxidise activity was blocked by immersion in 3% H₂O₂ (v/v) for 30 minutes at room temperature. Sections were washed twice in TBS and treated with a 1 in 6 solution of goat serum containing 4 drops/ml of Avidin block (Vector Laboratories, Peterborough, UK) for 20 minutes and incubated overnight with rabbit anti-survivin polyclonal antibodies (4 µg/ml; Santa Cruz Biotechnology) or an isotype control in TBS with 1% bovine serum albumin and 4 drops/ml of Biotin block. Following another wash, samples were treated with a 1:200 dilution of biotinylated seconddary antibody (Dako, Cambridshire, UK), washed again and immersed in a 1:200 dilution of Streptavadin HRP (Dako) for 30 minutes at room temperature. After washing, DAB peroxidise substrate (Vector Laboratories) was added for ten minutes. Samples were rinsed in H₂O, counterstained with Gill's Haematoxylin, dehydrated in Xylene and mounted using a Sakura Coveraid system.

2.4. Quantitative Reverse Transcription Real-Time PCR

Quantitative reverse transcription real time-PCR was performed on an Applied Biosystems (Foster City, CA) 7300 real time PCR machine as we have reported previously [31]. Relative quantitation was based on the $\Delta\Delta C_T$ method and used predesigned primer-probe sets obtained from Applied Biosystems TaqMan Gene Expression Assays for human Survivin (BIRC5; product number Hs00153353_m1).

2.5. SDS Page Electrophoresis and Western Blotting

Whole cell lysates were collected and subjected to SDS-PAGE electrophoresis and Western blotting as previously described [20]. All Western blots were stripped and re-probed for GAPDH.

2.6. Densitometry

Western blots were analyzed using the public domain NIH ImageJ program available at http://rsbweb.nih.gov/ij/ as previously described [20]. The relative expression of survivin was determined by dividing the survivin band density (adjusted for the background) by the density of the corresponding GAPDH. This ratio of survivin to GAPDH was then normalized by dividing these values by the average ratio in the "normal" fibroblasts, such that the average of all normal (or untreated) lung fibroblasts is equal to 1.0 and the IPF samples (or the TGF- β 1 treated samples) represent the fold change

2.7. Apoptosis

Apoptosis was assessed by ELISA-based detection of histone-associated DNA fragments using the Cell Death Detection ELISA Kit (Roche Applied Science, Indianapolis, IN) as previously described [20]. To allow comparisons across experiments using different IPF fibroblast cell lines, the apoptosis data is represented as the percentage of the absorbance of the positive control sample (provided by the manufacturer) that was run on each ELISA plate. Each IPF sample was run in triplicate.

2.8. Statistical Analyses

Statistical analyses were done using ANOVA with Tukey post-test for experiments involving multiple comparisons (apoptosis assays). An unpaired T-test with Welch's correction was used for comparisons between normal and IPF samples (Graphpad Prism software version 5.01 for Windows). The relationship between apoptosis induced by Fasactivating ligand combined with YM155 or CAY10625 was determined using a Spearman correlation (r).

3. RESULTS

3.1. Survivin Is Expressed in IPF Lung Tissue

To determine if survivin was increased in fibroblasts residing in areas of active fibrosis, we used immunohistochemistry to examine survivin expression in normal and IPF lung biopsies (Figure 1). In normal lung tissue, survivin staining was observed in subsets of alveolar epithelial cells (Figure 1(a), arrows) and in the normal bronchial epithelium (data not shown). In IPF tissue, survivin expression was identified in interstitial cells within thickened alveolar septae (Figure 1(b), arrowhead) and in the alveolar epithelium (Figure 1(b), arrow). Marked survivin expression was also observed in mesenchymal cells residing within the IPF fibroblastic foci and in the epithelium overlying the fibroblastic foci (Figure 1(c)).

3.2. Survivin Expression Is Increased in IPF Lung Fibroblasts

Having shown expression of survivin within fibroblastic foci of IPF tissue, we next sought to compare survivin expression in primary lung fibroblasts from patients with normal lung architecture to fibroblasts isolated from IPF lung tissue. Using quantitative real-time PCR of banked RNA from 12 patient-derived normal lung fibroblast cell lines and 21 IPF fibroblast cell lines, we found a signifycant (2-fold) increase in survivin expression in the IPF fibroblasts (Figure 2). While the overall difference was statistically significant, the distribution of survivin expression in IPF fibroblasts was heterogeneous and appeared to segregate into two distinct groups, with approximately half (12/21) of the IPF cell lines

We next assessed the relative survivin protein levels in 12 IPF fibroblast cell lines compared to 5 normal lung fibroblast cell lines (Figure 3). Consistent with the RNA findings, survivin expression was significantly (approximately 2.5-fold) higher in the IPF fibroblasts. Survivin protein expression in IPF fibroblasts was also heterogeneous, with some cell lines indistinguishable from the average expression in normal lung fibroblasts (white diamonds), and others demonstrating more than a 3-fold increase in survivin expression (black diamonds).

3.3. TGF-_{β1} Increases Survivin Expression in Normal Fibroblasts

TGF- β 1 is strongly implicated in the pathogenesis of fibrosis and has been shown to promote myofibroblast differentiation, ECM deposition, and the induction of an apoptosisresistant phenotype [22,23,34]. To determine if TGF- β 1 enhanced survivin expression in normal lung fibroblasts, we treated CCL-210 (normal adult lung) fibroblasts with TGF- β 1 and assessed survivin expression by Western blotting. Consistent with the induction of an apoptosis-resistant phenotype in normal fibroblasts, TGF- β 1 induced a robust increase in survivin expression within six hours of treatment (Figure 4). A similar response to TGF- β 1 was observed in IMR-90 fibroblasts (data not shown). In combination with our prior study showing induction of survivin by ET-1, these findings suggest that survivin induction represents a conserved response of normal lung fibroblasts to soluble pro-fibrotic mediators [20].

3.4. Survivin Inhibition Enhances IPF Fibroblast Susceptibility to Apoptosis

In normal lung fibroblasts, enhanced survivin expression by ET-1 is associated with decreased susceptibility to apoptosis [20]. In contrast, the suppression of survivin in response to treatment with PGE_2 is associated with enhanced susceptibility to apoptosis [20,24]. Finding that survivin is expressed within the fibroblastic foci in IPF tissue, that IPF fibroblasts demonstrate increased levels of survivin, and that TGF-B1 induced survivin expression in normal lung fibroblasts, we sought to determine if survivin inhibition could modulate susceptibility to apoptosis in IPF fibroblasts. Eight patient derived IPF fibroblast cell lines were treated with/without Fas-activating antibody in the presence/absence of two mechanistically distinct inhibitors of survivin (CAY10625 or YM155) and apoptosis was assessed at 16 hours (Figure 5). Consistent with prior reports, treatment with Fas-activating ligand alone did not significantly increase IPF lung fibroblast apoptosis [18,34,35]. However, treatment with the Fas-activating antibodies in the presence of either survivin inhibitor was associated with a statistically significant increase in IPF fibroblast apoptosis. As with survivin expression levels, the IPF fibroblast cell lines demonstrated heterogeneity in their apoptotic responses to Fas-activation in the presence of the survivin inhibittors, with enhanced apoptosis observed in only half of the IPF fibroblast cell lines. Neither inhibitor significantly increased apoptosis in the absence of Fas-activateing antibodies. Finally, there was strong correlation between apoptosis induced by Fas-activation with YM155 and Fasactivation with CAY1062 (Figure 5(b)), suggesting that the increased susceptibility to apoptosis was specifically related to survivin inhibition and not the result of an off-target effect of the inhibitors used.

4. DISCUSSION

In the current study, we examined the role of survivin in the regulation of IPF lung fibroblast apoptosis. We show that survivin expression is increased in the mesenchymal cells within the fibroblastic foci of IPF lungs, supporting a role for this IAP in the apoptosis-resistant phenotype of these cells. Further, we demonstrate increased survivin expression in IPF lung fibroblasts compared to normal lung fibroblasts, although the expression of survivin in these IPF fibroblasts is heterogeneous. Additionally, we show that TGF- β 1, like ET-1, induces survivin expression in normal lung fibroblasts [20]. Finally, we demonstrate that pharmacologic inhibition of survivin enhances the apoptotic susceptibility of a subset of IPF lung fibroblasts. Collectively, these findings indicate that increased survivin expression represents one mechanism by which IPF fibroblasts acquire resistance to apoptosis.

IPF lung tissue is characterized by the heterogeneous distribution of fibroblastic foci composed of fibroblasts and myofibroblasts which, in contrast to the overlying epithelium, lack evidence of active apoptosis [2,14,15, 17,18]. The mechanisms leading to the lack of apoptosis in IPF fibroblasts have not been fully elucidated. Given the heterogeneity seen in IPF, it is possible that different stimuli converge on overlapping signaling pathways and effector mechanisms to promote pro-fibrotic fibroblast phenotypes in different patients. Supporting this concept of mechanistic heterogeneity with phenotypic convergence in complex disease states such as fibrosis, we have shown that TGF- β 1 and ET-1, two soluble mediators implicated in fibrosis *in-vivo*, independently promote fibroblast resistance to apoptosis through activation of the pro-survival tyrosine kinases, focal adhesion kinase (FAK) and phosphatidylinositol 3-hydroxy kinase/AKT [23,34].

We recently demonstrated a mechanistic role for survivin in the regulation of normal lung fibroblast apoptosis, showing that FAK and PI3K/AKT activation are required for the upregulation of survivin by ET-1 and that survivin inhibition attenuates ET-1 mediated resistance to apoptosis in these cells [20]. In this study, we extend these findings into primary fibrotic lung fibroblasts, showing that a significant subset of IPF fibroblast cell lines have markedly increased levels of survivin. Moreover, while all of the IPF lung fibroblast cell lines studied were resistant to Fas-mediated apoptosis, survivin inhibition enhanced apoptosis in only one-half of the cell lines. The implication of this observation is that alternative mechanisms that are not dependent on survivin mediate apoptosis resistance in the other half of the IPF fibroblast cell lines. Such segregation of IPF fibroblasts into similar apoptosis-resistant phenotypes via heterogeneous mechanisms is reminiscent of a prior study demonstrating that diminished expression of the EP2 receptor accounted for the lack of IPF fibroblast responsiveness to the anti-fibrotic mediator PGE_2 in 4/9 cell lines while the remaining IPF fibroblasts had decreased responsiveness to PGE₂ despite normal levels of the EP2 receptor [36]. Interestingly, we have also shown that PGE₂ suppresses survivin expression in normal lung fibroblasts, suggesting a potential mechanistic link between PGE₂ resistance and increased survivin expression in the IPF fibroblasts.

Although the mechanisms by which survivin regulates cell apoptosis are controversial, there is abundant evidence demonstrating that survivin inhibition can enhance the susceptibility of cancer cells to apoptotic stimuli, making survivin an attractive target for the anti-cancer treatments [27,30,33,37–39]. We employed two mechanistically distinct inhibitors to assess the role of survivin in IPF fibroblast resistance to Fas-mediated apoptosis [32, 33,40]. Similar to the distribution of survivin expression in IPF fibroblasts, inhibition of survivin significantly enhanced Fas-mediated apoptosis in approximately half of the IPF fibroblasts studied. While all pharmacologic inhibitors can have off target effects, we found a high correlation between the effects of YM155 and CAY10625 on cell susceptibility to apoptosis (r = 0.95, p = 0.002). This high correlation between the effects of two mechanistically

distinct inhibitors, combined with the lack of any significant effect on apoptosis when cells were treated with the inhibitors alone, suggests that the enhanced apoptosis was specific to the inhibition of survivin and did not represent an off-target effect of either inhibitor.

Studies exploring a relationship between survivin and fibrosis are lacking. Our previous in vitro studies support a link between survivin and a pro-fibrotic fibroblast phenotype [20,36]. Additionally, several studies have linked survivin expression with PI3K/AKT activation, findings consistent with the established roles of PI3K/AKT signaling in pulmonary fibrosis [18,41–44]. *In vivo*, only one prior study has shown a link between survivin and fibrosis [45]. In that study, which employed a murine model of reversible liver fibrosis, survivin expression was significantly increased during the establishment of fibrosis. Moreover, the resolution phase of this fibrosis model, which is dependent on hepatic stellate cell (myofibroblast) apoptosis, coincided with the reduction of survivin expression to pre-fibrosis levels. However, a mechanistic link between survivin and fibrosis was not examined, and no reports have shown that survivin is required for the establishment of fibrosis. Together with the current demonstration that inhibition of survivin can restore susceptibility of IPF fibroblasts to Fas-mediated apoptosis, these findings suggest that survivin may represent a potential target for anti-fibrotic therapies.

In conclusion, this study is the first to show that survivin is expressed in the fibroblastic foci of IPF tissue. Moreover, it is the first study to examine survivin expression in IPF fibroblasts and to assess the impact of survivin inhibition on fibroblast susceptibility to apoptosis. Our findings demonstrate that survivin expression is increased in a subset of IPF fibroblasts and that it is possible to enhance the susceptibility of these cells to Fas-mediated apoptosis by inhibiting survivin. The heterogeneity of survivin expression in IPF fibroblasts, and the heterogeneous responses to apoptotic stimuli seen in the presence of survivin inhibitors, may be a reflection of the biologic and clinical heterogeneity observed in patients with IPF. Further studies are required to elucidate the mechanisms by which survivin regulates apoptosis susceptibility in these cells, and to determine if survivin inhibition can modulate fibrosis *in vivo*.

Acknowledgments

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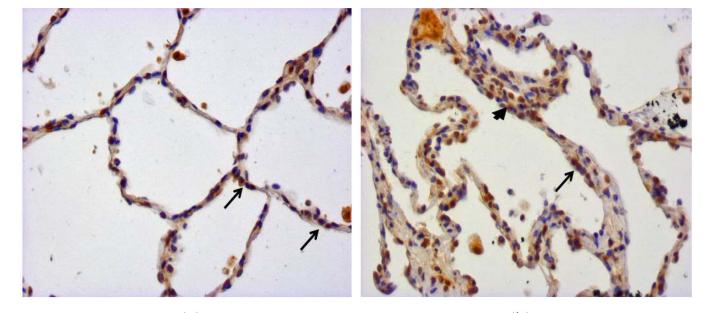
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Sisson et al.



(b)

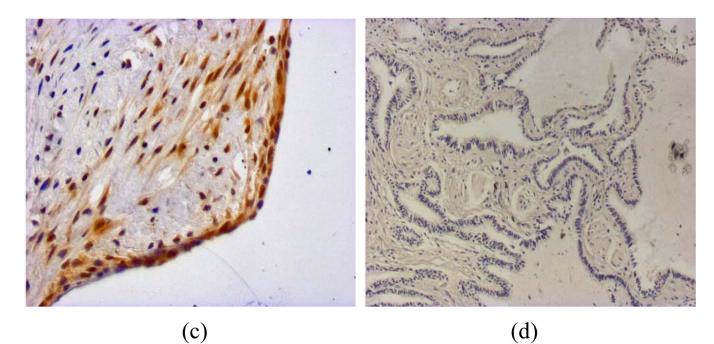


Figure 1.

Biopsies from architecturally normal (a) and IPF (b)–(d) lung were immunohistochemically stained with an antibody to survivin (a)–(c) or with an isotype control antibody (d). (a) Architecturally normal lung tissue from a patient without pulmonary fibrosis shows heterogeneous expression of survivin in alveolar epithelial cells (arrows); (b) An area of alveolar septal thickening in a patient with IPF shows survivin expression in alveolar epithelial cells (arrow) and in cells embedded within the thickened interstitium (arrowhead); (c) A fibroblastic focus from a patient with IPF shows survivin expression in the mesenchymal cells within a fibroblastic focus and in the alveolar epithelial cells overlying the fibroblastic focus; (d) Isotype control staining of IPF lung tissue.

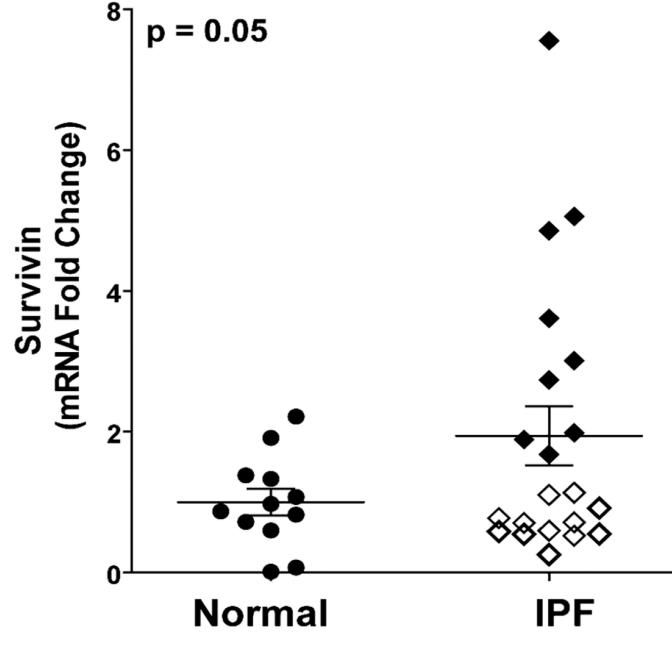


Figure 2.

Quantitative reverse transcription real-time PCR for survivin expression in RNA isolated from explanted lung fibroblasts derived from 12 normal and 21 IPF lungs. Expression levels in the IPF fibroblasts appeared to segregate into a cluster that was indistinguishable from the normal lung fibroblasts (white diamonds) and group that had markedly increased levels of survivin mRNA (black diamonds) p = 0.05.

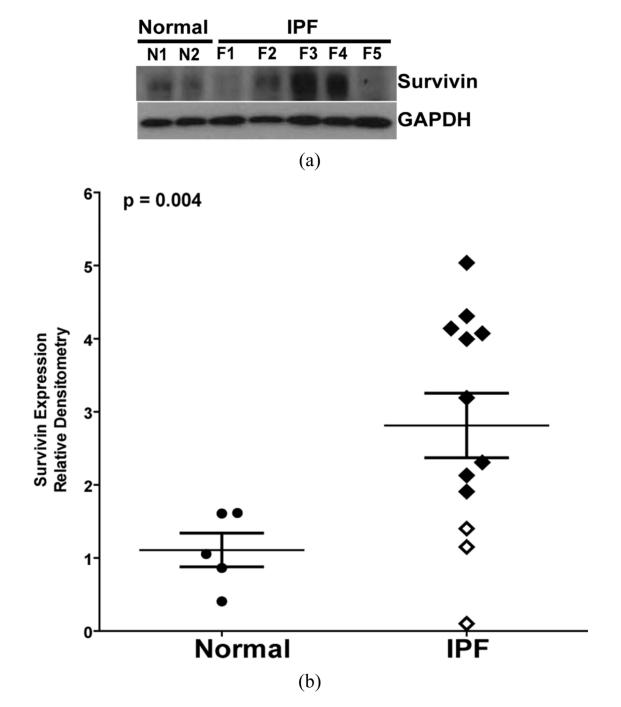


Figure 3.

(a) Cell lysates were obtained from normal and IPF lung fibroblasts and survivin expression assessed by Western blotting. The representative Western blot shows two different normal lung fibroblast cell lines (N) and five different IPF lung fibroblast cell lines (F). The blots were stripped and re-probed for GAPDH; (b) Relative survivin expression in the IPF fibroblasts in comparison to the normal lung fibroblasts was determined by densitometric analysis of indexed to the ratio of survivin to GAPDH in normal lung fibroblasts. Survivin expression in the IPF fibroblasts segregated in to low (white diamonds), and high (black diamonds) levels relative to the normal lung fibroblasts. p = 0.004.

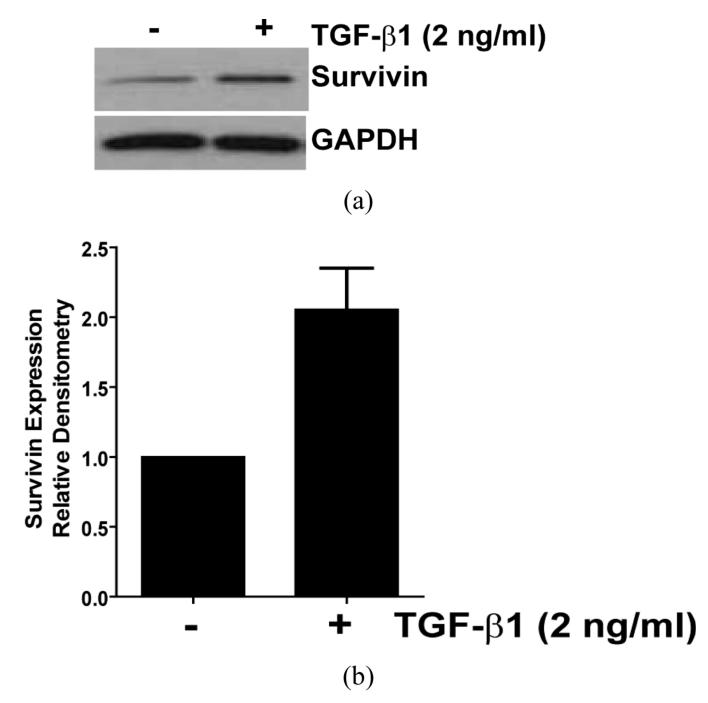


Figure 4.

Normal adult lung fibroblasts (CCL-210) were cultured to 60% confluence in DMEM supplemented with 5% fetal bovine serum and growth arrested in serum-free DMEM for 24 hours prior to treatment with TGF- β 1 (2 ng/ml) for 6 hours. (a) Survivin expression was assessed by Western blotting. The blot was stripped and re-probed for GAPDH, and is representative of three independent experiments; (b) Densitometric analysis of three independent replicates of this experiment, indexed to the ratio of survivin to GAPDH in untreated CCL-210 cells.

Sisson et al.

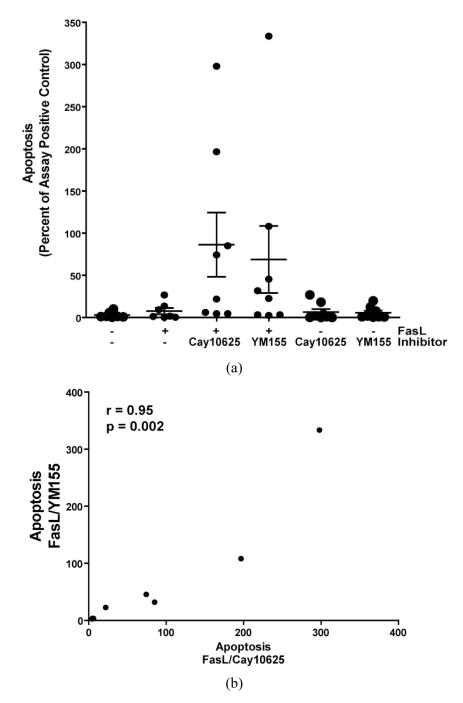


Figure 5.

IPF fibroblasts from 8 different patients were cultured to 60% confluence in DMEM supplemented with 10% fetal bovine serum and growth-arrested for 24 hours in serum-free DMEM prior to treatment for 16 hours with/without the Fas-activating antibody CH11 (FasL; 250 ng/ml) in the presence/absence of the survivin inhibitors CAY10625 (5 μ M) or YM155 (10.0 μ M), or with the survivin inhibitors alone. (a) Apoptosis was assessed using ELISA-based detection of histone associated DNA fragments. To allow relative comparisons of apoptosis across experiments, the data for each cell line is expressed as the percentage of the assay positive control that was included on each ELISA plate. The scatter plot shows the distribution of responses in individual cell lines along with the mean \pm

standard error for each treatment group. p = 0.038 overall (ANOVA). p = 0.027 for FasL compared to FasL/CAY10625, and p = 0.046 for FasL compared to FasL/YM155; (b) The correlation of apoptosis between IPF fibroblasts treated with FasL/CAY10625 and FasL/YM155. Each point represents a different IPF cell line. The Spearman correlation (r) = 0.95.