

A Selective Transforming Growth Factor- β Ligand Trap Attenuates Pulmonary Hypertension

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Abstract

Rationale: Transforming growth factor- β (TGF- β) ligands signal via type I and type II serine-threonine kinase receptors to regulate broad transcriptional programs. Excessive TGF- β -mediated signaling is implicated in the pathogenesis of pulmonary arterial hypertension, based in part on the ability of broad inhibition of activin-like kinase (ALK) receptors 4/5/7 recognizing TGF- β , activin, growth and differentiation factor, and nodal ligands to attenuate experimental pulmonary hypertension (PH). These broad inhibition strategies do not delineate the specific contribution of TGF- β versus a multitude of other ligands, and their translation is limited by cardiovascular and systemic toxicity.

Objectives: We tested the impact of a soluble TGF- β type II receptor extracellular domain expressed as an immunoglobulin-Fc fusion protein (TGFBR2-Fc), serving as a selective TGF- β 1/3 ligand trap, in several experimental PH models.

Methods: Signaling studies used cultured human pulmonary artery smooth muscle cells. PH was studied in monocrotaline-treated Sprague-Dawley rats, SU5416/hypoxia-treated Sprague-Dawley rats, and SU5416/hypoxia-treated C57BL/6 mice. PH, cardiac function, vascular remodeling, and valve structure were assessed by

ultrasound, invasive hemodynamic measurements, and histomorphometry.

Measurements and Main Results: TGFBR2-Fc is an inhibitor of TGF- β 1 and TGF- β 3, but not TGF- β 2, signaling. *In vivo* treatment with TGFBR2-Fc attenuated Smad2 phosphorylation, normalized expression of plasminogen activator inhibitor-1, and mitigated PH and pulmonary vascular remodeling in monocrotaline-treated rats, SU5416/hypoxia-treated rats, and SU5416/hypoxia-treated mice. Administration of TGFBR2-Fc to monocrotaline-treated or SU5416/hypoxia-treated rats with established PH improved right ventricular systolic pressures, right ventricular function, and survival. No cardiac structural or valvular abnormalities were observed after treatment with TGFBR2-Fc.

Conclusions: Our findings are consistent with a pathogenetic role of TGF- β 1/3, demonstrating the efficacy and tolerability of selective TGF- β ligand blockade for improving hemodynamics, remodeling, and survival in multiple experimental PH models.

Keywords: transforming growth factor- β ; pulmonary artery; vascular smooth muscle cells; vascular remodeling; pulmonary hypertension

Pulmonary arterial hypertension (PAH) is a highly morbid condition characterized by elevated pulmonary vascular resistance and arterial pressures, driven by a progressive

pulmonary vasculopathy that leads to right ventricular hypertrophy (RVH), and ultimately, RV failure and death (1, 2). In PAH, arteriolar remodeling is characterized

by medial hypertrophy, neointimal and obstructive lesions consisting of proliferating myofibroblast and endothelial lineages, and multichanneled plexiform

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At a Glance Commentary

Scientific Knowledge on the

Subject: Human and experimental pulmonary hypertension are characterized by an excess of transforming growth factor- β (TGF- β)-mediated signaling, and in particular, an imbalance between TGF- β and bone morphogenetic protein signaling. Previous work suggests that broad inhibition of TGF- β family signaling may be beneficial in pulmonary hypertension, but it is unclear if this strategy is generalizable to different etiologies of disease or if this strategy will be tolerable. We tested whether or not more selective inhibition of TGF- β signaling might be tolerable and effective in several distinct animal models of pulmonary hypertension.

What This Study Adds to the

Field: We found that selective blockade of TGF- β 1 and - β 3 was helpful in reducing pulmonary hypertension and pulmonary vascular remodeling, and in improving survival, using three mechanistically distinct animal models, all of which exhibited signatures of excessive TGF- β signaling. These findings implicate TGF- β 1/3 ligands as drivers of disease, and further investigation of this strategy is warranted.

lesions that are pathognomonic of this disease. Idiopathic and hereditary forms of PAH are associated with heterozygous mutations in *BMPR2* encoding the bone morphogenetic protein (BMP) type II receptor (BMPRII) (3, 4), a member of the transforming growth factor- β (TGF- β) signaling family that has essential functions in the physiologic homeostasis of the vascular endothelium and smooth muscle, as well as other tissues (5–8). The TGF- β family includes a structurally diverse set of more than 33 cytokines that regulate the differentiation, proliferation, migration, and survival of diverse cell types, and include among their members BMPs, activins, inhibins, growth and differentiation factors (GDFs), lefty, nodal, and anti-Mullerian hormone, as well as TGF- β isoforms 1, 2, and 3 (9–11).

Although PAH-associated mutations of *BMPR2* result in loss of BMP signaling function, it has long been observed that lung tissues from human idiopathic PAH are marked by enhanced activity of the TGF- β pathway (12). Multiple animal models of pulmonary hypertension (PH), including those induced by hypoxia, monocrotaline (MCT)-induced injury, or infection with schistosomiasis, have shown similar evidence of elevated TGF- β ligand expression and downstream transcriptional activity (13–16). Enhanced TGF- β signaling in these models has been associated with PH accompanied by smooth muscle hypertrophy, perivascular fibrosis, and extracellular matrix remodeling, all of which can be ameliorated with pharmacologic inhibitors of the TGF- β type I receptor kinases ALK5 (13, 17); these are inhibitors that also inhibit the highly homologous receptors ALK4 and ALK7 and their cognate ligands. However, the therapeutic potential of this strategy has been limited by the observation that ALK4/5/7 inhibitors have caused cardiovascular toxicity in the form of hemorrhagic valve necrosis, as well as physal hypertrophy and dysplasia in the femoral tibial joints of adolescent animals (18). It is unclear which of the individual ligands of the TGF- β family may be primarily responsible for these therapeutic or toxic effects, because ALK4, ALK5, and ALK7 collectively transduce the signals of nearly 20 ligands, including activins, GDFs, and nodal, as well as TGF- β 1, 2, and 3. In the present study, we sought to determine the effects of selective TGF- β ligand blockade in pulmonary vascular remodeling using a recombinant TGFBR2-Fc extracellular domain fusion protein. This ligand trap binds TGF- β 1 and β 3, but does not bind TGF- β 2, whose signaling requires coreceptor TGFBR3, also known as betaglycan (19). We propose that this ligand trap strategy might ameliorate several aspects of pulmonary vascular remodeling and PH that are specifically caused by the activities of TGF- β 1 and TGF- β 3, without incurring the toxic liabilities due to the inhibition of TGF- β 2. TGF- β 2 is known to be unique among TGF- β ligands for its essential roles in cardiac valve morphogenesis, and regulating endothelial and epithelial-to-mesenchymal transition in a variety of tissues (20–28). Part of the present study has been previously reported in abstract form (29).

Methods

Reagents

Recombinant TGFBR2 was expressed as a fusion protein with an IgG Fc domain (TGFBR2-Fc) in CHO cells and purified with two rounds of affinity column chromatography, similar to that previously described in another study (30). Additional information on reagents is provided in the online supplement.

Experimental PH Models

Adult male Sprague-Dawley rats (150–170 g) and male C57BL/6J mice (20–25 g) were purchased from Charles River Laboratory (Wilmington, MA). Experimental protocols were approved by Harvard Institutional Animal Care and Use Committee. Animals were housed at 24°C during a 12-hour light–dark cycle in which food and water were accessible *ad libitum*. In rats, PH was induced either by a single subcutaneous injection of MCT (40 mg/kg) followed by 3 to 5 weeks of normoxia, or a single subcutaneous injection of vascular endothelial growth factor receptor antagonist (SU5416, also known as SUGEN; 20 mg/kg) followed by 3 weeks of normobaric hypoxia (FiO₂ = 10%), then followed by 3 weeks of normoxia. Mice received weekly subcutaneous injections of SU5416 (20 mg/kg) and were exposed to normobaric hypoxia (FiO₂ = 10%) for 3 weeks. Mortality and the total number of animals studied are listed in Table E1 in the online supplement.

Prophylaxis in MCT-treated Rats

Twenty-four hours after administration of MCT, rats were randomized to receive TGFBR2-Fc or vehicle for 21 days. At Day 14, ventricular function and RV hypertrophy (RVH) were examined by echocardiography. At Day 21, rats were subjected to invasive hemodynamic measurements, and tissues were harvested.

Rescue in MCT-treated Rats

To test the ability of TGFBR2-Fc to reverse established PH, starting 18 days after MCT exposure, rats were randomized to receive TGFBR2-Fc or vehicle. Invasive hemodynamic measurements and tissue harvest were performed on Day 35.

Prophylaxis in SU5416/Hypoxia-treated Mice

Twenty-four hours after treatment with SU5416/hypoxia, mice were randomized to receive TGFBR2-Fc or vehicle. Invasive hemodynamic measurements and tissue harvest were performed on Day 21.

Rescue in SU5416/Hypoxia-treated Rats

Rats were treated with SU5416/hypoxia for 3 weeks. Rats were returned to normoxia and randomized to receive TGFBR2-Fc or vehicle for 3 weeks. Echocardiographic, invasive hemodynamic measurements, and tissue harvest were performed on Day 42. Additional details are provided in the online supplement.

Invasive Hemodynamic Measurements

Rats were anesthetized with pentobarbital (50 $\mu\text{g}/\text{kg}$ i.p.) and intubated intratracheally for mechanical ventilation (tidal volume = 8 ml/kg, frequency = 80/min). Invasive hemodynamic measurements of RV pressures were obtained by cannulation via the RV apex for MCT experiments as described previously (31), or alternatively for SU5416/hypoxia experiments, by a minimally invasive closed chest approach using a curved tip 2F pressure transducer catheter (Millar, Houston, TX; SPR-513) inserted into the RV through the right internal jugular vein. Invasive hemodynamic measurements of RV pressure in anesthetized mice were performed using a 1.2-F pressure catheter (Transonic Scisense Inc., Ithaca, NY) via the internal jugular vein.

Statistical Analysis

Measurements and analysis of physiological parameters and vascular remodeling were performed in blinded fashion. Data are presented as mean \pm SEM and compared between groups using the Student's *t* test, with the Bonferroni correction for multiple tests, or analysis of variance as appropriate; $P < 0.05$ was considered statistically significant.

Results

Recombinant TGFBR2-Fc selectively inhibited TGF- β 1- and TGF- β 3-mediated transcriptional activity, measured in HEK293 cells stably transfected with the

CAGA-luciferase reporter (Figure 1A). TGFBR2-Fc failed to suppress, and in fact modestly enhanced, the activity of TGF- β 2. TGFBR2-Fc did not inhibit BMP-mediated transcriptional activity, measured in C2C12 cells stably transfected with the BMP response element (BRE)-luciferase reporter, but enhanced the activity of several ligands in this assay (Figure 1B). These results were consistent with selective activity of this ligand trap against a subset of TGF- β ligands, and suggested a feedback mechanism in which suppression of TGF- β signaling resulted in enhanced BMP and TGF- β 2 signaling. TGFBR2-Fc inhibited TGF- β 1- and TGF- β 3-induced phosphorylation of Smads 2/3 (Figure 1C and Figure E1 in the online supplement) and inhibited the increase in *Pai-1* expression induced by TGF- β 1 or TGF- β 3 (Figure 1D) in human pulmonary artery smooth muscle cells, without affecting the activity of TGF- β 2. TGF- β 1 potently elicited the expression of smooth muscle contractility genes *Caldesmon*, *Smoothelin*, and *Calponin* in human pulmonary artery smooth muscle cells, all of which were inhibited by TGFBR2-Fc (Figures 1E and 1F). Taken together, TGFBR2-Fc acts as a potent antagonist of TGF- β 1 and TGF- β 3 signaling and abrogates the impact of these ligands on vascular smooth muscle cell plasticity *in vitro*.

Rats treated with MCT developed progressive elevation of RV systolic pressures (RVSPs) (Figure E2a) and Fulton's ratio, which was consistent with RVH (Figure E2b) and evolved in a time-dependent fashion over 3 weeks after MCT treatment. In the lung tissues of MCT rats, we observed that the development of PH over 3 weeks was associated with progressively diminished expression of *Bmpr2*, progressively diminished BMP-mediated signaling, and gene transcription based on the expression of *Id1*, accompanied with a marked increase in TGF- β -mediated signaling and transcription based on the expression of *Pai-1* (Figures E2c–E2f) (32). Disease severity, measured by RVSP and Fulton's ratio, were positively correlated with TGF- β signaling activity, based on *Tgfb1* and *Pai-1* expression, and negatively correlated with *Bmpr2* expression and downstream transcriptional activity based on *Id1* (Figure E3).

We tested the hypothesis that TGFBR2-Fc could inhibit TGF- β 1/3 signaling activity *in vivo* to modulate

disease progression in experimental PH. Treatment with a low dose of TGFBR2-Fc (5 mg/kg i.p., twice weekly, starting 1 day after MCT injection) resulted in a trend toward decreased RVSP, RVH, and muscularization of small pulmonary vessels, and significantly decreased medial wall thickness index after MCT treatment (Figure 2). Testing whether or not there might be a dose-dependent effect, treatment with higher doses of TGFBR2-Fc (15 mg/kg i.p., twice weekly, starting 1 day after MCT injection) was found to significantly alter RVSP (33.0 ± 2.1 vs. 46.7 ± 4.7 mm Hg in vehicle-treated control animals) (Figure 3A) and RVH (0.41 ± 0.02 vs. 0.52 ± 0.04 in vehicle-treated control animals) (Figure 3B). TGFBR2-Fc treatment under this regimen effectively prevented vascular remodeling in MCT rats (Figures 3C–3E). Whole-lung tissues of MCT rats that received TGFBR2-Fc exhibited decreased mRNA levels of *Pai-1* and unexpectedly exhibited reduced levels of *Tgfb1* (Figures 3F and 3G). The previously observed reduction in *Bmpr2* and *Id1* expression observed in MCT rats was unaffected by TGFBR2-Fc treatment (Figure E4). TGFBR2-Fc treatment also reduced the expression of *Il6* (Figure 3H), *Il1b* (Figure 3I), and intercellular adhesion molecule (*Icam*) (Figure 3J) mRNA levels in the lungs of MCT-treated rats. Consistent with enhanced TGF- β signaling activity, MCT-induced PH was associated with increased expression of phosphorylated Smad2 as detected by immunohistochemistry, primarily in the nuclei of vascular and perivascular cells, whereas treatment with TGFBR2-Fc attenuated this increase in phosphorylated Smad2 (Figure 3K). These data were confirmed by Western blot (Figure 3L and Figure E4e). MCT treatment robustly decreased the expression of phosphorylated Smad1/5/8 in lung extracts. However, in contrast to the reflexive increases in BMP signaling observed *in vitro*, there were no gains in the phosphorylation of Smad1/5/8 in the lung tissues of MCT rats that received TGFBR2-Fc.

Using echocardiography to provide a complementary measurement of RVH, treatment with TGFBR2-Fc was found to attenuate the increase in RV free-wall thickness observed in MCT rats at Day 14 (0.77 ± 0.15 vs. 0.59 ± 0.11 mm; $P < 0.05$) (Table E2). However, the decreased pulmonary acceleration time (PAT)

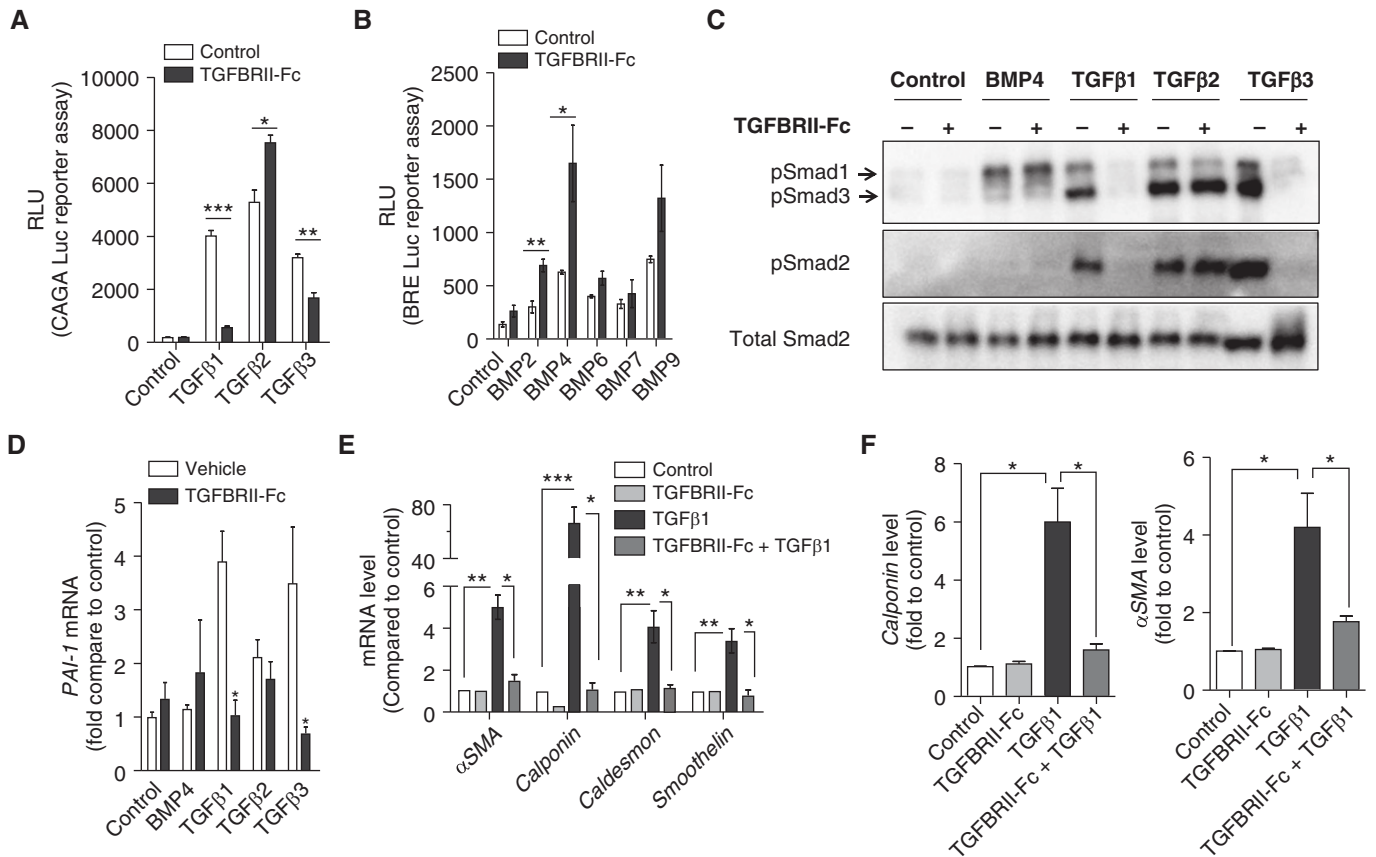


Figure 1. TGFBR2-Fc selectively inhibits transforming growth factor-β1 (TGF-β1)- and TGF-β3-induced signaling and blocks TGF-β-induced phenotypic modulation of smooth muscle cells. (A) HEK293 cells stably expressing a TGF-β-responsive CAGA-luciferase (Luc) reporter transgene and (B) C2C12 cells stably expressing a bone morphogenetic protein (BMP) response element reporter (BRE)-Luc reporter transgene were incubated in the presence or absence of TGFBR2-Fc (2 μg/ml) for 30 minutes before stimulation with various BMP ligands (10 ng/ml) or TGF-β1, -2, or -3 (1 ng/ml) overnight, revealing selective inhibition by TGFBR2-Fc of TGF-β1- and TGF-β3- but not TGF-β2-induced activation of CAGA-luciferase activity, and slightly increased BRE-luciferase activity in response to BMP2 and BMP4 in the presence of TGFBR2-Fc. Human pulmonary artery smooth muscle cells were deprived of serum overnight, pretreated with TGFBR2-Fc (2 μg/ml), followed by incubation with BMP4 (10 ng/ml), TGF-β1, -2, or -3 (1 ng/ml of each for 30 min), and analyzed by immunoblot for (C) phosphorylated Smads (pSmad) 1, 2, and 3 and (D) for mRNA expression of the TGF-β transcriptional target *PAI-1* by quantitative reverse transcriptase polymerase chain reaction. TGFBR2-Fc inhibited SMAD2 and SMAD3 activation, CAGA-Luc reporter activity, and *Pai-1* mRNA expression via TGF-β1 and TGF-β3, but not TGF-β2. After stimulation with TGF-β1, the relative levels of (E) mRNA and (F) protein expression of smooth muscle cell contractile markers were examined at 24 and 72 hours, respectively. Data are expressed as mean ± SEM. **P* < 0.05; ***P* < 0.01; ****P* < 0.001 as indicated. αSMA = α-smooth muscle actin; PAI-1 = plasminogen activator inhibitor 1; RLU = relative light units.

observed in MCT rats (19.8 ± 2.6 vs. 32.6 ± 2.2 ms; *P* < 0.05) (Table E2) was not significantly affected by TGFBR2-Fc treatment. Importantly, echocardiographic M-mode, pulsed-wave Doppler, and two-dimensional color Doppler assessment did not reveal evidence of valvular dysfunction or degeneration due to TGFBR2-Fc administration in normal or MCT rats (Table E2). Similarly, TGFBR2-Fc treatment was not associated with morphological changes in mitral valve structure, with no evidence of sclerotic or degenerative remodeling in sectioned valve tissues in rats (Figure E5). TGFBR2-Fc treatment did not significantly affect body weight in rats (Figure E6).

To assess whether or not modulation of TGF-β signaling might affect the progression of established PH, we tested the impact of TGFBR2-Fc (15 mg/kg i.p., three times per week) starting at 18 days after MCT exposure by assessing hemodynamics and pulmonary arteriolar remodeling at 35 days. When administered in this delayed fashion, TGFBR2-Fc significantly reduced RVSP (33.3 ± 0.8 vs. 43.2 ± 3.5 mm Hg in vehicle-treated control animals; *P* < 0.01) and attenuated arteriolar muscularization (*P* < 0.05) in MCT rats (Figure 4). Consistent with its impact on hemodynamics and remodeling, delayed treatment with TGFBR2-Fc improved survival compared with the vehicle

(Figure 4C). In addition to reducing the mRNA expression of *Tgfb1* and *Pai-1* (Figures 4F–4J), delayed administration of TGFBR2-Fc significantly decreased the expression of *Il6*, *Il1b*, and *Icam* in the lungs of MCT rats, again suggesting an antiinflammatory effect of this treatment. As in previous experiments, delayed treatment with TGFBR2-Fc in MCT rats decreased levels of phosphorylated Smad2 (Figure 4K).

To explore the role of TGF-β-mediated signaling in a mechanistically distinct model of PH, we tested the effects of TGFBR2-Fc in the SU5416/hypoxia mouse model, in which mice were treated with vascular endothelial growth factor receptor

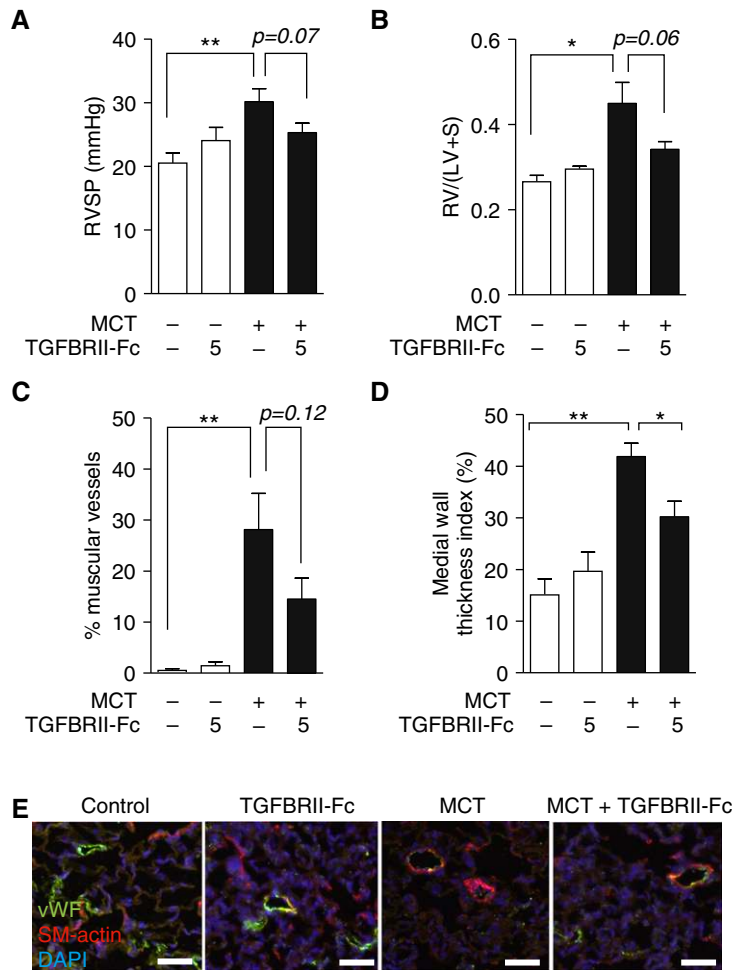


Figure 2. Lower-dose treatment with TGFBR2-Fc elicits trends toward improved pulmonary hypertension, right ventricular hypertension (RVH), and pulmonary vascular remodeling. (A) Three weeks after treatment with monocrotaline (MCT) with or without TGFBR2-Fc (5 mg/kg, twice weekly i.p.), rats were analyzed in a blinded fashion to determine right ventricular systolic pressure (RVSP). (B) The degree of RVH was assessed based on measurement of Fulton's ratio [RV/(LV + S)]. (C) Muscularization of distal intraacinar vessels (10–50 μ m diameter) was quantified, and the percentages of fully (circumferentially) muscularized vessels were calculated. (D) Medial wall thickness was calculated for all fully muscularized intraacinar vessels (10–50 μ m diameter). Wall thickness index was calculated as index = (external diameter – internal diameter)/external diameter \times 100. TGFBR2-Fc treatment under this dosing regimen trended toward decreased frequency of muscularized vessels, and significantly reduced medial wall thickness, based on 100 to 150 vessels per treatment group from 6 to 8 rats each. * $P < 0.05$ or ** $P < 0.01$ versus control animals or otherwise as shown. (E) Immunofluorescence of lung sections for von Willebrand's factor (vWF) to mark vessels and smooth muscle actin (SM-actin) revealed qualitatively increased muscularization of small (<50 μ m) arterioles following MCT treatment, which was reduced by the administration of TGFBR2-Fc (scale bars = 50 μ m). Nuclei were counterstained with 4',6-diamidino-2-phenylindole (DAPI). Data are shown as mean \pm SEM ($n = 6$ –8). * $P < 0.05$ and ** $P < 0.01$ as indicated. LV = left ventricle; RV = right ventricle; S = septum.

inhibitor SU5416 and exposed to chronic hypoxia for 3 weeks. Combined treatment of SU5416 and hypoxia triggered more severe and consistent PH and RVH than that seen in mice exposed to hypoxia alone (see Figures E8a and E8b in the online supplement). Prophylactic treatment with TGFBR2-Fc significantly reduced RVSP

(35 \pm 6 mm Hg vs. 43 \pm 6 mm Hg in vehicle-treated control animals; $P < 0.05$) (Figure 5A) and RVH (0.24 \pm 0.02 mm vs. 0.29 \pm 0.01 mm in vehicle-treated control animals; $P < 0.05$) (Figure 5B) in SU5416/hypoxia mice. TGFBR2-Fc treatment decreased pulmonary arteriolar muscularization in the lungs of SU5416/

hypoxia mice (Figure 5E). Administration of TGFBR2-Fc elicited a trend toward decreased expression of *Tgfb1* (Figure 5C) and significantly reduced *Pai-1* mRNA expression in the lungs of SU5416/hypoxia mice (Figure 5D). TGFBR2-Fc treatment qualitatively reduced the frequency of Ki67-positive cells in the lungs of SU5416/hypoxia mice, which was seen primarily in the adventitial, airway epithelial, and alveolar cells in vehicle-treated mice (see Figure E8c in the online supplement).

Finally, we tested the efficacy of TGFBR2-Fc in ameliorating PH in an SU5416/hypoxia-induced rat model of PH, which was shown previously to closely recapitulate vascular remodeling features observed in patients with severe PAH (33). Rats were treated with SU5416 during 3 weeks of chronic normobaric hypoxia ($F_{I_{O_2}} = 10\%$), after which they were returned to normoxia for 3 weeks, and randomized to receive either TGFBR2-Fc or vehicle. At the end of the 6-week period, SU5416/hypoxia-treated rats that received vehicle had significantly higher RVSP (38.5 \pm 4.4 mm Hg vs. 20.8 \pm 1.4 mm Hg; $P < 0.01$) (Figure 6) and a significantly shorter PAT (13.4 \pm 0.5 ms vs. 21.6 \pm 1.3 ms in vehicle-treated control animals; $P = 0.001$; Figure 6 and Figure E9 in the online supplement). TGFBR2-Fc-treated animals exhibited significant improvement in PH demonstrated by a normalization of RVSP (17.2 \pm 1.9 mm Hg; $P < 0.01$ vs. SU5416/hypoxia-treated animals) (Figure 6) and PAT (20.8 \pm 1.4 ms; $P < 0.01$ vs. SU5416/hypoxia-treated animals) (Figure 6 and Figure E9 in the online supplement). SU5416/hypoxia-treated rats exhibited severe RVH based on an increased ratio of the weight of the right ventricle to that of the left ventricle and septum [RV/(LV + S)] (0.46 \pm 0.1 vs. 0.21 \pm 0.1; $P < 0.01$) and increased RV end-diastolic wall thickness by echocardiography (1.3 \pm 0.1 vs. 0.5 \pm 0.02 mm in vehicle-treated control animals; $P < 0.001$; Figure 6 and Figure E10). These rats developed RV dysfunction as demonstrated by decreased tricuspid annular plane systolic excursion, a noninvasive echocardiographic surrogate metric of RV function (0.9 \pm 0.1 mm vs. 1.8 \pm 0.2 mm in vehicle-treated control animals; $P = 0.001$; Figure 6D). TGFBR2-Fc-treated rats

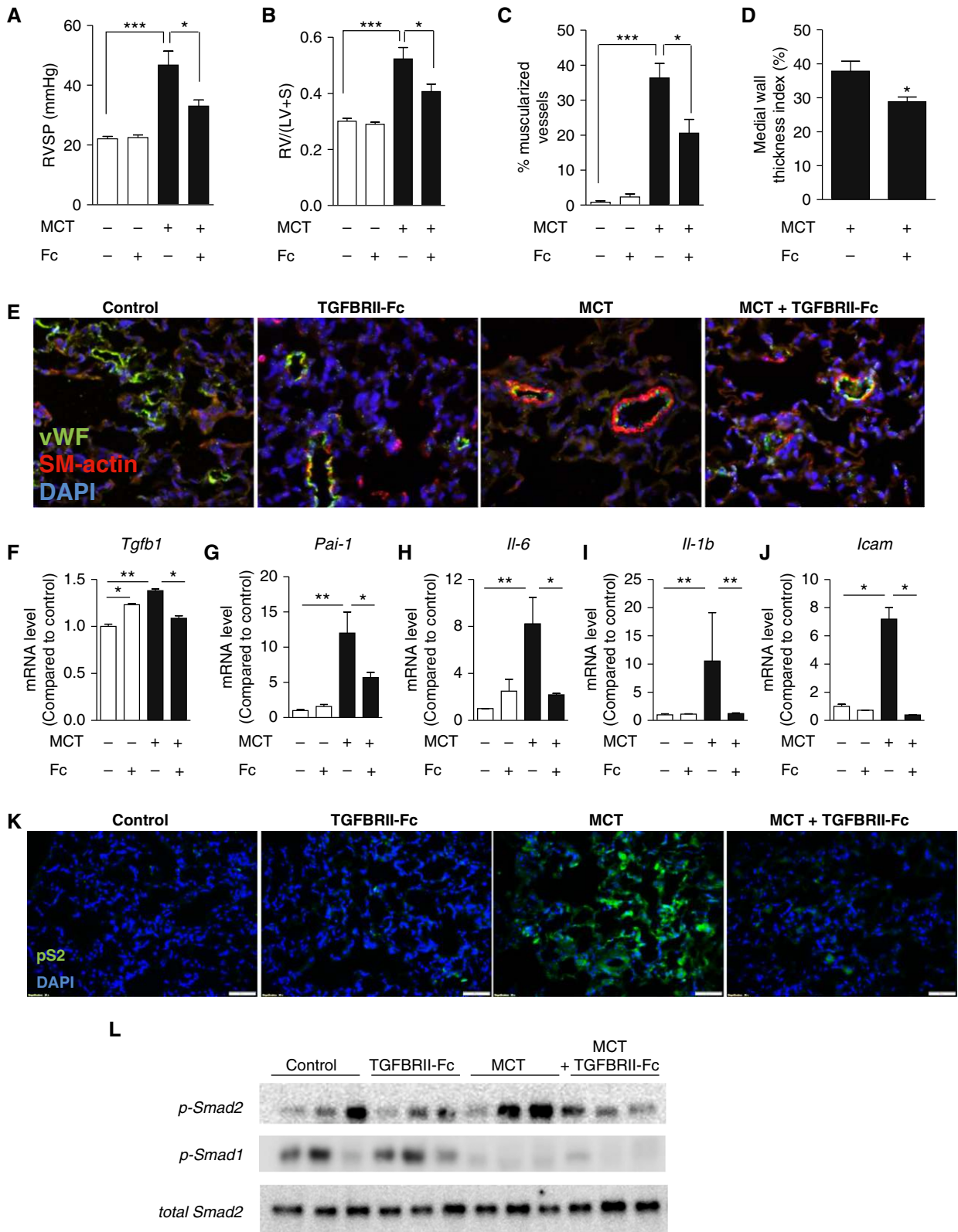


Figure 3. Higher-dose treatment with TGFBR11-Fc prevented pulmonary hypertension and vascular remodeling. Adult rats treated with monocrotaline (MCT) (40 mg/kg s.c. \times 1) were administered TGFBR11-Fc (15 mg/kg i.p. twice weekly, starting 1 day after MCT injection) or vehicle for 3 weeks. Treatment with TGFBR11-Fc significantly attenuated (A) right ventricular systolic pressure (RVSP) and (B) right ventricular hypertrophy in comparison to vehicle ($n = 6-8$). (C and D) TGFBR11-Fc decreased the percentage of fully muscularized vessels (10–50 μ m diameter) (C) and medial wall thickness, calculated as

exhibited improved RVH on the basis of decreased RV/(LV + S) (0.30 ± 0.03 ; $P < 0.05$ vs. SU5416/hypoxia-treated rats; Figure 6) and decreased RV end-diastolic wall thickness by echocardiography (0.7 ± 0.1 mm; $P < 0.001$ vs. SU5416/hypoxia-treated rats; Figure 6 and Figure E10 in the online supplement), as well as improved RV function demonstrated by increased tricuspid annular plane systolic excursion (1.2 ± 0.1 mm; $P = 0.01$ vs. SU5416/hypoxia-treated rats; Figure 6D). In addition, TGFBR11-Fc treatment markedly attenuated vascular remodeling based on muscularization and medial wall thickness (Figures 6F, 6G, and 6I), reduced vascular cell proliferation based on the frequency of Ki67-positive cells in the adventitia and intima (Figures 6H and 6J), and attenuated RV fibrosis in SU5416/hypoxia-treated rats (Figure S11). Consistent with the observed improvements in PH, RVH, and PAT, there was an observed trend toward improved survival among TGFBR11-Fc-treated rats (Figure E12).

Discussion

Based on the known importance of TGF- β signaling in pulmonary vascular disease, and the ability of recombinant ligand traps to modulate the activity of selected ligands *in vivo*, we tested whether or not a selective TGF- β ligand trap could affect experimental PH and pulmonary vascular remodeling. We characterized recombinant TGFBR11-Fc fusion protein as a potent inhibitor of TGF- β 1 and TGF- β 3 signaling that did not affect the signaling of TGF- β 2, which is an important regulator of endothelial-to-mesenchymal transition, extracellular matrix remodeling, and cardiac valve formation (24, 26). Consistent with this activity, TGFBR11-Fc blocked canonical Smad 2/3 signaling via TGF- β 1 and TGF- β 3, but did not block TGF- β 2, and prevented TGF- β 1-mediated switching of

pulmonary artery smooth muscle cells from a synthetic to a contractile phenotype *in vitro* (34, 35). Hypothesizing that TGF- β 1 and/or TGF- β 3 contribute to pulmonary vascular remodeling seen in PH, we found that TGFBR11-Fc treatment corrected excessive TGF- β signaling activity and thereby attenuated pulmonary vascular remodeling, pulmonary hypertension, and RV hypertrophy in three distinct models of experimental PH. TGFBR11-Fc not only prevented the progression of established PH and RVH in rats treated with MCT or SU5416/hypoxia, but also improved their survival. These present data are the first demonstration that selective inhibition of specific TGF- β ligands may attenuate PH, using multiple, mechanistically distinct experimental models of PH, by applying a strategy that appears to be more tolerable than nonselective approaches.

TGF- β family ligands orchestrate numerous processes during embryogenesis and play critical roles in the pathophysiologic remodeling seen in vascular and fibrotic diseases, roles that have generated tremendous interest in their blockade for therapy (36–41). Broad blockade of TGF- β signaling via small molecules IN-1333, SD-208, and SB-525334 was shown to attenuate MCT-induced PH in rats, purportedly by inhibiting TGF- β signaling via the activity of its type I receptor ALK5 (13, 17, 32). However, these molecules demonstrated potent activity for closely related type I receptors ALK4 and ALK7 (42–44), and would thus affect the activity of not only TGF- β 1 and TGF- β 3 but nearly 20 other ligands, including activins, GDFs, and nodal, whose functions may or may not be concordant with TGF- β in the vasculature, and many of which may serve important roles in tissue homeostasis. These potent ALK4/5/7 inhibitors appear to have a class-wide effect of causing hemorrhagic valve necrosis in animals (18), possibly due to their lack of selectivity, a property

that has limited their deployment for human therapy. It was recently shown that a neutralizing monoclonal antibody (1D11) that recognizes TGF- β ligands TGF- β 1, - β 2, and - β 3 was also able to attenuate PH in a MCT rats (1, 31). The use of a similar pan-TGF- β neutralizing antibody in humans, fresolimumab, although generally well-tolerated, has been associated with dose-related side effects, including skin rashes and lesions, epistaxis, gingival bleeding, and fatigue (36). Moreover, TGF- β 2 is essential in cardiac valve morphogenesis, which is consistent with its central role in regulating endothelial- and epithelial-to-mesenchymal transition in a variety of tissues (20–28), and raising the possibility that its inhibition might contribute to the valvular toxicity seen with small molecule inhibitors of ALK4/5/7. Consistent with the concept that the TGF- β type II receptor does not transduce TGF- β 2 signals in the absence of the type III co-receptor betaglycan (19, 45–47), soluble TGFBR11-Fc did not inhibit TGF- β 2 or cause any observable changes in cardiac valve structure or function. In contrast to the effects of ALK4/5/7 inhibitors seen in toxicology studies (18), TGFBR11-Fc was well tolerated, even when higher doses were used in rescue studies with established PH. Although it cannot be ruled out that toxicity might be observed at higher or longer exposures, the doses used in this study were sufficient to inhibit TGF- β -induced Smad2 activation and *Pai-1* expression in lung tissues.

A limitation of previous studies investigating the role of TGF- β in PH has been the reliance on MCT-induced PH in rats, a preclinical model known to exhibit a molecular signature of exaggerated TGF- β signaling and profibrotic transcriptional activity, suppression of BMPR2 expression, and downstream BMP signaling activity (32), with evidence of diffuse inflammation and lung injury (48), as were observed in the

Figure 3. (Continued). (external diameter – internal diameter)/external diameter $\times 100$ (D), based on 89 to 127 vessels per treatment group from 6 to 8 rats each. (E) TGFBR11-Fc treatment reduced muscularization evident by smooth muscle actin staining of von Willebrand (vWF)⁺ small vessels. (F–J) TGFBR11-Fc treatment normalized the expression of *Tgfb1* (F), *Pai-1* (G), *Il-6* (H), *Il1b* (I), and *Icam* (J) in the lung tissues of MCT-treated rats (n = 3–5). (K) Immunohistochemistry of lung sections and (L) immunoblotting of whole-lung lysates demonstrated qualitatively increased *p-Smad2* (pS2) expression in the lungs of MCT-treated animals, which was normalized by treatment with TGFBR11-Fc. Conversely, treatment with MCT robustly decreased expression of *p-Smad1/5/8*, which was not significantly affected by treatment with TGFBR11-Fc. Nuclei were counterstained with 4',6-diamidino-2-phenylindole (DAPI). Data are expressed as mean \pm SEM. * $P < 0.05$; ** $P < 0.01$; *** $P < 0.001$ as indicated. Scale bars = 50 μ m. *Icam* = intercellular adhesion molecule; i.p. = intraperitoneal; LV = left ventricle; *Pai-1* = plasminogen activator inhibitor 1; RV = right ventricle; S = septum; s.c. = subcutaneous; *Tgfb1* = transforming growth factor- β 1.

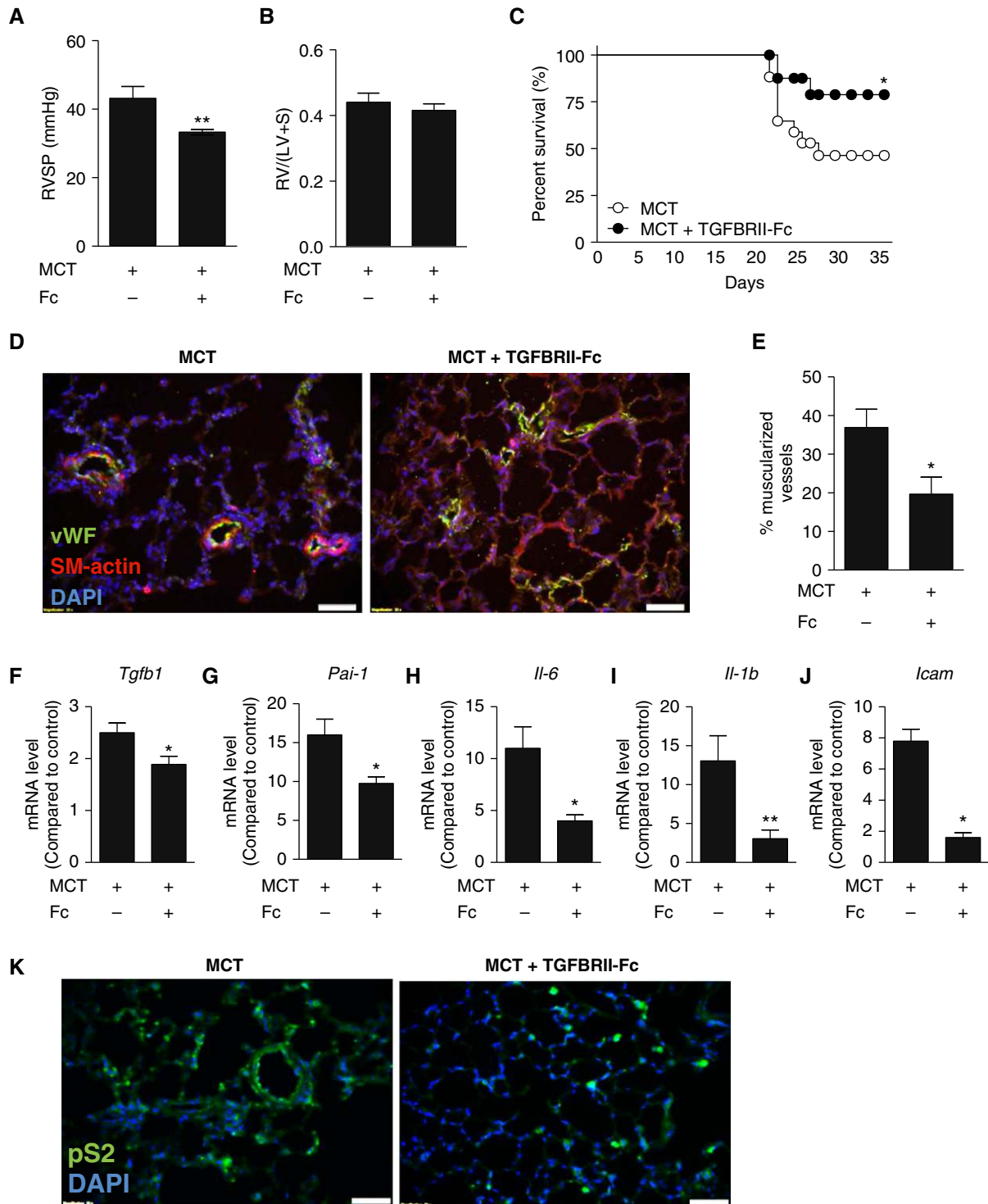


Figure 4. Treatment with TGFBR11-Fc after establishment of pulmonary hypertension (PH) is associated with partial rescue of PH and mortality. After monocrotaline (MCT) treatment, rats were administered TGFBR11-Fc (15 mg/kg three times weekly) in a delayed fashion starting on Day 17, after the establishment of PH. Among surviving animals at 35 days, there was (A) significantly decreased right ventricular systolic pressure (RVSP) with TGFBR11-Fc treatment, but (B) no significant difference in right ventricular hypertension. Values are shown as mean \pm SEM (n = 8–11 per group). (C) Kaplan-Meier analysis revealed improved survival in the TGFBR11-Fc-treated group (n = 18 per group). (D and E) TGFBR11-Fc treatment attenuated pulmonary vascular remodeling in rats with established PH. (F–J) Delayed treatment with TGFBR11-Fc reduced the expression of *Tgfb1* (F), *Pai-1* (G), *Il6* (H), *Il1b* (I), and *Icam* (J) in the lung tissues of MCT-treated rats (n = 3–5). (K) TGFBR11-Fc treatment reduced elevated phospho-Smad2 (pS2) levels in MCT lungs. Nuclei were counterstained with 4',6-diamidino-2-phenylindole (DAPI). Data are shown as mean \pm SEM. * $P < 0.05$; ** $P < 0.01$ as indicated. Scale bars = 50 μ m. *Icam* = intercellular adhesion molecule; LV = left ventricle; *Pai-1* = plasminogen activator inhibitor 1; RV = right ventricle; S = septum; SM-actin = smooth muscle actin; *Tgfb1* = transforming growth factor- β 1; vWF = von Willebrand factor.

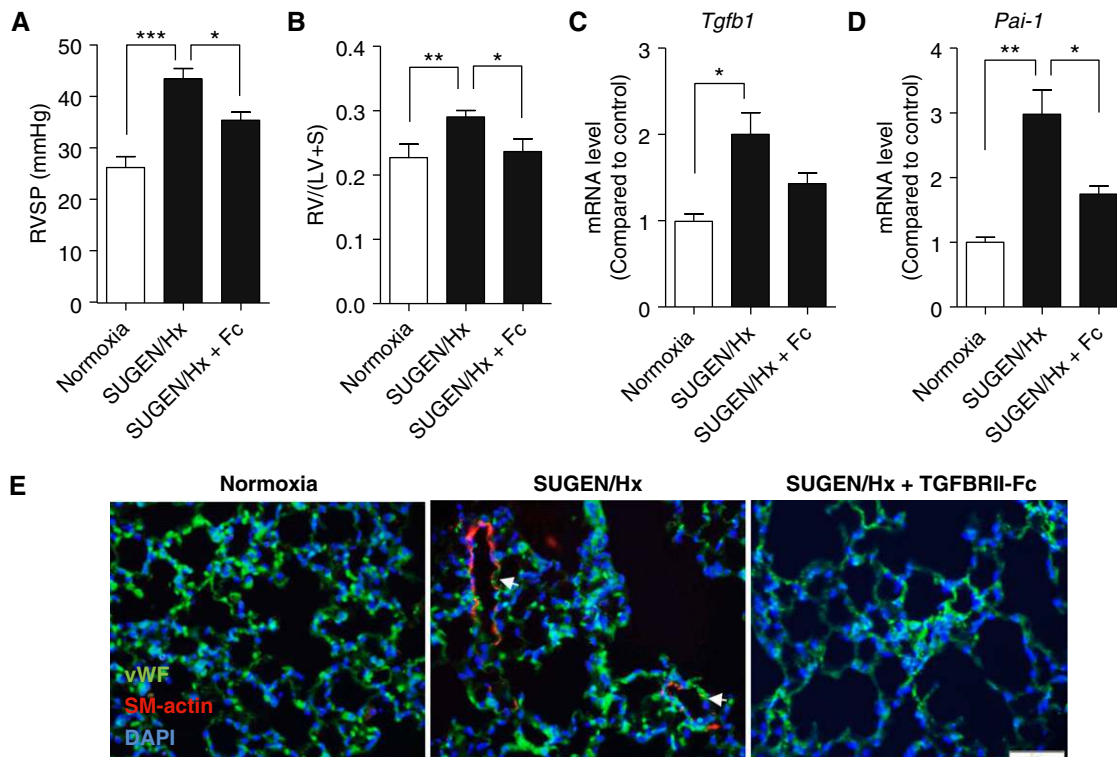


Figure 5. Efficacy of TGFBR11-Fc in a murine model of pulmonary hypertension. Adult male mice were treated with SU5416 (SUGEN) and exposed to hypoxia for 3 weeks. (A and B) TGFBR11-Fc treatment (15 mg/kg, three times per week) reduced right ventricular systolic pressure (RVSP) (A) and prevented right ventricular hypertrophy (B) compared with vehicle-treated mice. (C and D) TGFBR11-Fc treatment trended toward reduced *Tgfb1* (C) and prevented the up-regulation of *Pai-1* (D) mRNA levels in lungs of SU5416/hypoxia-treated mice. (E) TGFBR11-Fc treatment ameliorated pulmonary vascular remodeling. Data are expressed as mean \pm SEM (n = 6–8). * P < 0.05; ** P < 0.01; *** P < 0.001 as indicated. Scale bars = 50 μ m. DAPI = 4',6-diamidino-2-phenylindole; Hx = hypoxia; LV = left ventricle; *Pai-1* = plasminogen activator inhibitor 1; RV = right ventricle; S = septum; SM-actin = smooth muscle actin; *Tgfb1* = transforming growth factor- β 1; vWF = von Willebrand factor.

present study. Although this phenotype might highlight potential effects of TGF- β inhibition, these characteristics may or may not reflect the spectrum of human PAH disease. Other animal models including hypoxia-induced PH in rats (49) and PH associated with the administration of *Schistosoma mansoni* (16) were found to exhibit signatures of enhanced TGF- β expression. Consistent with a role of TGF- β in human PAH, elevated levels of TGF- β 1 have been reported in the transpulmonary flux of individuals with PAH (50), as well as in the circulation and lungs of individuals with idiopathic PAH (51), and increased Smad2 activity was found in the lung tissues of individuals with heritable PAH (52). Similarly, functional polymorphisms regulating the expression of *TGFB1* affect the age at diagnosis and penetrance of PAH associated with *BMPR2* mutations, consistent with the concept that BMP and TGF- β signaling imbalance contributes to disease (52). In the mechanistically

distinct SU5416/hypoxia-induced mouse model (53), we found evidence of increased TGF- β signaling activity that was responsive to TGFBR11-Fc with improved RVSPs and RVH. In SU5416/hypoxia-treated rats, we found that TGFBR11-Fc treatment reversed RVH and fibrosis, improved RV function, and normalized hemodynamic and echocardiographic measures of PH. Taken together, our current and previous data indicate that the pathogenetic role of TGF- β may be broadly generalizable across several experimental PH models and human PAH.

TGFBR11-Fc inhibited TGF- β 1-induced smooth muscle phenotypic modulation *in vitro*, although its administration in animal models of PH blocked TGF- β -mediated transcription and the expression of inflammatory signaling molecules previously implicated in PH (54, 55). The effects of TGF- β signaling in PH have been attributed to recruitment of myogenic and fibrogenic

transcriptional programs in vascular medial hypertrophy and medial extension, but the present findings suggest TGF- β may also potentiate inflammation in affected lungs. Unexpectedly, we observed that administration of TGFBR11-Fc decreased the expression of *Tgfb1* mRNA itself when analyzed in severe stages of disease (Figures 3–5). Dampening of ligand expression by a TGF- β ligand trap suggests TGFBR11-Fc may disrupt positive feedback mechanisms of TGF- β signaling that may be intrinsic to the TGF- β pathway or to profibrotic or proinflammatory cascades triggered by TGF- β 1. Positive feedback or signal amplification is a motif frequently observed in BMP/TGF- β signaling, and the present observations are reminiscent of the recent report that administration of exogenous BMP9 ligand stimulates the BMP pathway while simultaneously up-regulating its cognate receptor BMPRII in animal models of PH (56).

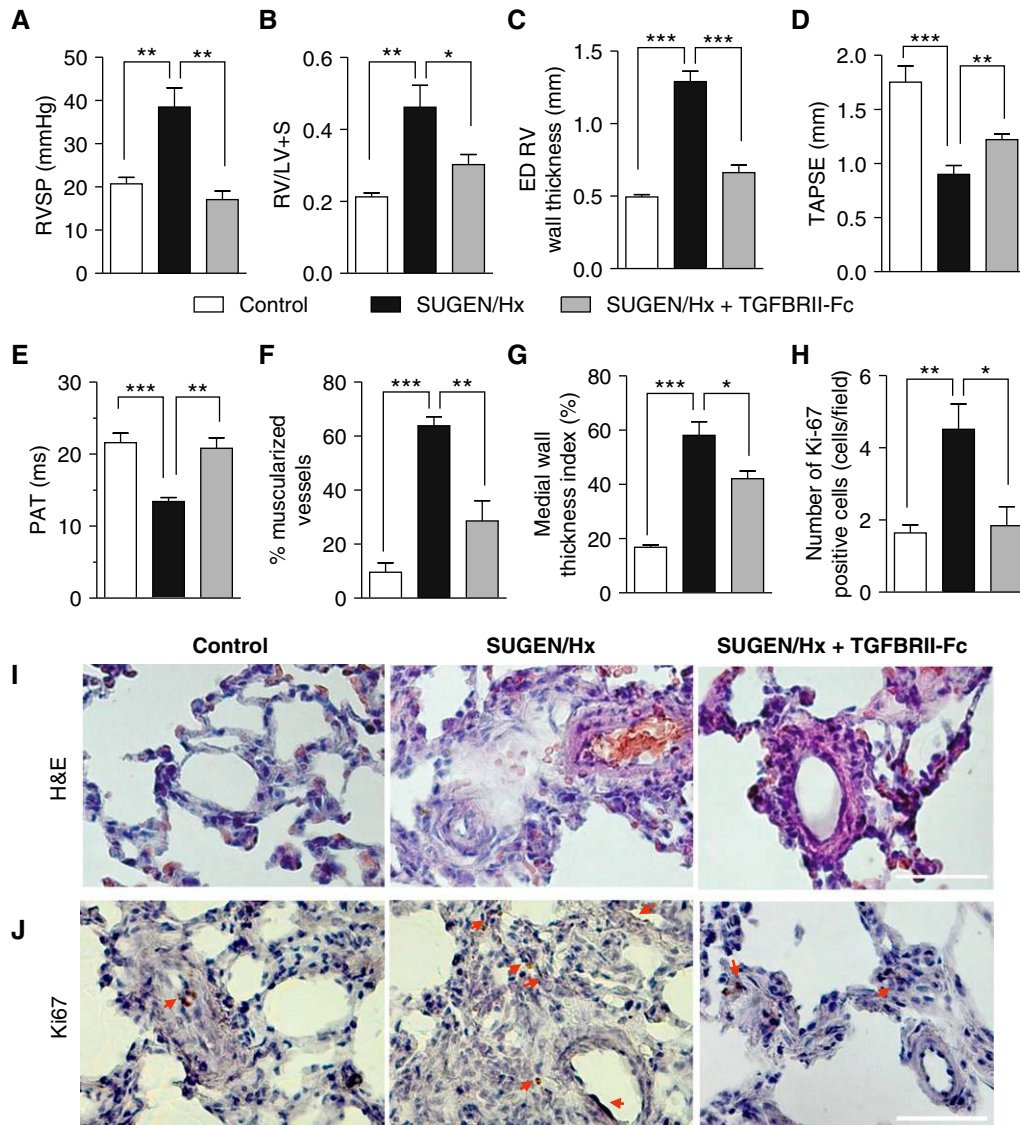


Figure 6. Treatment with TGFBR11-Fc reverses established pulmonary hypertension in SU5416 (SUGEN)/hypoxia-treated rats. Male adult Sprague-Dawley rats received a single subcutaneous injection of SU5416 and were subjected to normobaric hypoxia ($F_{I_{O_2}} = 0.10$) for 3 weeks, followed by maintenance in normoxia for 3 weeks, during which rats were randomized to receive either vehicle or TGFBR11-Fc (15 mg/kg, three times per week). (A–C) TGFBR11-Fc treatment reduced right ventricular systolic pressure (RVSP) (A), attenuated right ventricular hypertension (B), and reduced echocardiographic end-diastolic (ED) right ventricular free wall thickness (C) compared with vehicle-treated rats. (D and E) TGFBR11-Fc treatment improved echocardiographic measures of right ventricular function via tricuspid annular plane systolic excursion (TAPSE) (D) and of pulmonary hypertension via pulmonary acceleration time (PAT) (E) in SU5416/hypoxia-treated rats. There were no differences in pulmonary ejection time among various experimental groups (data not shown). (F, G, and H) TGFBR11-Fc decreased the percentage of fully muscularized vessels (10–50 μ m diameter) and medial wall thickness, calculated as (external diameter – internal diameter)/external diameter \times 100, based on 50 to 85 vessels per treatment group from 5 to 6 rats each. (I and J) TGFBR11-Fc also reduced the frequency of Ki67-positive cells in the intima and adventitia (indicated by red arrows) seen in SU5416/hypoxia-treated rats. Data are expressed as mean \pm SEM (n = 5–6). * $P < 0.05$; ** $P < 0.01$; *** $P < 0.001$ as indicated. Scale bars = 50 μ m. H&E = hematoxylin and eosin; Hx = hypoxia; LV = left ventricle; RV = right ventricle; S = septum.

In summary, these data demonstrate direct contributions of TGF- β 1/3 to PH and remodeling, showing that the receptor fusion protein TGFBR11-Fc acts as a selective ligand trap to attenuate vascular remodeling and hemodynamics, and improve survival in

established disease. Antagonism of specific TGF- β ligands may be an effective and tolerable approach to mitigating PAH and other disease processes that arise from the profibrotic and promyogenic effects of TGF- β ligands. ■

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