Glycogen Synthase Kinase 3beta Contributes to Proliferation of Arterial Smooth Muscle Cells in Pulmonary Hypertension

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Abstract

Rationale: Pulmonary arterial hypertension (PAH) is a rare progressive pulmonary vascular disorder associated with vascular remodeling and right heart failure. Vascular remodeling involves numerous signaling cascades governing pulmonary arterial smooth muscle cell (PASMC) proliferation, migration and differentiation. Glycogen synthase kinase 3beta (GSK3β) is a serine/threonine kinase and can act as a downstream regulatory switch for numerous signaling pathways. Hence, we hypothesized that GSK3β plays a crucial role in pulmonary vascular remodeling.

Methods: All experiments were done with lung tissue or isolated PASMCs in a well-established monocrotaline (MCT)-induced PAH rat model. The mRNA expression of Wnt ligands (Wnt1, Wnt3a, Wnt5a), upstream Wnt signaling regulator genes (Frizzled Receptors 1, 2 and secreted Frizzled related protein sFRP-1) and canonical Wnt intracellular effectors (GSK3β, Axin1) were assessed by real-time polymerase chain reaction and protein levels of GSK3β, phospho-GSK3β (ser 9) by western blotting and localization by immunohistochemistry. The role of GSK3β in PASMCs proliferation was assessed by overexpression of wild-type GSK3β (WT) and constitutively active GSK3β S9A by [3H]-thymidine incorporation assay.

Results: Increased levels of total and phosphorylated GSK3β (inhibitory phosphorylation) were observed in lungs and PASMCs isolated from MCT-induced PAH rats compared to controls. Further, stimulation of MCT-PASMCs with growth factors induced GSK3β inactivation. Most importantly, treatment with the PDGFR inhibitor, Imatinib, attenuated PDGF-BB and FCS induced GSK3β phosphorylation. Increased expression of GSK3β observed in lungs and PASMC isolated from MCT-induced PAH rats was confirmed to be clinically relevant as the same observation was identified in human iPAH lung explants. Overexpression of GSK3β significantly increased MCT-PASMCs proliferation by regulating ERK phosphorylation. Constitutive activation of GSK3β (GSK3β S9A, 9th serine replaced to alanine) inhibited MCT-PASMCs proliferation by decreasing ERK phosphorylation.

Conclusion: This study supports a central role for GSK3β in vascular remodeling processes and suggests a novel therapeutic opportunity for the treatment of PAH.

Introduction

Pulmonary arterial hypertension (PAH) is a progressive pulmonary vascular disorder with high morbidity and mortality [1,2]. The pulmonary vascular remodeling may typically involve numerous molecular signaling cascades governing endothelial dysfunction, neovascularization of small pulmonary arteries, pulmonary arterial smooth muscle cell (PASMC) and adventitial fibroblast (PAAF) migration and proliferation [3,4]. However, 3 of the currently approved therapies targeting vasoconstrictive/vasodilatory abnormalities in PAH, like endothelin, nitric oxide or prostacyclin, show beneficial effects and improved quality of life [5,6,7], but do not appear to reverse or modify disease progression.

To this end, in the last 5 years, enormous progress has been made by our group and authors in developing new anti-proliferative and pro-apoptotic therapeutic strategies to reverse the disease progression. The role of growth factors, tyrosine/threonine kinase receptors in PAH has been extensively studied at the cellular, preclinical and clinical levels. Accumulation of growth factors such as platelet derived growth factor (PDGF),
Materials and Methods

**Ethics Statement**

The study protocol for human tissue donation was approved by the ethics committee (Ethik Kommission am Fachbereich Humannmedizin der Justus Liebig Universität Gießen) of the University Hospital Gießen (Gießen, Germany) in accordance with national law and with “Good Clinical Practice/International Conference on Harmonisation” guidelines (AZ 31/93). A written informed consent was obtained from each individual patient or the patient’s next of kin.

The University Animal Care Committee and the Federal Authorities approved all animal studies for Animal Research of the Ethik Kommission am Fachbereich Humannmedizin der Justus Liebig Universität Gießen (Hessen, Germany) (Az. GI 20/10 Nr. 09/2010).

**Materials**

GSK3β total, phospho-GSK3β (ser 9), AKT, phospho-AKT antibodies were purchased from Cell Signaling (Denver, USA). Axin1, phospho-ERK1/2, ERK total and H1 histone antibodies were obtained from Santa Cruz (Heidelberg, Germany). GAPDH antibody was obtained from Novus (Littleton, USA). Dulbecco’s Modified Eagle’s medium, nutrient mixture F-12 (DMEM-F12), fetal calf serum (FCS), Streptomycin/Penicillin, penicillins, and non-essentials amino acids were obtained from Gibco (Karlsruhe, Germany). Platelet derived growth factor BB (PDGF-BB) was purchased from PeproTech (Hamburg, Germany). Wnt3A ligand was purchased from R&D Systems (Minneapolis, USA). PDGFRβ inhibitor Imatinib (Gleevec) was obtained from Novartis (Switzerland). Im Prom reverse transcriptase and Taq polymerase PCR Kit were obtained from Promega (Mannheim, Germany). RIPA buffer was obtained from Santa Cruz (Heidelberg, Germany). ECL detection kit was obtained from Amersham Biosciences (Piscataway, USA). Histostain SP Rabbit Primary (AEC) was purchased from ZYMED Laboratories (Carlsbad, USA). MaxiPrep Plasmid Isolation Kit was obtained from PeqLab (Erlangen, Germany) and MaxiPrep Plasmid Isolation Kit, PCR purification Kit was purchased from QIAGEN (Hilden, Germany), pGEMT-Easy vector Kit was obtained from Promega, (Mannheim, Germany), pcDNA3.1 Directional TOPO Expression Kit, Lipofectamine 2000, T4 Ligase, Platinum Taq DNA Polymerase High Fidelity, Platinum SYBR Green qPCR SuperMix-UDG was purchased from Invitrogen (Karlsruhe, Germany), restriction enzymes (BamHI, BsmI, EcoRV) was purchased from New England Biolabs (Frankfurt, Germany). Gel Extraction Kit was obtained from QIAGEN (Hilden, Germany), Opti-DMEM medium was purchased from Gibco (Karlsruhe, Germany).

**Animal experiments**

Experiments were performed on male CD rats of 300–350 g body weight (Charles River, Sulzfeld, Germany). Pulmonary arterial hypertension was induced by a single subcutaneous injection of monocrotaline (MCT, 60 mg/kg, Sigma, Deishofen, Germany), dissolved in 0.1 M NaOH, adjusted to pH 7.4 with 0.1 M HCl, as described previously [24]. In these experiments 3 study groups were used: healthy rats, MCT-injected rats sacrificed either 3 weeks or 5 weeks after injection.

**Pulmonary arterial smooth muscle cells isolation**

Rat PASMCs were isolated from pulmonary arteries using the explant method as described previously [8] and the characterization and purity was assessed as described [25]. Briefly, characterization of PASMCs was done using immunocytochemical staining for a-smooth muscle actin and desmin. The presence of endothelial cells and fibroblasts were excluded by staining for CD31 and FSP1. Further in order to determine the purity of PASMCs, flow cytometric analysis was performed with anti-desmin and anti-a-smooth muscle actin antibodies. Usually, the purity of isolated PASMCs is 91–94%. Cultures were maintained at 37°C in a humidified 5% CO2/95% O2 atmosphere. All experiments using PASMCs were performed between 2–4 passages. Expression analysis studies were done at early passages to minimize the influence of phenotypic alterations.

**Cell culture**

Primary rat PASMCs were maintained at 37°C in a humidified 5%CO2/95%O2 atmosphere. Cells were grown on 10 cm2 dishes, 6 and 48-well plates in DMEM-F12 supplemented with 10% FCS, 5% streptomycin/penicillin, 5% glutamate. Cells were cultured from the time of isolation in 10% FCS medium and subsequently in experiments with 0.1% FCS medium supplemented with PDGF-BB (60 ng/ml) or Wnt3A (100 ng/ml) for 6 and 24 hrs. Additionally, PASMCs were stimulated with PDGF-BB (60 ng/
Reverse transcriptase PCR and Real-Time PCR

Total RNA was isolated from tissue rat’s lung homogenates and primary rat pulmonary arterial smooth muscle cells (PASMC) with Trizol® Reagent. The quantity and quality of RNA was determined by NanoDrop (PeqLab, Erlangen, Germany). Equal amounts of RNA from each sample were used as templates for reverse transcription reaction for generation of cDNA using Im Prom Reverse Transcriptase and Taq polymerase PCR Kit with oligo(dT)₁₈ primers according to the supplier’s instructions. Later, quantitative Real-Time PCR analysis was performed as described previously [26]. Briefly, 1 µl cDNA was placed into 25 µl reaction volume containing Platinum SYBR Green qPCR SuperMix-UDG and sequence-specific oligonucleotide primers. The thermal cycle conditions used for all reactions were as follows: activation, 50°C for 2 min; denaturation, 95°C for 10 min; and cycle, 95°C for 10 s and 99°C for 30 s (40 times). Specific primers used for sequence detection both in tissue homogenates as well as in the cells were: Wnt1 (Forward 5’ GTC ACT TGT ATC AGG ACC ACA3’, Reverse 5’ ATG TCC GGT CTT CAC TGC TCA3’), Wnt3a (Forward 5’ ATT TGG AGG AAT GGT CTC TCG3’, Reverse 5’ GCA GGT CTT CAC TTC GCA AC3’), Wnt5a (Forward 5’ GCC ACT TGT ATC AGG ACC ACA3’, Reverse 5’ GGC ATT TAG CAC TCC AGC AG3’), sFRP-1 (Forward 5’ GCT AGA GAG GAC CCC TGA AAA T3’, Reverse 5’ TCG CCA CTC GAG TAG AAG AAA3’), Axin1 (Forward 5’ GTA TCC TTC ATC GGC ACA TC3’, Reverse 5’ TGA GCC ACT TGT AGT TGTT TG3’), Reverse 5’ ATG TCC GGT AAG GGC GGC3’), PBGD (Forward 5’ TGC ACT GAG TGG AGT GTG TTT TG3’, Reverse 5’ TGC ACT TGG CAC TGG AGC AG3’), Frizzled 1 receptor (Forward 5’ GTA TCC TTC CCA GGT TCT CTC3’, Reverse 5’ GGT TAG AGC ACG GAG AAG AGC3’), Frizzled 2 receptor (Forward 5’ GTG TAG TGG ACG AGC GAG AAG AGC3’, Reverse 5’ TCT GCT TTT ATG AGC GCA ACT3’), Axin1 (Forward 5’CCA CAG AAA TAG TAG GGC ACA3’, Reverse 5’ GGA GGA AGA AGA AAA GAG AGC3’), GSK3β (Forward 5’ TCG CCA CTC GAG TAG AAG AAA3’, Reverse 5’ ACT TTG TGA CTC AGG AGA ACT3’), PBGD (Forward 5’ CAA GGT TTT CAG CAT CGC TAC3’, Reverse 5’ ATG TCG GGT AAG GGC GGC3’).

Protein isolation and Western blotting

Lung tissue and PASMC’s samples were homogenized using tissue homogenizer (for tissues only) or lysed in RIPA buffer (Santa Cruz) with addition of a protease inhibitor cocktail and PMSF. Tissue and cell lysates were equalized with SDS 5× sample buffer and electrophoretically separated on 10% polyacrylamide gels and transferred for 1 h on to nitrocellulose membranes. Subsequently membranes were blocked 1 h with 5% non-fat dry milk in Tris-buffered saline/0.1% Tween 20. After blocking, membranes were probed with primary antibodies diluted as follows: GSK3β total (1:1000), phospho-GSK3β (ser 9) (1:1000), AKT (1:1000), phospho-AKT (1:1000), Axin1 (1:500), phospho-ERK1/2 (1:1000), ERK total (1:1000), GAPDH (1:5000). Samples were normalized to GAPDH housekeeping gene and total protein of interest (when examining protein phosphorylation) and further quantification was performed. After primary antibody incubation, membranes were incubated with secondary goat anti-rabbit antibody (1:30000) or rat anti-mouse (1:50000) HRP-conjugated antibodies (Sigma). Signals were then detected by ECL detection system and further quantified using specific software as described [27].

Immunohistochemistry

Paraffin-embedded lung sections were cut to 3 µm thickness by microtome and incubated 45 minutes at 65°C and subsequently deparaffinized in xylene three times, each 3 minutes. Samples were hydrated with 100% to 80% ethanol concentration washed with water and the antigen was unmasked by microwaving 20 minutes in the retrieval buffer. Sections were incubated in 3% hydrogen peroxide in methanol for 20 minutes at room temperature to block endogenous peroxidase activity, washed and blocked for 30 minutes in 2% bovine serum albumin. Then sections were probed overnight at 4°C with primary anti-rabbit antibody (GSK3β, Cell Signaling, 1:50 dilution). After overnight incubation sections was washed with PBS and incubated with secondary biotinylated horse anti-rabbit IgG Antibody (Vector Laboratories) for 30 min, followed by incubation with the ABC reagent kit (avidin and biotinylated peroxidase, Vector Laboratories) for 30 min. Development of the target-bound peroxidase for detection of GSK3β was carried out with Vector NovaRed.
substrate kit for peroxidase according to manufacturer’s instruction (Vector Laboratories). Finally, sections were counterstained with haematoxylin (Zymed Laboratory, Cambridge, UK), mounted with VectaMount (Vector Laboratories) and visualized by light microscopy (bright-field microscopy).

**Cloning**

Human GSK3β wild type full length insert was obtained on the human PASMCs cDNA template by PCR reaction using Platinum Taq DNA Polymerase High Fidelity with the following primers: Forward 5′CCT AAC ACC CCA AGA TAA AGA CA3′, Reverse 5′GTA ACT GGT GGT TTT TCC TGT GC3′. Human GSK3β WT insert was subsequently ligated into the pGEMT-Easy Vector and then electroporated to the DH5α competent strain of E.Coli, becoming the template for following active GSK3β mutant construction. Later GSK3β full length and mutant constructs was transferred to mammalian pcDNA3.1 expression vector TOPO cloning system and transformed to JM-109 competent E.Coli strain following heat shock procedure. After transformation constructs were digested with BamHI and verified on a 0.7% agarose gel. Pure plasmid GSK3β constructs were obtained using MaxiPrep Plasmid Isolation Kit in extensive amounts for subsequent transient transfection and further sent for sequencing to AGOWA GmbH sequencing service (Berlin, Germany).

**Mutagenesis and Sequencing**

Wild type pGEMT- GSK3β was used as a template for creating mutants of GSK3β which are S9A- constitutively active mutant of GSK3β where serine 9 residue was substituted with alanine. To prepare constitutively active constructs of GSK3β, point mutations were done with long template PCR technique with use of following mutagenesis primers: GSK3β S9A (Sense 5′GGC CCA GAA CCA CGG CAT TCG CGG AGA GCT GCA A3′; Antisense 5′TTG CAG CTC TCC GCG AAT GCG GTG GTT CTG GGC C3′) in the following set up (95°C-90 s; 95°C-30 s; 62°C-30 s; 68°C-900 s; 68°C-1200 s; 4°C). Digestion of methylated template was done by 3 h incubation in 37°C with use of DpnI.

**Figure 2. Increased GSK3β and its phosphorylated form in MCT-induced PAH rat lungs.** Protein expression as analyzed by (A) western blotting and subsequent (B) densitometric quantification of GSK3β in control (white bar) and after 3 weeks (grey bar) and 5 weeks (black bar) of MCT-induced PAH in rats. Phosphorylation analysis by (C) western blotting and subsequent (D) densitometric quantification of pGSK3β (serine 9) in control (white bar) and 5 weeks (black bar) lungs of MCT-induced PAH in rats. GAPDH was used as a loading control. Values were presented significant as **P<0.01, ***P<0.001 vs control lungs. All values were expressed as mean ± SEM (n = 3). (E) Immunohistochemical localization of GSK3β in the healthy lungs (a) and lungs 5 weeks after MCT injury (b). Magnification 40×.

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enzyme. Subsequently precipitated plasmid was transformed to JM-109 strain of E.Coli and produced in extensive amounts for subsequent transfections.

Transient transfection of PASMCs with pcDNA3.1 TOPO-GSK3β construct

PASMCs were plated in 6 or 48-well plates until they reach 60 to 80% confluence at the time of transfection. Empty pcDNA3.1 vector and pcDNA3.1-GSK3β construct (GSK3β WT, S9A) were mixed with Opti-DMEM F12 medium and combined with Lipofectamine 2000 transfection reagent and added to cultured MCT PASMCs for 6 hrs as described [28]. After this time cells were washed with PBS and stimulated with 10% FCS to allow cell recovery for next 18 hrs and later depending on further assays adequately stimulated with 0.1 or 10% FCS supplied DMEM F12 medium.

Proliferation Assay

For assessment of proliferation, rat PASMCs from passage 2 were seeded in 48-well plates. Primary cells were starved by incubation for 24 hrs in DMEM containing 0.1% FCS. Subsequently, they were stimulated with 10% FCS/DMEM or PDGF-BB (60 ng/ml) to induce cell cycle reentry. During the last 4 hrs of the stimulation period, cells were pulsed with 1.5 μCi per well [3H] thymidine (Amersham Pharmacia Biotech Ltd). The [3H] thymidine content of cell lysates was determined by scintillation counting as described previously [8]. Data were expressed as counts per minute (cpm) and normalized to the amount of cells per well. In our study all of [3H] thymidine uptake experiments were carried out on 48-well plates and the amount of PASMCs seeded on one well was around 30,000.

Patient characteristics

Human lung tissue was obtained from donors and iPAH patients undergoing lung transplantation. Donor lung tissue was from non-transplanted lung tissue of transplant donors. Donor lungs were explanted according to a standard European explant protocol (Eurotransplant), using cold perfused with preservation buffer and stored inflated on ice until use. Non-transplanted donor lung was not transplanted because i) the complete lung would not

**Figure 3. Increased GSK3β and its phosphorylated form in primary PASMCs isolated from lungs of control and MCT-induced PAH rats.**

(A) Proliferation capacity of primary rat MCT-PASMCs compared to healthy control-PASMCs isolated from rat lungs 5 weeks post MCT injury in 10% FCS conditioned media was assessed by [3H]-thymidine incorporation (n = 5). Data were obtained as counts per minute (cpm) and normalized to the amount of cells per well. All values were expressed as the percentage of proliferation capacity (mean ± SEM). Values were presented significant as ***P<0.001 vs control. (B) mRNA expression of Wnt1, Wnt5a (not expressed), Wnt5a, Frizzled 1, Frizzled 2, sFRP-1, Axin 1 and GSK3β in primary control PASMCs and PASMCs from MCT-induced PAH rats isolated from the lungs 5 weeks after MCT injection, as analyzed by quantitative real-time PCR. Rat PASMCs were maintained in culture media supplemented with 10% FCS. All values were normalized to Porphobilinogen deaminase (PBGD) and were presented as fold of gene regulation with a control set as 1. Values were presented significant as **P<0.01 vs control PASMCs. All values were expressed as mean ± SEM (n=4). Protein expression as analyzed by (C) western blotting and subsequent (D) densitometric quantification of total GSK3β/GAPDH, phosphorylation of GSK3β at serine 9 residue (pGSK3β S9/total GSK3β) in primary PASMCs isolated from control (grey bar) and MCT-induced PAH rats (black bar). GAPDH was used as a loading control. All values were expressed as mean ± SEM (Control PASMCs, n = 4; MCT-PASMCs n = 5). Values were presented significant as ***P<0.001 vs control PASMCs. Both mRNA and protein were isolated from healthy control- and MCT-PASMCs at passage 3 in 10% FCS conditioned media.

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fit into the recipient thorax and a part was resected or ii) due to on site decision of the transplant surgeon not to use the lung, based on edema, pulmonary thrombi or obvious pneumonia. In either case, lung tissue was snap-frozen directly after transplantation.

Results

GSK3β is dysregulated in total lung homogenates of monocrotaline-induced pulmonary arterial hypertension in rats

Expression of Wnt ligands (Wnt1, Wnt3a, and Wnt5a) and Wnt signaling upstream regulator genes (Frizzled1 and 2 receptors and sFRP-1) and intracellular effectors (Axin1 and GSK3β) were investigated in pulmonary hypertensive rat lungs, 3 and 5 weeks after MCT injury. Real-time RT-PCR demonstrated a decrease in Wnt canonical ligands (Wnt1 and Wnt3a) with no significant changes in Frizzled receptors expression in 5 weeks of MCT-PAH rat lungs (Figure 1). Parallel increase in mRNA expression of GSK3β and Axin1 was observed (Figure 1). Western blotting demonstrated a significant increase in total GSK3β expression in MCT lungs 3 and 5 weeks post MCT injury (Figure 2A, B) as well as significant increase in its phosphorylation in 5 weeks MCT-PAH rat lungs (Figure 2C, D). However, phosphorylation of GSK3β was not altered 3 weeks after MCT treatment (data not shown). Immunohistochemical analysis suggested an immunoreactivity of GSK3β was observed in the medial layer of pulmonary arteries of both 5 weeks MCT-PAH rat lungs and control PAH rat lungs (Figure 2E).

Increased GSK3β and its phosphorylated form in pulmonary arterial smooth muscle cells isolated from MCT-PAH rats

Interestingly PASMCs isolated from the rat lungs 5 weeks after MCT injury displays much stronger proliferative phenotype as compared to healthy control-PASMCs in 10% FCS condition (Figure 3A) and PDGF-BB (60 ng/ml) supplemented media (Figure S1). Therefore, we tested the expression profile of GSK3β-related genes (Figure 3B) in these hyper-proliferative PASMCs. Similarly to the mRNA level in the lungs after MCT injury we found a decrease in canonical Wnt1 (Figure 3B). On the other hand, reduction in Frizzled receptor 2 is significant in isolated MCT-PASMCs, but not in MCT injured lungs as compared to healthy controls (Figure 3B). Additionally, we observed discrepancies in Axin1 mRNA expression level which was significantly upregulated in lungs 5 weeks after MCT injury, but its mRNA level was significantly reduced in MCT-PASMCs. Downregulation of upstream regulators of GSK3β and GSK3β itself (Figure S2), on mRNA level, in MCT-PASMCs as compared to healthy controls (Figure 3B) suggests dysregulation of canonical Wnt signaling pathway. Interestingly, further analysis of GSK3β and its phosphorylated form at serine 9 residue in primary isolated pulmonary arterial smooth muscle cells (PASMCs) revealed an increase in total protein expression of GSK3β and significant increase in its phosphorylated (inactive) form in PASMCs isolated from MCT-PAH rats compared to control rats (Figure 3C, D).

Growth factors signaling regulates GSK3β phosphorylation in primary rat PASMCs

To investigate upstream regulators of GSK3β, quiescent PASMCs were stimulated with PDGF-BB (60 ng/ml) for 6 (Figure S3) and 24 hrs (Figure 4) or 10% FCS (Figure 5) for 24 hrs. In addition, to verify the specificity of PDGF-BB mediated signaling, primary rat PASMCs were also simultaneously treated with Imatinib (1 and 5 μM). Proteins from untreated and stimulated cells were isolated and western blotting for AKT, GSK3β, phospho-AKT and phospho-GSK3β ser 9 and ERK and phospho-ERK were performed.

Interestingly, in all cases PDGF-BB and 10% FCS stimulation caused a significant increase in phospho-AKT and phospho-GSK3β [Ser 9] in combination with significant activation of ERK (Figure 4A–C, 5A–C, Figure S3). The increased phosphorylation of AKT, GSK3β at ser 9 site and ERK observed after stimulation with growth factors were significantly abrogated by Imatinib treatment (Figure 4A–C, 5A–C, Figure S3). Decreased phosphorylation status of AKT, GSK3β and ERK after Imatinib treatment were accompanied with significant decrease in proliferation capacity of these cells after 24 hrs stimulation with PDGF-BB as well as 10% FCS (Figure 4D, 5D). This suggest that GSK3β is a possible crucial player in growth factors signaling and abnormal proliferation of PASMCs in experimental PAH.

Overexpression of GSK3β and phosphorylation deficiency of GSK3β influences PASMCs proliferation

To further evaluate the contribution of GSK3β to vascular remodeling processes, we generated pcDNA 3.1 TOPO-cloning constructs carrying GSK3β wild type (WT) cDNA and GSK3β S9A (where serine at residue 9 is replaced to alanine) mutant. Sequence analysis performed on each construct confirmed the intended point mutations at specific sites in GSK3β S9A (data not shown).

Transient overexpression of GSK3β WT and GSK3β S9A mutant in PASMCs resulted in a significantly enhancement of GSK3β expression 24 hrs post transfection. Empty vector (EV) transfection caused no change in GSK3β protein expression (Figure 6A). Importantly, serum stimulation did not cause GSK3β phosphorylation in GSK3β S9A mutant overexpressing cells while another tyrosine 216 residue was phosphorylated (Figure 6A). The gain of GSK3β WT expression was coupled to a significantly increase of [3H]-thymidine uptake as compared to cells transfected with either empty pcDNA3.1 vector or Lipofectamine 2000 reagent (LF2000) only, 24 hrs post serum stimulation (Figure 6B). On the other hand, GSK3β S9A mutant expression decreased [3H]-thymidine uptake, suggested that GSK3β inactivation is involved in PASMCs proliferation (Figure 6B).

Overexpression of GSK3β and phosphorylation deficiency of GSK3β influences proliferation via ERK phosphorylation

Transient overexpression of GSK3β WT significantly increased ERK phosphorylation compared to empty pcDNA3.1 vector transfection (Figure 7A, B). In contrary, GSK3β S9A mutant
decreased ERK1/2 phosphorylation, in serum induced proliferation of PASMCs (Figure 7A, B).

**Increase in GSK3β protein levels in human PAH lung explants**
Expression of GSK3β was investigated also in iPAH patients lung explants. While mRNA levels of GSK3β was not regulated in iPAH patient lungs (Figure S4), western blotting analysis showed significant upregulation of total GSK3β protein in iPAH lung homogenates compared to healthy donor lung homogenates (Figure 8A, B).

**Discussion**
This study has 4 salient features. First, GSK3β and the phosphorylated form of GSK3β (inactivation) are increased in

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Figure 5. Serum regulates GSK3β, Akt, ERK phosphorylation and proliferation in primary rat MCT-PASMCs. (A) Western blot analysis and subsequent (B, C) quantification of Akt, GSK3β, ERK and phosphorylation status in primary rat MCT-PASMCs stimulated with 10% FCS in combination with two doses of Imatinib (1 and 5 μM) for 24 hrs. GAPDH was used as reference loading control. (D) Proliferation of primary rat MCT-PASMCs was assessed by [3H]-thymidine incorporation (n = 6). Data were expressed as counts per minute (cpm) and normalized to the amount of cells per well. All values were expressed as mean ± SEM. Values were presented significant as *** P<0.001 vs control, ††† P<0.001 vs 10% FCS. doi:10.1371/journal.pone.0018883.g005

Figure 6. GSK3β overexpression and S9A modification influences proliferation of primary rat PASMCs. (A) Primary rat PASMCs were transiently transfected with GSK3β wild type (GSK3β WT), constitutively active GSK3β (GSK3β S9A), empty vector (pcDNA3.1 TOPO). 24 hrs post transfection, expression and phosphorylation of GSK3β were analyzed by western blotting. (B) Primary rat PASMCs transiently transfected with GSK3β wild type (GSK3β WT), constitutively active GSK3β (GSK3β S9A), empty vector (pcDNA3.1 TOPO) and proliferation was assessed after stimulation with 10% FCS for 24 hrs by [3H]-thymidine incorporation. All values are expressed as mean ± SEM (n = 6, WT or S9A n = 12). Data were expressed as counts per minute (cpm) and normalized to the amount of cells per well. Values were presented significant as ***P<0.001 vs 0.1% Empty Vector pcDNA3.1 and ††† p<0.001 vs GSK3β WT transfected cells. doi:10.1371/journal.pone.0018883.g006
MCT lungs and PASMCs isolated from monocrotaline induced pulmonary arterial hypertensive rats compared to control rats. Second, both PDGF and FCS stimulation induced GSK3β phosphorylation (inactivation) in PASMCs. Third, treatment with the PDGFR inhibitor, Imatinib, attenuated growth factors-induced GSK3β and ERK phosphorylation. Fourth, overexpression of wild type and constitutively activate form of GSK3β (GSK3β S9A; 9th serine replaced to alanine) influenced serum induced MCT-PASMCs proliferation by regulating ERK phosphorylation. This study supports a central role for GSK3β in vascular remodeling processes and suggests a novel therapeutic opportunity for the treatment of pulmonary hypertension.

The dynamic process of pulmonary vascular remodeling involves numerous molecular signaling cascades governing PASMCs proliferation, differentiation and migration [4,29]. We hypothesized that GSK3β signaling plays a critical regulatory role in pulmonary vascular remodeling. In accord with our hypothesis, we demonstrated significant increase in total GSK3β expression levels as well as its phosphorylation status in lungs and PASMCs in response to MCT injury and total GSK3β in human IPAH lung explants. Interestingly, the increase in GSK3β and its phosphorylated form, observed in primary PASMCs isolated from MCT-PAH rats as well as in rat and human lungs, suggest that aberrant GSK3β signaling may trigger the proliferative phenotype of PASMCs. In line with this notion, aberrant GSK3β signaling was recently implicated in various vascular- and fibro-proliferative diseases [23,30,31].

GSK3β remains increased and inactivated 5 weeks after MCT injury, which is accompanied by decreases in mRNA expression of canonical Wnt ligands (Wnt1 and Wnt3a) and an increase in non-canonical Wnt5a ligand, 3 and 5 weeks after MCT injury. However, GSK3β was downregulated at the mRNA level in MCT-PASMCs compared to control-PASMCs. We assume that downregulation of GSK3β mRNA concurrent with upregulation of protein expression may occur when a protein’s half-life is increased due to stabilization-components involved with the protein’s normal turnover may be disrupted. It is also possible that the protein may become stabilized through protein-protein interactions. Additionally, this can be explained by transcriptional repression when high levels of proteins accumulate. Furthermore as delineated in our study, GSK3β phosphorylation is regulated by growth factors, i.e. PDGF-BB that were highly upregulated in PAH. Hence we believe that GSK3β is clearly dependent on multiple lines of regulation in addition to the phosphorylation state of GSK3β [32,33,34].

In our study, upregulation of non-canonical [35,36] Wnt5a mRNA in MCT-lungs is in line with previously published data, showing involvement of non-canonical Wnt signaling in human IPAH [37]. Moreover, downstream targets of the non-canonical Wnt pathway, like Rho-kinases or calcium signaling, were shown to significantly contribute to PASMCs proliferation and vasoconstriction and demonstrated therapeutic potential in pulmonary hypertension [38,39,40].

Our data indicate that MCT-PASMCs are hyper-proliferative in the presence of FCS and PDGF compared to control-PASMCs. Interestingly, our results showed that PDGF and FCS stimulation on MCT-PASMCs, acting via PI3-kinase-dependent activation of AKT, causes GSK3β (Ser 9) phosphorylation and GSK3β inactivation followed by ERK activation, which is potentially suppressed by Imatinib. A similar role of growth factors, such as PDGF, IGF and EGF, mediated GSK3β inactivation was described previously [41,42]. Furthermore, several kinases were also shown to be capable of mediating ser 9 phosphorylation, including AKT, PKA, PKC and Wnts [16,32,43]. Considering the crucial role of PDGF signaling in pulmonary vascular remodeling [8,9], the increased presence of growth factors signaling in human and experimental PAH [8,10,13] and PDGF and FCS mediated alteration of GSK3β activity, collectively suggests GSK3β plays an important role in the pathogenesis of PAH.

The effects of GSK3β are also regulated by Wnt signaling pathway protein complex formation, a process involved in modulating β-Catenin levels [18,44]. Future studies are needed to study the regulation of canonical Wnt signaling and the multitude of factors regulating β-Catenin expression in the pathogenesis of pulmonary hypertension. Albeit recent studies suggest that recruitment of both canonical and non-canonical Wnt pathways promote pulmonary arterial endothelial cell proliferation, survival, and migration. In addition, it was demonstrated that both canonical and non-canonical Wnt pathways are required for BMP-2-mediated angiogenesis in...
severe combined immunodeficient (SCID) mice [45]. These findings may help better understand the pathogenesis of pulmonary hypertension, a disease that featured with the loss of small precapillary arteries.

In the present study, overexpression of GSK3β significantly increased expression of GSK3β that was accompanied by increased proliferation capacity of MCT-PASMCs proliferation after FCS stimulation. Constitutive activation of GSK3β significantly reduced expression of phospho-GSK3β and PASMCs proliferation. This effect can be due to GSK3β phosphorylation of a diverse group of substrates or by inhibition of transcription factor activation such as p53, CREB and β-Catenin [17,18,19]. In addition, GSK3β constitutively activation (S9A) has also been shown to directly effects cyclin D1 expression, independent of β-Catenin [46]. Here, we show that overexpression of wild-type GSK3β significantly influenced the proliferation capacity of MCT-PASMCs via regulating phosphorylation of ERK. In a recent study by Wang et al., it was shown that GSK3β acts as a negative regulator of ERK in human colon cancer cells [47]. Our study shows that constitutive activation of GSK3β significantly reduced phospho-GSK3β levels and PASMCs proliferation that was accompanied by a significant decrease in ERK phosphorylation. These results collectively suggest that modification of GSK3β can significantly influence highly dysregulated growth factors signaling associated with abnormal proliferation of PASMCs in PAH.

Although mRNA levels of GSK3β is downregulated in MCT-PASMCs and in human iPAH patient lungs, along with other canonical Wnt signaling dependent genes we found that protein levels of GSK3β increased in a time dependent fashion with disease progression in the MCT-induced PAH model (lungs and MCT-PASMCs) and also in explanted iPAH patient lungs, suggesting a role for GSK3β in disease progression, potentially independent of canonical Wnt signaling. Recently a crucial role for GSK3β in systemic vascular remodeling was reported [48,49]. Authors demonstrated for the first time in vivo that active GSK3β gene transfer results in a significant reduction in neointima formation in the restenosis model of balloon injury in rat carotid arteries. These effects were attributable, at least in part to the ability of GSK3β to inhibit smooth muscle proliferation and to promote sustained apoptosis. This concept was extended to demonstrate that GSK3β plays a significant role in VSMCs proliferation and apoptosis in vascular remodeling after balloon injury [49]. Similarly, in our study the inactivation of GSK3β by serine 9 phosphorylation was observed 5 weeks after MCT injury in rats, both in PASMCs and lung homogenates. This suggest that the introduction of active GSK3β (S9A) may also prove beneficial for regression of vascular remodeling in experimental PAH.

![Figure 8. Increased GSK3β and its phosphorylated form in human lungs of healthy donor and iPAH patients. (A) Protein expression as analyzed by western blotting and subsequent (B) densitometric quantification of GSK3β and in human lungs of healthy donor and iPAH patients. GAPDH was used as a loading control. Values were presented significant as ***P<0.001 vs control lungs. All values were expressed as mean ± SEM (n=7). doi:10.1371/journal.pone.0018883.g008](image-url)
To our knowledge this is the first study to demonstrate that GSK3β is significantly altered in the pathogenesis of experimental as well as human PAH and the regulatory role for GSK3β in pulmonary arterial smooth muscle cell proliferation. This study supports a central role for GSK3β in vascular remodeling processes and suggests a novel therapeutic opportunity for the treatment of pulmonary arterial hypertension.

Supporting Information

Figure S1 MCT-PASMCs display significant increase of PDGF-BB-induced proliferation capacity as compared to healthy control-PASMCs. Proliferation capacity of primary rat MCT-PASMCs compared to healthy control PASMCs isolated from rat lungs 5 weeks post MCT injury in 10% FCS conditioned media was assessed by [3H]-thymidine incorporation (n = 5). Data were obtained as counts per minute (cpm) and normalized to the amount of cells per well. All values were expressed as percentage of proliferation capacity (mean ± SEM). Values were presented significant as *** P<0.001 vs control. (TIF)

Figure S2 Expression of GSK3β in rat MCT-PASMCs. mRNA expression of GSK3β in MCT-PASMCs after 5 weeks of MCT-induced PAH rats, as analyzed by quantitative real-time PCR. All values were given as the mean ± SEM (n = 3) and were normalized to Porphobilinogen deaminase (PBGD). Values were presented significant as * P<0.05, ** P<0.01, *** P<0.001 vs PASMCs isolated from healthy rat lungs. Healthy controls were set as 1 on X axis and expression profile from 5 weeks MCT-PASMCs were presented as fold of gene regulation. (TIF)

Figure S3 PDGF regulates GSK3β, Akt and ERK phosphorylation in primary rat MCT-PASMCs. (A) Western blot analysis and subsequent (B, C) quantification of Akt, GSK3β and ERK phosphorylation status in primary rat MCT-PASMCs stimulated with PDGF-BB (60 ng/ml) alone or in combination with two doses of Imatinib (1 and 5 μM) for 6 hrs. All values were expressed as mean ± SEM (n = 4). Values were presented significant as *** P<0.001 vs control, ††† P<0.001 vs PDGF-BB. GAPDH was used as reference loading control. (TIF)

Figure S4 GSK3β is not significantly regulated in iPAH patient lungs on mRNA level. mRNA expression of GSK3β in donor lungs and iPAH patient lungs as analyzed by quantitative real-time PCR. All values were normalized to Porphobilinogen deaminase (PBGD) and determined as fold of gene regulation. All values were expressed as mean ± SEM (n = 5). (TIF)

Author Contributions
Conceived and designed the experiments: PS RTS FG WS SSP. Performed the experiments: PS XT. Analyzed the data: PS SSP. Contributed reagents/materials/analysis tools: DS NW HAG TK. Wrote the paper: PS RTS SSP.

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