

# TGF $\beta$ acts through PDGFR $\beta$ to activate mTORC1 via the Akt/PRAS40 axis and causes glomerular mesangial cell hypertrophy and matrix protein expression

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Interaction of transforming growth factor- $\beta$  (TGF $\beta$ )-induced canonical signaling with the noncanonical kinase cascades regulates glomerular hypertrophy and matrix protein deposition, which are early features of glomerulosclerosis. However, the specific target downstream of the TGFB receptor involved in the noncanonical signaling is unknown. Here, we show that TGFB increased the catalytic loop phosphorylation of plateletderived growth factor receptor  $\beta$  (PDGFR $\beta$ ), a receptor tyrosine kinase expressed abundantly in glomerular mesangial cells. TGFβ increased phosphorylation of the PI 3-kinase-interacting Tyr-751 residue of PDGFR $\beta$ , thus activating Akt. Inhibition of PDGFR<sup>B</sup> using a pharmacological inhibitor and siRNAs blocked TGFβ-stimulated phosphorylation of proline-rich Akt substrate of 40 kDa (PRAS40), an intrinsic inhibitory component of mTORC1, and prevented activation of mTORC1 in the absence of any effect on Smad 2/3 phosphorylation. Expression of constitutively active myristoylated Akt reversed the siPDGFRβ-mediated inhibition of mTORC1 activity; however, co-expression of the phospho-deficient mutant of PRAS40 inhibited the effect of myristoylated Akt, suggesting a definitive role of PRAS40 phosphorylation in mTORC1 activation downstream of PDGFRB in mesangial cells. Additionally, we demonstrate that PDGFRB-initiated phosphorylation of PRAS40 is required for TGFB-induced mesangial cell hypertrophy and fibronectin and collagen I ( $\alpha$ 2) production. Increased activating phosphorylation of PDGFRB is also associated with enhanced TGF $\beta$  expression and mTORC1 activation in the kidney cortex and glomeruli of diabetic mice and rats, respectively. Thus, pursuing TGFB noncanonical signaling, we identified how TGFB receptor I achieves mTORC1 activation through PDGFR\beta-mediated Akt/PRAS40 phosphorylation to spur mesangial cell hypertrophy and matrix protein accumulation. These findings provide support for targeting PDGFR $\beta$  in TGF $\beta$ -driven renal fibrosis.

Chronic kidney disease affects 10% of the world's population, including 20 million Americans, and causes increased risk of cardiovascular diseases and loss of renal function, leading to end-stage renal disease with significant public health costs (1, 2). Thus, understanding the progression of the disease process is of prime importance for its prevention and arrest. The pathologic correlates of progressive renal function impairment include renal structural changes with initial renal hypertrophy, especially glomerular hypertrophy, that leads to hyperfiltration and microalbuminuria, followed by accumulation of matrix proteins and a greater degree of proteinuria (3, 4). A significant pathologic characteristic of chronic kidney disease is glomerulosclerosis. One-third of the cell population in the glomerulus is made up of mesangial cells, which have the capacity to communicate with the endothelial cells; they can also regulate glomerular filtration because of their contractile nature (5, 6). Transforming growth factor- $\beta$  (TGF $\beta$ ) contributes to glomerulosclerosis and albuminuria during progressive kidney injury, especially in diabetic kidney disease (4, 7). Liver-specific TGF $\beta$ transgenic mice with increased circulating levels of cytokine developed mesangial expansion with augmented glomerular immune deposits and matrix proteins (8). In the disease milieu or in vitro when mesangial cells are exposed to TGFB, they undergo hypertrophy and acquire a myofibroblast-like phenotype to synthesize larger amounts of matrix proteins (9, 10). Also, anti-TGF $\beta$  antibody decreases mesangial and fibrotic protein expression in rodent models of nephropathy in type 1 and type 2 diabetes where TGF $\beta$  contributes to the pathology (11, 12).

The dimeric active TGF $\beta$  directly binds to TGF $\beta$  receptor II, which is a constitutively active kinase. After TGF $\beta$  binding, TGF $\beta$ RI is recruited into the complex and undergoes phosphorylation by TGF $\beta$ RII. Activated TGF $\beta$ RI then binds and phosphorylates the receptor-specific Smad, Smad 2, and Smad 3, which form a complex with a common Smad, Smad 4, for translocation to the nucleus and regulation of gene transcription (13). Apart from this canonical signaling, TGF $\beta$  also initiates other noncanonical signal transduction pathways involving mitogen-activated protein kinases, PI 3-kinase/Akt/mTORC1, and Rho GTPase, which contribute to the renal complications under disease conditions (13–17).

Among many hormones, growth factors, and cytokines, platelet-derived growth factor (PDGF) contributes to significant injury in different renal, especially glomerular, diseases (5). Four different isoforms of PDGF (A, B, C, and D) form a total of five homo- or heterodimers (18). These dimers bind to three

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**Figure 1. TGF** $\beta$  **receptor activates PDGFR** $\beta$ . *A*, TGF $\beta$  increases activating phosphorylation of PDGFR $\beta$  at Tyr-857. Mesangial cells were incubated with 2 ng/ml TGF $\beta$  for the indicated duration. The cleared cell lysates were immunoblotted with phospho-PDGFR $\beta$  (Tyr-857) and PDGFR $\beta$  antibodies. *B–J*, mesangial cells were treated with 5  $\mu$ M SB (*panels B–D*) or 0.1  $\mu$ M JNJ (*panels E–G*) or transfected with siRNAs against PDGFR $\beta$  or scrambled RNA (*panels H–J*) prior to incubation with 2 ng/ml TGF $\beta$  for 2 h. The cell lysates were immunoblotted against the indicated antibodies. Molecular weight markers are shown in the *left margins*.

dimeric PDGF receptors (PDGFRs),  $\alpha\alpha$ ,  $\beta\beta$ , and  $\alpha\beta$ , with variable affinity (19). For example, whereas PDGF AA and CC show affinity for PDGFR $\alpha$ , CC also binds to PDGFR $\alpha\beta$ . PDGF DD binds to PDGFR $\alpha\beta$  at a low affinity, but it interacts with PDGFR $\beta\beta$  with significantly higher affinity. On the other hand, PDGF AB and BB bind to both receptors. All these PDGF dimers and three dimeric receptors are expressed in mesangial cells (19). In glomerulonephritis, PDGF BB, CC, and DD play prominent roles and stimulate proliferation of mesangial cells, whereas PDGF AA does not (5, 20-22). The importance of PDGF BB in glomerular mesangial cell biology is established because mice deficient in either the B-chain or its receptor  $\beta$  lack these glomerular cells (23, 24). In fibrotic disorders such as diabetic nephropathy where  $TGF\beta$  contributes to the pathology, expression of PDGF BB and PDGFR $\beta$  is increased, especially in mesangial cells (25, 26). Whether PDGFR $\beta$  contributes to the signal transduction of TGF $\beta$  in mesangial cell pathology through a cross-talk has not been investigated. In our studies, we show that  $TGF\beta$ activates PDGFR $\beta$  to initiate Akt signaling to inactivate

PRAS40, which results in the increased mTORC1 activity necessary for mesangial cell hypertrophy and expression of matrix proteins fibronectin and collagen I ( $\alpha$ 2).

#### Results

## Activation of TGF $\beta$ RI stimulates PDGFR $\beta$ autophosphorylation in mesangial cells

Previously, we demonstrated that TGF $\beta$  increases the tyrosine phosphorylation of several proteins, including a protein at 190 kDa to induce noncanonical signaling in mesangial cells (15). To characterize the tyrosine kinase involved in the action of TGF $\beta$ , we considered PDGFR $\beta$  because it is a receptor tyrosine kinase, it is one of most abundant receptors expressed in mesangial cells, and it elicits pathology in glomerulosclerosis (5, 19, 27). Because activation of PDGFR $\beta$  requires its phosphorylation at Tyr-857 in the activation loop, we examined this in glomerular mesangial cells. TGF $\beta$  increased the phosphorylation of PDGFR $\beta$  in a time-dependent manner (Fig. 1*A*). Because TGF $\beta$  utilizes its type I serine/threonine kinase receptor for its signal transduction, we used a specific TGF $\beta$ RI inhibitor,



**Figure 2. TGF** $\beta$ **-stimulated PDGFR** $\beta$  **activates Akt signaling.** *A* and *D*, TGF $\beta$  increases phosphorylation of PDGFR $\beta$  at the PI 3-kinase binding site Tyr-751 to increase Akt phosphorylation. Mesangial cells were incubated with 2 ng/ml TGF $\beta$  for the indicated duration. The cleared cell lysates were immunoblotted with indicated antibodies. *B*, *C*, and *E*–*H*, mesangial cells were treated with 5  $\mu$ M SB (*panel B*) or 0.1  $\mu$ M JNJ (*panels C*, *E*, and *F*) or transfected with siRNAs against PDGFR $\beta$  or scrambled RNA (*panels G* and *H*) prior to incubation with 2 ng/ml TGF $\beta$  for 2 h. The cell lysates were immunoblotted against the indicated antibodies. *B* Nolecular weight markers are shown in the *left margins*.

SB-431542 (SB), to test its involvement in PDGFR $\beta$  phosphorylation. Similar to the effect on phosphorylation of Smad 2 and Smad 3, SB significantly inhibited TGF $\beta$ -stimulated PDGFR $\beta$ Tyr-857 phosphorylation (Fig. 1, *B*–*D*). PDGFR $\beta$  undergoes autophosphorylation at this site (13). Therefore, we used a specific PDGFR $\beta$  ATP competitive inhibitor, JNJ-10198409 (JNJ), in mesangial cells. JNJ blocked the autophosphorylation of PDGFR $\beta$ in response to TGF $\beta$  (Fig. 1*E*). Importantly, JNJ did not have any effect on phosphorylation of Smad 2 and Smad 3 in response to TGF $\beta$  (Fig. 1, *F* and *G*). Similarly, siRNAs against PDGFR $\beta$  to block its expression (Fig. 1*H*) showed no effect on TGF $\beta$ -induced phosphorylation of Smad 2 and Smad 3 (Fig. 1, *I* and *J*). These data indicate that TGF $\beta$  activates PDGFR $\beta$ ; however, PDGFR $\beta$ does not control the canonical Smad 2/3 phosphorylation by TGF $\beta$ RI.

#### TGF $\beta$ uses PDGFR $\beta$ for Akt kinase activation

We and others demonstrated previously that TGF $\beta$  regulates PI 3-kinase–dependent Akt activation (15, 28, 29). For activation of PI 3-kinase, the SH2 domain of the lipid kinase has to interact with the tyrosine phosphorylated proteins (30). In the

case of activated PDGFRB, PI 3-kinase associates with the phosphorylated Tyr-751 (31). Therefore, we determined the phosphorylation of PDGFR $\beta$  at this site in mesangial cells. TGF $\beta$  increased the phosphorylation of PDGFR $\beta$  at Tyr-751 in a time-dependent manner similar to its catalytic loop Tyr-857 phosphorylation (Fig. 2A). Both TGF $\beta$ RI and PDGFR $\beta$ inhibitors SB and JNJ significantly blocked the phosphorylation of PDGFRβ at Tyr-751 (Fig. 2, *B* and *C*). Activation of PI 3-kinase results in the phosphorylation of its downstream target Akt at the catalytic loop Thr-308 and hydrophobic motif Ser-473 sites (32). Therefore, to determine the activation of PI 3-kinase, we measured the phosphorylation of these Akt residues. TGF $\beta$  increased the phosphorylation of both these residues in a time-dependent manner (Fig. 2D). JNJ significantly inhibited the TGFB-induced phosphorylation of Akt at both of these sites (Fig. 2E). Because phosphorylation of Akt increases its kinase activity, we determined the phosphorylation of one of its endogenous substrates, glycogen synthase kinase  $3-\beta$  (GSK3 $\beta$ ). JNJ significantly inhibited GSK3 $\beta$  phosphorylation by TGF $\beta$  (Fig. 2*F*). To confirm this observation, we used siRNAs to down-regulate PDGFR $\beta$  in





**Figure 3. TGFβ-stimulated PDGFRβ regulates mTORC1signaling.** Mesangial cells were treated with 0.1 μM JNJ (*panels A, B, E, G,* and *I*) or transfected with siRNAs against PDGFRβ or scramble RNA (*C, D, F, H*, and *J*) prior to incubation with 2 ng/ml TGFβ. The cleared cell lysates were immunoblotted with indicated antibodies.

mesangial cells. siRNAs against PDGFR $\beta$  inhibited the TGF $\beta$ induced phosphorylation of Akt and its endogenous substrate GSK3 $\beta$  (Fig. 2, *G* and *H*).

#### PDGFRβ regulates TGFβ-induced mTORC1 activation

We and others demonstrated that TGF $\beta$  regulates mTORC1 activity (14, 16, 33). The exclusive mTORC1 subunit PRAS40 acts as an inhibitor of mTORC1 activity (34). Similarly, the GTPase-activating protein tuberin inhibits mTORC1 by acting on the Rheb GTPase (35). When phosphorylated by Akt, both of these proteins undergo inactivation (36–38). We determined the effect of PDGFR $\beta$  inhibition on the phosphorylation of

these proteins. JNJ significantly inhibited the TGF $\beta$ -stimulated phosphorylation of PRAS40 and tuberin (Fig. 3, *A* and *B*). Similarly, PDGFR $\beta$  siRNAs blocked phosphorylation of PRAS40 and tuberin in response to TGF $\beta$  (Fig. 3, *C* and *D*). Because phosphorylation/inactivation of PRAS40 and tuberin activate mTORC1, we determined activity of the latter by using phosphorylation of its substrates S6 kinase and 4EBP1. We also examined the phosphorylation of ribosomal protein S6 (rps6), which undergoes phosphorylation upon S6 kinase activation. TGF $\beta$  increased the phosphorylation of S6 kinase and 4EBP1 at mTORC1 sites and rps6 at S6 kinase sites. Both JNJ and siPDGFR $\beta$  markedly inhibited phosphorylation at these sites by TGF $\beta$  (Fig. 3, *E*–*J*).



**Figure 4.** PDGFR $\beta$  does not regulate TGF $\beta$ -induced ERK1/2 and STAT3 phosphorylation. *A* and *B*, mesangial cells were incubated with 2 ng/ml TGF $\beta$  for the indicated periods of time. *C*–*F*, mesangial cells were treated with 0.1  $\mu$ M JNJ (*panels C* and *D*) or transfected with siRNAs against PDGFR $\beta$  (*panels E* and *F*) prior to incubation with TGF $\beta$  (2 ng/ml) for 2 h. The cell lysates were immunoblotted with the indicated antibodies.

#### PDGFR $\beta$ -independent signaling by TGF $\beta$

Along with Akt/mTORC1, TGF $\beta$  also increases phosphorylation/activation of extracellular signal-regulated kinase (ERK) 1/2 and STAT3 (39, 40). TGF $\beta$  increased phosphorylation of ERK1/2 and STAT3 with similar kinetics (Fig. 4, *A* and *B*) as autophosphorylation of PDGFR $\beta$  (Fig. 1*A*). We examined whether activation of PDGFR $\beta$  is necessary for ERK1/2 and STAT3 phosphorylation. Interestingly, incubation of mesangial cells with JNJ did not have any effect on TGF $\beta$ -induced phosphorylation of ERK1/2 and STAT3 (Fig. 4, *C* and *D*). Similarly, siRNAs against PDGFR $\beta$  did not inhibit phosphorylation of ERK1/2 and STAT3 in response to TGF $\beta$  (Fig. 4, *E* and *F*). These results indicate that PDGFR $\beta$  is involved in TGF $\beta$ induced Akt/mTORC1 activation but not in the regulation of ERK1/2 and STAT3.

#### TGFβ regulates PDGF B expression

Our results above demonstrate that TGF<sup>β</sup> activates PDGFR<sup>β</sup>. To determine the mechanism of activation of PDGFR $\beta$ , we considered the expression of the PDGFR $\beta$  ligand PDGF B, which forms BB homodimer to activate the receptor (41-43). TGF $\beta$ time-dependently increased the expression of PDGF B in mesangial cells (Fig. 5A). These results indicate that a TGF $\beta$ -induced increase in PDGF expression may regulate PDGFR<sup>β</sup> activation. We tested this hypothesis. We used siRNAs against PDGF B. siPDGF B inhibited TGF\beta-stimulated phosphorylation of PDGFR $\beta$  (Fig. 5B). Also, TGF $\beta$ -induced phosphorylation of Akt was inhibited by siRNAs against PDGF B (Fig. 5C). These data conclusively demonstrate that PDGF B regulates the activation of PDGFR $\beta$  and the downstream activation of Akt/mTORC1 by TGF<sup>β</sup>. We tested this notion by using PDGF BB ligand directly in mesangial cells. PDGF BB increased the phosphorylation of PDGFR $\beta$  in a time-de-

pendent manner (Fig. 6A) similar to the kinetics of PDGF B expression by TGF $\beta$  (Fig. 5A). Also, PDGF BB enhanced the phosphorylation of Akt (Fig. 6B), resulting in phosphorylation of PRAS40 and tuberin (Fig. 6, C and D), which led to the activation of mTORC1 as judged by phosphorylation of S6 kinase, rps6, and 4EBP1 (Fig. 6, E-G). Furthermore, increased phosphorylation of PDGFRB was observed when PDGF BB ligand was added as compared with TGF $\beta$  alone (Fig. S1A). No further increase in phosphorylation was found when both ligands were added (Fig. S1A). Importantly, this increased PDGFR<sup>β</sup> phosphorylation resulted in a similar increase in Akt phosphorylation leading to mTORC1 activation (Fig. S1, B-G). Similar to the observations with TGFβ, PDGFRβ inhibitor JNJ blocked PDGF BB-stimulated phosphorylation of Akt, PRAS40, and tuberin (Fig. S2, A-C). JNJ also blocked activation of mTORC1 in response to PDGF BB (Fig. S2, D-F).

#### PDGF regulates TGFβ signaling

The above results show that TGF $\beta$  utilizes PDGF ligand to activate PDGFR $\beta$ . We examined whether there is a crosstalk between PDGF and TGF $\beta$ . Incubation of mesangial cells with PDGF increased the expression of TGF $\beta$  in a time-dependent manner (Fig. 7*A*). Inhibition of PDGFR $\beta$  with JNJ blocked PDGF-stimulated TGF $\beta$  expression (Fig. 7*B*). To test if PDGF signals through TGF $\beta$ , we examined phosphorylation of Smads. PDGF increased phosphorylation of Smad 2 and Smad 3, and the TGF $\beta$  receptor kinase inhibitor SB blocked these phosphorylations (Fig. 7, *C* and *D*). Interestingly, TGF $\beta$  antibody inhibited PDGF-stimulated Smad 2 and Smad 3 phosphorylation (Fig. 7, *E* and *F*). These data demonstrate the presence of a positive feedback loop between TGF $\beta$  and PDGF.





**Figure 5. TGF** $\beta$  **increases PDGF B expression to augment activating phosphorylation of PDGFR** $\beta$  **and Akt.** *A*, mesangial cells were incubated with TGF $\beta$  (2 ng/ml) for the indicated period of time. *B* and *C*, mesangial cells were transfected with siPDGF B or scrambled RNA prior to incubation with TGF $\beta$  (2 ng/ml) for 2 h. The cell lysates were immunoblotted with the indicated antibodies.



Figure 6. PDGF increases activating phosphorylation of PDGFRβ leading to Akt/mTORC1 activity. Mesangial cells were incubated with 20 ng/ml PDGF for indicated period of time. The cell lysates were immunoblotted with the indicated antibodies.

# PRAS40 regulates TGF $\beta$ -induced mTORC1 activity downstream of PDGFR $\beta$

Our results above show that TGF $\beta$ -induced PDGFR $\beta$  activation contributes to phosphorylation of Akt and PRAS40, resulting in mTORC1 activation (Fig. 2 and Fig. 3). To determine whether Akt-mediated phosphorylation of PRAS40 regulates the activation of mTORC1 downstream of PDGFR $\beta$ , we transfected mesangial cells with siRNAs against PDGFR $\beta$ , along with constitutively active myristoylated Akt (myr-Akt) and the non-phosphorylatable mutant PRAS40 T246A. Expression of myr-Akt alone increased the phosphorylation of S6 kinase, rps6, and 4EBP1, similar to TGF $\beta$  treatment (data not shown). However, constitutively active Akt kinase reversed the inhibition of phosphorylation of S6 kinase, rps6, and 4EBP1 by siPDGFR $\beta$  in the presence of TGF $\beta$  (Fig. 8, *A*–*C*; compare 4th lanes with 3rd lanes). Importantly, the phospho-deficient



**Figure 7. PDGF increases expression of TGF** $\beta$  **to induce its signaling.** *A*, mesangial cells were incubated with PDGF (20 ng/ml) for the indicated period of time. *B*–*F*, mesangial cells were treated with 0.1 µM JNJ (*panel B*) or 5 µM SB (*panels C* and *D*) or 1 µg/ml IgG or TGF $\beta$  antibody (*panels E* and *F*) prior to incubation with PDGF for 2 h. The cell lysates were immunoblotted with the indicated antibodies.

mutant of PRAS40 abrogated the reversal of phosphorylation of these mTORC1 and S6 kinase substrates by constitutively active Akt (Fig. 8, A-C; compare *5th lanes* with *4th lanes*). These results conclusively demonstrate that Akt-mediated phosphorylation-dependent inactivation of PRAS40 downstream of PDGFR $\beta$  regulates TGF $\beta$ -induced mTORC1 activation in mesangial cells.

# Activation of Akt kinase by PDGFR $\beta$ regulates TGF $\beta$ -induced mesangial cell hypertrophy and matrix protein expression

We demonstrated that TGF\beta-stimulated Akt kinase regulates mesangial cell hypertrophy and expression of matrix proteins (14, 44). Because PDGFR $\beta$  contributes to the activation of Akt by TGF $\beta$ , we investigated the involvement of this receptor tyrosine kinase in mesangial cell hypertrophy. JNJ significantly inhibited TGFβ-induced protein synthesis and hypertrophy of mesangial cells (Fig. 9, A and B). Similarly, siRNAs against PDGFR $\beta$ attenuated both these phenomena (Fig. 9, C and D). Because PRAS40 phosphorylation by Akt regulates its downstream signaling, we determined the involvement of Akt. Expression of constitutively active myr-Akt restored the siPDGFR<sub>β</sub>-mediated inhibition of mesangial cell protein synthesis and hypertrophy in the presence of TGF $\beta$  (Fig. 9, *E* and *F*). Furthermore, expression of phospho-deficient PRAS40 T246A significantly prevented the myr-Akt-mediated reversal of siPDGFR<sub>B</sub>-induced inhibition of protein synthesis and hypertrophy (Fig. 9, *E* and *F*).

TGF $\beta$  regulates renal fibrosis, including glomerulosclerosis, by increasing the synthesis and deposition of matrix proteins fibronectin and collagen (3). TGF $\beta$  increased the expression of fibronectin and collagen I ( $\alpha$ 2) in mesangial cells (44–46). Both JNJ and siRNA against PDGFR $\beta$  inhibited TGF $\beta$ -stimulated expression of these matrix proteins (Fig. 10, *A*–*D*). We showed above that TGF $\beta$ -induced expression of PDGF B regulates activation of PDGFR $\beta$  and Akt kinase (Fig. 5). Therefore, we tested the effect of siPDGF B on expression of fibronectin and colla-

gen I ( $\alpha$ 2). Inhibition of PDGF B blocked fibronectin and collagen I ( $\alpha$ 2) expression in response to TGF $\beta$  (Fig. 10, *E* and *F*). To further evaluate the mechanism of matrix expression, we determined the role of Akt because it is activated downstream of TGF $\beta$  by PDGFR $\beta$  activation (Fig. 2D, Fig. 6B, and Fig. S1B). We used an Akt inhibitor, MK, which blocked Akt phosphorylation by TGF $\beta$  (Fig. S3A). MK inhibited TGF $\beta$ -induced expression of fibronectin and collagen I ( $\alpha$ 2) (Fig. S3, *B* and *C*). We showed above that siPDGFR<sup>β</sup> blocked TGF<sup>β</sup>-induced expression of fibronectin and collagen I ( $\alpha$ 2) (Fig. 10, C and D). Expression of myr-Akt reversed the siPDGFR<sub>B</sub>-mediated inhibition of TGFβ-stimulated expression of both of these matrix proteins (Fig. 10, G and H; compare 4th lanes with 3rd lanes). Importantly, PRAS40 T246A phospho-deficient mutant inhibited this myr-Akt effect on fibronectin and collagen I ( $\alpha$ 2) expression (Fig. 10, G and H; compare 5th lanes with 4th lanes). Together, these data show a significant role of Akt downstream of PDGFRβ in mesangial cell matrix protein expression. Furthermore, our results demonstrate a critical role for the Akt and its substrate PRAS40 in PDGFR<sub>β</sub>-mediated signaling by TGF $\beta$  in mesangial cell pathology.

# PDGFRβ-stimulated mTORC1 regulates TGFβ-induced mesangial cell hypertrophy and matrix protein expansion

We demonstrated previously that activation of mTORC1 by TGF $\beta$  contributes to mesangial cell hypertrophy and matrix protein expression (14, 46, 47). Our results above show a role of PDGFR $\beta$  in TGF $\beta$ -stimulated mTORC1 activation in mesangial cells (Fig. 3). Therefore, we examined whether mTORC1 downstream of PDGFR $\beta$  regulates mesangial cell hypertrophy. To test this, we used a vector expressing raptor, which is a constitutive component of mTORC1 and is essential for its kinase activity (48, 49). Expression of raptor in mesangial cells increased mTORC1 activity as judged by phosphorylation of S6







**Figure 8. PDGFRβ regulates TGFβ-induced Akt/PRAS40-mediated mTORC1 activation.** *A*, mesangial cells were transfected with siRNAs against PDGFRβ or scrambled RNA or HA-tagged myr-Akt and PRAS40 (Thr-246A) mutant expression vectors as indicated. Transfected cells were incubated with 2 ng/ml TGFβ. The cleared cell lysates were immunoblotted with indicated antibodies.

kinase, rps6, and 4EBP1 (Fig. S4, A-C). Raptor reversed the siPDGFR<sub>B</sub>-mediated inhibition of protein synthesis and hypertrophy of mesangial cells in the presence of TGF $\beta$  (Fig. 11, A and B). Similarly, raptor restored the inhibition of fibronectin and collagen I ( $\alpha$ 2) by siPDGFR $\beta$  in the TGF $\beta$ -stimulated cells (Fig. 11, C and D). To confirm these observations, we used a mutant of mTOR, which delivers hyperactive mTORC1 (49, 50). Expression of this mutant increased the phosphorylation of S6 kinase, rps6, and 4EBP1 in the absence of any effect on phosphorylation of Akt at Ser-473 in mesangial cells (Fig. S5, A-D), indicating that it acts as hyperactive mTORC1 and not mTORC2. Expression of this hyperactive mTORC1 reversed the siPDGFRβ-mediated inhibition of protein synthesis, hypertrophy, and matrix proteins fibronectin and collagen I ( $\alpha$ 2) expression (Fig. S6, A–D). These results show a positive role of mTORC1 downstream of PDGFRB in the TGF $\beta$ -induced mesangial cell pathology.

# Phosphorylation of PDGFR $\beta$ in the kidneys of type 1 diabetic OVE26 mice and streptozotocin-induced rats

Diabetic nephropathy is a fibrotic disorder. In this disease, mesangial expansion proceeds to glomerular hypertrophy and accumulation of matrix proteins, which lead to glomerulosclerosis (51). Hyperglycemia produces multiple fibrotic growth factors, including TGF $\beta$ , which contribute to the pathogenesis of diabetic nephropathy (4, 51). Our results described above show a role of PDGFRB in TGFB-induced mesangial cell hypertrophy and matrix protein expression. To investigate the in vivo relevance of our results, we used OVE26 type 1 diabetic mice. We demonstrated recently that the blood glucose levels in 3-month-old OVE26 mice are significantly increased and are associated with renal hypertrophy (52). Renal cortical lysates were used to examine the expression of TGF $\beta$ . In diabetic renal cortical samples, a significant increase in  $TGF\beta$  was detected (Fig. 12, A and B). This increased TGF $\beta$  correlated with an increase in phosphorylation of Smad 2 and Smad 3, two downstream TGFβ receptor-specific Smads (Fig. 12, C-E). Furthermore, diabetic mice showed a significantly increased expression of PDGF B (Fig. 12, F and G). The level of phosphorylation of PDGFR $\beta$  at Tyr-857, which is required for its tyrosine kinase activity, was significantly enhanced (Fig. 12, H and I). Also, the level of phosphorylation at Tyr-751 of PDGFR<sup>β</sup> was elevated in the diabetic renal cortex (Fig. 12, J and K). Our results above show that PDGFRβ is necessary for TGFβ-induced mTORC1 activation (Fig. 3). We determined the phosphorylation of S6 kinase, rps6, and 4EBP1 as readout of mTORC1 activation in renal cortical lysates. A significant increase in phosphorylation of S6 kinase, rps6, and 4EBP1 was observed in the diabetic mice, indicating activation of mTORC1 (Fig. 12, L-R). To



**Figure 9. PDGFR** $\beta$  **regulates TGF** $\beta$ -**induced protein synthesis and hypertrophy of mesangial cells.** Mesangial cells were treated with 0.1  $\mu$ M JNJ (*panels A* and *B*) or transfected with siRNAs against PDGFR $\beta$  or scrambled RNA (*C* and *D*) along with myr-Akt and PRAS40 (Thr-246A) as indicated (*E* and *F*) prior to incubation with 2 ng/ml TGF $\beta$  for 24 h. Protein synthesis and hypertrophy were measured as described in "Experimental procedures." Mean  $\pm$  S.D. of triplicate (*A*-*D*) or quadruplicate (*E* and *F*) measurements is shown. In *panels A*-*D*, \**p* < 0.0001–0.002 *versus* control; \*\**p* < 0.001–0.006 *versus* TGF $\beta$ . In *panels E* and *F*, \**p* < 0.0001 *versus* control; \*\**p* < 0.01 *versus* TGF $\beta$ ; #*p* < 0.01 *versus* TGF $\beta$  + siPDGFR $\beta$ ; @*p* < 0.01 *versus* TGF $\beta$  + myr-Akt.

confirm these observations in mice, we used a streptozotocin (STZ)-induced rat model of type 1 diabetes, which showed early pathologic changes of diabetic nephropathy (44). From kidney cortex, we prepared glomeruli, which contains the mesangial cells. Glomerular lysates were used to determine the expression of TGF $\beta$ . In the diabetic glomeruli, significantly increased TGF $\beta$  was detected, which correlated with enhanced Smad 2 and Smad 3 phosphorylation (Fig. S7, A-E). Furthermore, increased expression of PDGF B and phosphorylation of PDGFR $\beta$  were observed in the diabetic glomeruli (Fig. S7, F-K). Additionally, diabetic rat glomeruli showed activation of mTORC1 as judged by phosphorylation of S6 kinase, rps6, and 4EBP1 (Fig. S7, L-R). These results show that activation of PDGFR $\beta$  is associated with increased expression of TGF $\beta$  in the diabetic kidney.

#### Discussion

We report activation of PDGFR $\beta$  by TGF $\beta$  via expression of PDGF, which also contributes to TGF $\beta$  signaling in glomerular

mesangial cells (Fig. 13). We show that TGF $\beta$ -induced Akt kinase activation requires PDGFR $\beta$  tyrosine kinase activity. Furthermore, we provide novel evidence that Akt-mediated phosphorylation of PRAS40 by TGF $\beta$  is mediated by PDGFR $\beta$  and is required for mesangial cell hypertrophy and matrix protein expression. Finally, we show activation of PDGFR $\beta$  and mTORC1 in kidneys of diabetic mice that are undergoing matrix expansion.

A role of TGF $\beta$  in human chronic kidney disease was established decades ago (53, 54). Also, in the animal models of renal fibrosis such as diabetic nephropathy, increased expression of TGF $\beta$  promotes the pathology (4, 17, 55). Similarly, in a rat model of glomerulonephritis, increased expression of TGF $\beta$  promotes fibrosis (56). Administration of anti-TGF $\beta$ antibody ameliorates the disease progression (57). Similarly, a murine monoclonal TGF $\beta$  antibody reduced matrix expansion and disease-associated histological changes in adriamycin- and podocyte ablation–induced nephropathies in mice





**Figure 10. PDGFR** $\beta$  **regulates TGF** $\beta$ -**induced fibronectin and collagen I** ( $\alpha$ **2**). Mesangial cells treated with JNJ (*panels A* and *B*) or transfected with siRNAs against PDGFR $\beta$  or scrambled RNA (*panels C* and *D*) or siRNAs against PDGF B (*panels E* and *F*) or myr-Akt and PRAS40 (*panels G* and *H*) and incubated with TGF $\beta$  for 24 h. The cleared cell lysates were immunoblotted with the indicated antibodies.

(58). Furthermore, in diabetic nephropathy associated with STZ-induced type 1 diabetes in mice and with type 2 diabetic db/db mice, neutralization of TGF $\beta$  with anti-TGF $\beta$  antibody ameliorates the glomerular hypertrophy and matrix protein expansion (11, 12). A hypomorphic mouse with  $TGF\beta$ expression at 10% the level of that in WT showed significant reduction in glomerular filtration rate and albuminuria. In fact, 3-fold overexpression of TGFB in mice causes glomerulosclerosis and albuminuria (59). More recently, administration of a soluble TGF $\beta$  receptor II by a gene therapy protocol to mice with renal fibrosis, where  $TGF\beta$  contributes to the pathology, showed significant beneficial effects (60). These results conclusively demonstrate a significant pathologic role of TGF $\beta$  in renal diseases, suggesting that blocking of TGF $\beta$ may be beneficial in renal fibrosis. However, the lack of progress is due to the concerns that inhibition of TGF $\beta$  may enhance the risk of autoimmune disease (17). Because  $TGF\beta$ has multiple roles in maintaining various homeostatic processes, direct targeting of this cytokine may elicit multi-organ side effects. Thus, an alternative therapeutic strategy to inhibit pathologic actions of TGF $\beta$  in glomerulos clerosis needs to be identified.

The role of receptor tyrosine kinases in renal disease is established (61). Both positive and negative regulatory roles of epidermal growth factor receptor (EGFR) in renal pathology have been reported. For example, in ischemic kidney injury, activation of EGFR ameliorates the disease progression (62). Furthermore, in an EGFR mutant mouse with significantly reduced tyrosine kinase activity, tubular damage was more severe with increased apoptosis after nephrotoxic injury than in the WT mice, demonstrating a protective role of EGFR (63). On the other hand, a pathologic role of this receptor in TGF\beta-mediated renal fibrosis was reported (17). EGFR is abundantly expressed in the renal proximal tubular epithelial cells and interstitial fibroblasts (62). Inhibition of EGFR and proximal tubule-specific deletion of this receptor tyrosine kinase ameliorate TGF<sub>β</sub>-mediated renal fibrosis (64, 65). Also, involvement of Src tyrosine kinase has been reported in angiotensin IIinduced kidney fibrosis (66). Thus, a role of tyrosine kinases in renal fibroblasts and tubular epithelial cells is established for





**Figure 11. PDGFR** $\beta$ -stimulated mTORC1 regulates TGF $\beta$ -induced protein synthesis, hypertrophy, and matrix proteins fibronectin and collagen I ( $\alpha$ 2) expression. Mesangial cells were transfected with siRNAs against PDGFR $\beta$  or scrambled RNA along with Myc-tagged raptor expression vector as indicated. Transfected cells were incubated with 2 ng/ml TGF $\beta$ . In *panels A* and *B*, respectively, protein synthesis and hypertrophy of mesangial cells were measured as described in the experimental procedures. Mean  $\pm$  S.D. of triplicate measurements is shown. \*p < 0.008 (*panel A*) and \*p < 0.001 (*panel B*) versus control; \*\*p < 0.013 (*panel A*) and \*\*p < 0.002 (*panel B*) versus TGF $\beta$ ; #p < 0.005 (*panels* A) and #p < 0.004 (*panel B*) versus TGF $\beta$  + siPDGFR $\beta$ . In *panels C* and *D*, the cleared cell lysates were immunoblotted with the indicated antibodies.

induction of fibrosis. In the present study, we found that TGF $\beta$  activates the PDGFR $\beta$  in the renal mesangial cells. These results are in contrast to the observations of a recent report in which TGF $\beta$  did not have any effect on PDGFR $\beta$  phosphorylation in fibroblasts, suggesting cell-specificity for this interaction (67). In fact, we showed that TGF $\beta$  increased the expression of PDGF, which binds to the PDGFR $\beta$  to induce autophosphorylation and its activation. Addition of PDGF directly to mesangial cells also increased phosphorylation/activation of PDGFR $\beta$ .

Our data in renal mesangial cells suggest that PDGFR $\beta$  may serve as the initiator of TGF $\beta$  noncanonical signaling. We and others previously reported that both Smad 3 and PI 3-kinase/ Akt signaling regulate mesangial cell pathology in response to TGF $\beta$  (15, 29, 45, 68). PI 3-kinase activation requires its association with tyrosine phosphorylated proteins (69). In the case of PDGFR $\beta$ , the specific residue was identified as Tyr-751 (31). We showed that TGF $\beta$  increased the tyrosine phosphorylation of this residue, which was sensitive to the inhibition of PDGFRβ tyrosine kinase activity, demonstrating autophosphorylation by PDGFR $\beta$ . These results indicate that TGF $\beta$  may activate PI 3-kinase via this direct interaction mechanism with PDGFRβ. In fact, we found that TGFβ-stimulated phosphorylation of Akt, a downstream target of active PI 3-kinase, depended upon PDGFRB and its tyrosine kinase activity. This observation was similar to that obtained with direct addition of PDGF to mesangial cells. Interestingly, we found that PDGF

increased the expression of TGF $\beta$  to induce Smad 2/3 phosphorylation. These results identify a positive feedback loop between TGF $\beta$  and PDGF signaling in mesangial cells (Fig. 13)

Recently, we and others demonstrated that mTORC1 controls renal cell pathology in rodent models of fibrosis where TGF $\beta$  plays important role (70–76). Three protein subunits of mTORC1, raptor, deptor, and PRAS40, regulate the activity of this kinase complex (48). Both nutrients and growth factors use independent mechanisms for mTORC1 activation. Amino acids promote formation of GTP-bound Rag proteins via Ragulator to recruit mTORC1 to GTP-bound Rheb for its activation (77). During growth factor receptor tyrosine kinase stimulation of cells, activated Akt kinase phosphorylates the raptor binding protein, PRAS40, which under basal state inhibits the substrate recruitment to mTORC1. Akt-mediated phosphorylation of PRAS40 results in its inactivation to increase the mTORC1 activity (34). For example, inactivation of PRAS40 in HeLa cells increased mTORC1 activity, leading to inhibition of their apoptosis. Interestingly, rapamycin did not reverse this anti-apoptotic effect, indicating that the effect of PRAS40 does not involve inhibition of mTORC1 (78). In 293 cells, PRAS40 inactivation showed inhibition of mTORC1 activity (79). These results are similar to those observed in Drosophila eyes where the hypomorphic allele of *lobe*, the fly ortholog of PRAS40, resulted in inhibition of mTORC1 and reduced eye size (80, 81). We reported previously that TGF $\beta$  stimulates phosphorylation





Figure 12. Increased level of TGF $\beta$  is associated with activation of PDGFR $\beta$  and mTORC1 signaling in OVE26 mice renal cortexes. Renal cortical lysates from control and OVE26 diabetic mice were immunoblotted with indicated antibodies. *Bottom panels* in each blot represent quantification of protein bands. Mean ± S.D. of 3–4 animals is shown. \*p < 0.001-0.03 versus control animals.

of PRAS40 in mesangial cells (46). Now we show that PDGFR $\beta$  tyrosine kinase is required for this phosphorylation of PRAS40 by TGF $\beta$ , similar to PDGF addition. In contrast to the results in 293 cells and in *Drosophila* described above, phosphorylation of PRAS40 by PDGFR $\beta$ -mediated activated Akt kinase enhances mTORC1 activity in mesangial cells by both TGF $\beta$  and PDGF.

Induction of mesangial cell hypertrophy and matrix protein accumulation correlate with increased expression of TGF $\beta$  in rodent models of fibrosis (17, 51, 82). Also, renal cells derived from TGF $\beta$  null mice show impaired hypertrophic response and fibronectin expression (83). Similarly, glomerular hypertrophy and mesangial matrix expansion are reduced in TGF $\beta$ receptor II heterozygous mice models of diabetes (84). We demonstrated previously that TGF $\beta$  regulates mesangial cell hypertrophy and fibronectin expression (14, 46, 47). A role of Akt kinase was initially identified in cell-size control in the fruit fly. Overexpression of Akt in the imaginal disc of *Drosophila* 

increases the cell size (85). Similarly, transgenic mice expressing constitutively active Akt in the heart show cardiomyocyte hypertrophy (86). Also, Akt kinase activity is required for TGF $\beta$ -induced mesangial cell hypertrophy (44). We showed that expression of both fibronectin and collagen I ( $\alpha$ 2) by TGF $\beta$  was meditated by Akt kinase. However, the specific substrate that regulates this TGF $\beta$  response is not clear. In mouse hearts, Akt-mediated phosphorylation of PRAS40 increases mTORC1 activity to induce pathological hypertrophy and fibrosis (87). We reported the requirement of mTORC1 in TGF $\beta$ -induced mesangial cell hypertrophy and matrix protein expression (14, 46, 47). Now we demonstrate that Akt-phosphorylated PRAS40 is required for mesangial cell hypertrophy and fibronectin and collagen I ( $\alpha 2$ ) expression in response to TGF $\beta$ . In fact, we provide evidence that PDGFR $\beta$  tyrosine kinase downstream of TGF $\beta$ RI acts as a mediator of these phenomena. To our knowledge, this is the first demonstration of



Figure 13. Schematic summarizing our results. Thin arrows indicate canonical TGF $\beta$  signaling. Bold arrows show noncanonical signal transduction described in the present study.

the involvement of PDGFR $\beta$  tyrosine kinase in TGF $\beta$ -stimulated Akt/PRAS40/mTORC1 signaling that contributes to mesangial cell hypertrophy and matrix protein expansion. Furthermore, our results demonstrate an *in vivo* relevance showing association of increased TGF $\beta$  expression with activation of PDGFR $\beta$  and mTORC1 in the renal cortex and glomeruli of diabetic mice and rats, respectively. Thus, in TGF $\beta$ -regulated fibrotic renal diseases such as diabetic nephropathy, attacking PDGFR $\beta$  may provide an attractive therapeutic option that bypasses the detrimental effects of directly targeting TGF $\beta$ .

#### **Experimental procedures**

#### Reagents

TGF $\beta$ 1 and PDGF BB were obtained from R&D Systems. Cell culture materials, including Opti-MEM medium, were purchased from Thermo Fisher Scientific. PMSF, Nonidet P-40, Na<sub>3</sub>VO<sub>4</sub>, protease inhibitor mixture, JNJ,  $\beta$ -actin (catalog no. A2066, lot 3082M4781) and fibronectin (catalog no. F3648, lot 103M4818V) antibodies were purchased from Sigma. The antibodies against phospho-PDGFR $\beta$  (Tyr-857) (catalog no.

3270S, lot 3), phospho-PDGFRβ (Tyr-751; catalog no. 3161S, lot 7), phospho-Smad 3 (Ser-423/425; catalog no. 9520S, lot 15), Smad 3 (catalog no. 9513S, lot 2), phospho-Smad 2 (Ser-465/467; catalog no. 3108S, lot 2), Smad 2 (catalog no. 5339S, lot 6), phospho-Akt (Ser-473; catalog no. 9271S, lot 14), phospho-Akt (Thr-308; catalog no. 9275S, lot 26), Akt, (catalog no. 9272S, lot 28) phospho-GSK3β (Ser-9; catalog no. 9323S, lot 9), GSK3B (catalog no. 12456S, lot 3), phospho-PRAS40 (Thr-246; catalog no. 2997S, lot 12), PRAS40 (catalog no. 2691S, lot 8), phospho-tuberin (catalog no. 3611S, lot 6), phospho-4EBP1 (Thr-37/46) (catalog no. 9459S, lot 10), phospho-4EBP1 (Ser-65; catalog no. 9456S, lot 5), 4EBP1 (catalog no. 9452S, lot 10), phospho-S6 kinase (Thr-389; catalog no. 9205S, lot 16), S6 kinase (catalog no. 9202S, lot 20), phospho-rps6 (Ser-240/244; catalog no. 5364S, lot 5), rps6 (catalog no. 2217S, lot 7), phospho-STAT3 (Tyr-705; catalog no. 9131S, lot 18), mTOR (catalog no. 3270S, lot 3), phospho-ERK1/2 (Thr-202/Tyr-204; catalog no. 9101S, lot 28), and ERK1/2 (catalog no. 9102S, lot 19) were obtained from Cell Signaling Technology. PDGFR $\beta$  (catalog no. 2972S, lot 6), tuberin (SC-893, lot K1703), STAT3 (catalog no. SC-482, lot E176), and Myc (catalog no. SC-40, lot A1201) antibodies and pooled siRNAs against PDGFR $\beta$  were purchased from Santa Cruz Biotechnology. Collagen I ( $\alpha$ 2) antibody (catalog no. 14695-1-AP) was obtained from Proteintech. PDGF B antibody was purchased from Millipore (catalog no. 06-127, lot 2585791). HA antibody (catalog no. MMS-101R-500, lot D13FF01646) was obtained from Babco (Princeton, NJ). TGFβ antibody (catalog no. ab27969, lot GR66929-7) was purchased from Abcam. The PVDF membrane to transfer proteins was obtained from PerkinElmer. The transfection reagent FuGENE HD was purchased from Promega. HA-tagged mvr-Akt, PRAS40 (Thr-246A), and Mvc-tagged raptor expression plasmids have been described previously (49, 88, 89). The mTOR mutant expression vector (SL1 + I2017T), which renders hyperactive mTORC1 activity, has been described (49, 50).

#### Cell culture

Human mesangial cells were originally prepared by Abboud and co-workers (90) from glomeruli isolated by differential sieving. The resuspended glomeruli were digested with collagenase. Mesangial cells were cultured from outgrowths of collagenase-treated glomeruli and characterized (90). The frozen cells were thawed and grown in DMEM in the presence of 10% FBS as described previously (47, 91–93). The cells were used between passages 9 and 12. Before performing experiments, the cells were grown in complete medium until they reached confluency. The cell monolayer was then starved in serum-free medium for 24 h prior to incubation with TGF $\beta$  (2 ng/ml) for indicated periods of time.

#### Animals

FVB mice overexpressing calmodulin transgene in the  $\beta$ -cell of pancreas, called OVE26 mice, were used and purchased from The Jackson Laboratory. At 3 days of age the OVE26 mice are hyperglycemic (94). At 2 months of age, these mice develop pathologies of diabetic nephropathy, including significant renal hypertrophy, glomerular hypertrophy, and albuminuria (52,



95). The control FVB and diabetic OVE26 mice had free access to food and water. At 3 months of age, the animals were sacrificed, the kidneys were removed, and renal cortical tissues were isolated and frozen as described previously (52).

To induce diabetes in rats (Sprague Dawley; 200–250 g), 55 mg/kg of body weight STZ in sodium citrate buffer (pH 4.5) was injected by tail vein. The blood glucose was measured after 24 h. The animals had free access to water and food. At 5 days after STZ injection, the rats were euthanized and the kidneys were removed to isolate cortical sections. Glomeruli were prepared from the kidney cortexes by a differential sieving technique as described (90). Both mice and rats were kept at the University of Texas Health Science Center animal facility. The University of Texas Health Science Center Institutional Animal Care and Use Committee approved both protocols.

#### Preparation of cell, cortical, and glomerular lysates

At the end of the incubation period, the mesangial cell monolayer was washed twice with PBS. The monolayer was then harvested in radioimmune precipitation assay buffer (RIPA; 20 mM Tris-HCl, pH 7.5, 150 mM NaCl, 5 mM EDTA, 1 mM Na<sub>3</sub>VO<sub>4</sub>, 1% Nonidet P-40, 1 mM PMSF, and 0.1% protease inhibitor mixture). The renal cortexes from control and diabetic mice and the glomeruli from the control and STZ-induced diabetic rats were harvested in the same RIPA buffer. The cells, cortexes, and glomeruli were lysed in RIPA at 4 °C for 30 min. After the incubation, the cell, cortical, and glomerular extracts were centrifuged at 12,000 × g for 30 min at 4 °C. The cleared supernatant was collected as cell, cortical, and glomerular lysates. Protein concentration was determined.

#### Immunoblotting

Cell lysates containing equal amounts of protein were separated by SDS-PAGE. The proteins were transferred to PVDF membrane. For immunoblotting, the membrane containing the proteins was incubated with indicated primary antibody at 4 °C. The primary antibody dilutions used for immunoblotting were 1:1000. The membrane was washed, followed by incubation with horseradish peroxidase-conjugated secondary antibody (1:10,000). The membrane was then incubated with enhanced chemiluminescence reagent to develop a specific protein band, which was visualized by exposing the membrane to X-ray film (91, 96).

#### Transfection

The semiconfluent mesangial cell monolayer was washed with PBS. Opti-MEM was added. The expression vector or siR-NAs against PDGFR $\beta$  were transfected in Opti-MEM with FuGENE HD as described previously (91). After 24 h, the cells were starved as described above prior to adding TGF $\beta$ .

#### Measurement of protein synthesis

Protein synthesis was determined as described previously (14, 89, 91). Briefly, during the last four hours of incubation with TGF $\beta$ , the cells were incubated with 1 µCi of [<sup>35</sup>S]methionine as described before (44). At the end of incubation, the cells

were lysed in RIPA as described above and the protein was estimated. Equal amount of protein was spotted onto 3MM filter paper and washed in boiling 10% trichloroacetic acid containing 0.1 gm/l methionine for 1 min. The filters were then dried and the radioactivity was counted using scintillation fluid.

#### Measurement of hypertrophy

The cells were trypsinized in the medium at the end of the incubation and counted using a hemocytometer (91, 96). The cells were then gently centrifuged at  $1,000 \times g$  for 5 min at 4 °C. The cell pellet was washed with PBS and resuspended in RIPA buffer to lyse, as described above. Protein concentration was measured in the cell lysate and the ratio of total protein to cell number was determined. The increase in the ratio was considered as cell hypertrophy (44, 89).

#### Statistics

Prism GraphPad software was used to determine the significance of the data. Analysis of variance followed by Tukey's multiple comparison test was used. A *p*-value of <0.05 was considered significant (97).

#### Data availability

All data are contained within the article and in the supporting information.

Author contributions—S. M. and F. D. data curation; S. M., F. D., and G. G.-C. formal analysis; G. G.-C. conceptualization; G. G.-C. supervision; G. G.-C. funding acquisition; G. G.-C. writing-original draft; G. G.-C. project administration; G. G.-C. writing-review and editing; B. S. K. analyzed *in vivo* data, took part in the discussion of the data, and edited the manuscript; N. G.-C. took part in discussion of the results and provided intellectual input.

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*Abbreviations*—The abbreviations used are: TGFβ, transforming growth factor- $\beta$ ; PDGF, platelet-derived growth factor; PDGFR, PDGF receptor; PI, phosphatidylinositol; PRAS40, proline-rich Akt substrate of 40 kDa; mTORC, mTOR complex; Akt, protein kinase B; TGF $\beta$ R, TGF $\beta$  receptor; SB, SB-431542; JNJ, JNJ-10198409; SH, src homology; GSK, glycogen synthase kinase; 4EBP, eukaryotic translation initiation factor 4E-binding protein; rps6, ribosomal protein S6; ERK, extracellular signal-regulated kinase; STZ, streptozotocin; EGFR, epidermal growth factor receptor; RIPA, radioimmune precipitation assay buffer.

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