# Rapamycin inhibits transforming growth factor β-induced peritoneal angiogenesis by blocking the secondary hypoxic response

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#### Abstract

Patients with end-stage kidney disease on peritoneal dialysis often develop progressive scarring of the peritoneal tissues. This manifests as submesothelial thickening and is associated with increased vascularization that leads to ultrafiltration dysfunction. Hypoxia induces a characteristic series of responses including angiogenesis and fibrosis. We investigated the role of hypoxia in peritoneal membrane damage. An adenovirus expressing transforming growth factor (TGF)  $\beta$  was used to induce peritoneal fibrosis. We evaluated the effect of the mTOR inhibitor rapamycin, which has been previously shown to block hypoxia-inducible factor (HIF) 1 $\alpha$ . We also assessed the effect of HIF1 $\alpha$  independently using an adenovirus expressing active HIF1 $\alpha$ . To identify the TGF $\beta$ 1-independent effects of HIF1 $\alpha$ , we expressed HIF1 $\alpha$  in the peritoneum of mice lacking the TGF $\beta$  signalling molecule Smad3. We demonstrate that TGF $\beta$ -induced fibroproliferative tissue is hypoxic. Rapamycin did not affect the early angiogenic response, but inhibited angiogenesis and submesothelial thickening 21 days after induction of fibrosis. In primary mesothelial cell culture, rapamycin had no effect on TGF $\beta$ -induced vascular endothelial growth factor (VEGF) but did suppress hypoxia-induced VEGF. HIF1 $\alpha$  induced submesothelial thickening and angiogenesis in peritoneal tissue. The fibrogenic effects of HIF1 $\alpha$  were Smad3 dependent. In summary, submesothelial hypoxia may be an important secondary factor, which augments TGF $\beta$ -induced peritoneal injury. The hypoxic response is mediated partly through HIF1 $\alpha$  and the mTOR inhibitor rapamycin blocks the hypoxic-induced angiogenic effects but does not affect the direct TGF $\beta$ -mediated fibrosis and angiogenesis.

**Keywords:** hypoxia-inducible factor • transforming growth factor • adenovirus • Smad • mTOR • rapamycin • vascular endothelial growth factor • angiogenesis

#### Introduction

The peritoneal membrane provides life support for patients who rely on peritoneal dialysis (PD) as their renal replacement therapy. The removal of salt and water by ultrafiltration (UF) depends on the maintenance of a small solute gradient between the hypertonic

\*Correspondence to: Dr. Peter J. MARGETTS, Department of Medicine, McMaster University, Division of Nephrology, St. Joseph's Hospital, 50 Charlton Ave E, Hamilton, Ontario, Canada L8P 4A6. Tel.: (905) 522-1155 ext 32299 Fax: (905) 540-6589 E-mail: margetts@mcmaster.ca dialysate and the intravascular space. Over time, most PD patients develop increased solute transport across the peritoneal membrane with a concomitant decrease in UF secondary to a rapid loss of the osmotic gradient [1]. This fast peritoneal solute transport status and UF dysfunction are associated with fibrosis and angiogenesis of the peritoneal tissues [2]. Increased peritoneal membrane solute transport is associated with increased mortality in patients on PD therapy [3]. Peritoneal fibrosis has been associated with the non-physiological nature of PD solutions [4], uraemia [5] and recurrent infections [6].

Transforming growth factor (TGF)  $\beta$  is involved in the wound healing response and is essential in progressive changes in the peritoneal membrane over time. Several studies have

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demonstrated an association between peritoneal effluent TGF $\beta$  concentration and peritoneal membrane injury in patients on PD therapy [7–9]. In an animal model, we have demonstrated that adenovirus-mediated gene transfer of TGF $\beta$  to the peritoneum results in functional and structural changes similar to those seen in patients on long-term PD [10]. Peritoneal fibrosis and angiogenesis have been demonstrated to be associated with epithelial-to-mesenchymal transition (EMT) of mesothelial cells [11] and we have shown that TGF $\beta$  induces a pattern of EMT similar to that which is seen in the peritoneal tissues of PD patients [12].

Hypoxia has been linked to angiogenesis [13], fibrosis [14] and EMT [15]. The cellular hypoxic response is mediated through several factors including hypoxia-inducible factors (HIF). HIF1 $\alpha$  protein expression is tightly regulated by oxygen sensitive enzymes and in a low oxygen environment, binds to the constitutively expressed HIF1 $\beta$ , translocates to the nucleus and induces a characteristic pattern of gene expression [16]. Studies have identified an association between hypoxia and fibrosis in the liver [17], lung [18] and kidney [19]. There is extensive evidence that hypoxia directly up-regulates TGF $\beta$ 1 and independently regulates vascular endothelial growth factor (VEGF) [20, 21].

HIF1 $\alpha$  has been demonstrated to have significant fibrogenic and angiogenic effects *in vivo*. Mice with a conditional hepatic deletion of HIF1 $\alpha$  demonstrated less fibrosis in a common model of cirrhosis [22]. Furthermore, HIF1 $\alpha$  has been linked to EMT in renal tubular epithelial cells [15]. The mechanism of HIF1 $\alpha$ -associated EMT appears to involve regulation of metalloproteinases and the zinc finger regulatory proteins SNAI1 (Snail) [23], SNAI2 (Slug) [24] and Twist [25].

Rapamycin is a commonly used immunosuppressive agent that inhibits the mammalian target of rapamycin (mTOR) and has been demonstrated to down-regulate hypoxia-induced HIF1 $\alpha$  [26]. Rapamycin therefore has anti-angiogenic effects through blockade of HIF1 $\alpha$ -induced VEGF expression [27]. Because of the association between HIF1 $\alpha$  and fibrosis, rapamycin has been demonstrated to have anti-fibrotic activity in a number of mouse models including the kidney [28, 29], liver [30] and two models of systemic sclerosis [31]. Everolimus, a related mTOR inhibitor, has been demonstrated to have a positive effect in a model of cyclohexidine gluconate-induced peritoneal fibrosis [32]. Furthermore, rapamycin blocks some of the EMT effects of TGF $\beta$  in mesothelial cell culture [33].

The interaction between fibrosis, angiogenesis and hypoxia is complex and likely organ specific. Peritoneal tissues are poorly vascularized and peritoneal injury appears to follow a classic wound healing response with induction of new blood vessels and associated deposition of connective tissue [34]. This is distinct from the fibrogenic response in highly vascularized organs such as the lung or kidney. In the lung, it appears as though fibrotic foci demonstrate angiogenic stimuli but a loss of blood vessels related to increased anti-angiogenic factors [18]. The same loss of blood vessels is observed in the renal interstitium after glomerular injury [35]. We hypothesize that in peritoneal tissue, fibrogenesis induces hypoxia, and this can potentiate angiogenesis and maintain and amplify the fibrotic response.

#### Materials and methods

#### **Recombinant adenoviruses**

The construction of the adenovirus vector AdTGF $\beta$ 1 has been previously described [36]. AdTGF $\beta$ 1 expresses biologically active TGF $\beta$ 1. AdHIF1 $\alpha$  (mut) was constructed as previously described [37]. Briefly, the oxygen-dependent degradation domain (ODDD, amino acids from 392 to 511) was excised from the full-length HIF1 $\alpha$ . Missense mutations (Pro567Thr and Pro658GIn) were introduced. These mutations are necessary to fully activate HIF1 $\alpha$  [38]. We also constructed an adenovirus expressing the full-length HIF1 $\alpha$ . A null adenovirus (AdDL) was used for control.

#### **Animal studies**

Animals were treated in accordance with guideline of the Canadian Council on Animal Care. C57BI/6 mice (Harlan, Indianapolis, IN, USA) were treated with AdTGF $\beta$ 1, AdHIF1 $\alpha$ , AdHIF1 $\alpha$  (mut) or AdDL at a dose of  $1.5 \times 10^8$  plaque forming units diluted in 100 µl in PBS by intraperitoneal (i.p.) injection and animals were killed on days 10 or 21 after infection. Rapamycin (generously provided by Wyeth Canada) or vehicle was administered by oral gavage to mice treated with AdTGF $\beta$ 1 or AdDL starting on day 3 and continuing daily at a dose of 2.5 mg/kg. Before the mice were killed, 5 ml of 4.25% glucose dialysis solution (Baxter Healthcare, McGaw Park, IL, USA) was administered by i.p. injection and then recovered 15 min. later. The entire anterior abdominal wall was removed and half was taken for histology and half for RNA extraction. Omental tissue was taken and frozen in liquid nitrogen for protein extraction. The *in vivo* effects of rapamycin were confirmed in a repeat experiment.

Smad3<sup>-/-</sup> mice were generously provided by Dr. A. Roberts. These mice were generated by removal of exon 8 of the Smad3 gene in mice of background 127SV/EV x C57BL/6 as per Yang and colleagues [39]. Smad3<sup>-/-</sup> and littermate Smad3<sup>+/+</sup> mice were infected with AdHIF1 $\alpha$  (mut) at a dose of 1.5  $\times$  10<sup>8</sup> plaque forming units diluted in 100  $\mu$ l in PBS by i.p. injection and animals were killed on day 21 after infection.

#### Immunohistochemical analysis

C57Bl/6 mice, 10 days after treatment with i.p. AdTGF $\beta$ 1 or AdDL, received a tail vein injection of 60 mg/kg of pimonidazole (HPI, Inc., Burlington, MA, USA) dissolved in 0.9% saline. One hour later, anterior abdominal wall tissue was taken and stained for pimonidazole using a mAb to pimonidazole (HPI, Inc.) and a secondary Fab antimouse antibody. To assess the effects of rapamycin treatment on angiogenesis, paraffin-embedded tissues of anterior abdominal wall from these mice were sectioned and immunohistochemical analysis was performed using vWf-related antigen (Dako Corp., Carpenteria, CA, USA). The density of blood vessels in the superficial 200  $\mu$ m layer of the submesothelium and submesothelial thickness of the parietal peritoneum were analysed using Northern Eclipse software (Empix Imaging, Mississauga, ON, Canada).

#### **Cell culture**

Mesothelial cells were isolated from rat peritoneal tissue and cultured as previously described [10]. Cells were exposed to media alone or rapamycin

(provided by Wyeth Canada) at concentrations of 1, 10 or 50 ng/ml. Cells were then exposed to 2.5 ng/ml of recombinant TGF $\beta$ 1 (R&D Systems, Minneapolis, MN, USA) for 24 hrs. In a separate experiment, cells were treated with rapamycin and then grown for 12 or 24 hrs in a hypoxic environment (1% 0<sub>2</sub>) or normoxia (HeraCell incubator, Thermo Scientific, Nepean, ON, Canada).

#### Quantitative PCR

mRNA was extracted from primary rat mesothelial cells or from the peritoneum of killed animals by immersing the parietal peritoneal surface in Trizol reagent (Invitrogen, Burlington, ON, Canada) for 20 min. RNA was extracted from Trizol according to the manufacturer's instruction. RNA (1  $\mu$ g) then was reverse-transcribed using a standard protocol (Invitrogen). Quantitative real-time PCR for plasminogen activator inhibitor (PAI)-1, VEGF,  $\alpha$ -smooth muscle actin (SMA), and SNAI1, SNAI2, type 1 collagen, and HIF1 $\alpha$  was performed on mRNA using an ABI Prism 7500 Sequence Detector (Applied Biosystems, Foster City, CA, USA). mRNA quantification was referenced to Glyceraldehyde-3-phosphate dehydrogenase (GAPDH) and 18S. Data were similar with each reference gene and mRNA. Results referenced to GAPDH are shown.

#### Protein analysis

Protein (25  $\mu$ g) was electrophoresed on an 8% SDS-PAGE gel, transferred and probed with an antibody to HIF1 $\alpha$  (R&D Systems) or  $\beta$ -actin (Sigma-Aldrich, Oakville, ON, Canada). Band density was measured using Scion Image Software (Scion Corp, Frederick MD, USA). VEGF in cell culture supernatant was measured by ELISA (R&D Systems).

#### Statistical analysis

Data are shown  $\pm$  standard deviation. Differences between groups were assessed by one-way ANOVA with Tukey's *post-hoc* test. In order to test the independent effects of TGF $\beta$ 1 or hypoxia and rapamycin in mesothelial cell culture experiments we used two-way ANOVA with Tukey's *post-hoc* test.

#### Results

#### TGFβ1-induced peritoneal fibrosis leads to a hypoxic response

Ten days after infection with an adenovirus expressing active TGF $\beta$ 1 (AdTGF $\beta$ 1), we demonstrated increased cellular localization of pimonidazole in the submesothelial fibroproliferative zone (Fig. 1A–C). Pimonidazole is retained in hypoxic tissue in the presence of oxygen tension below 10 mmHg [40]. Tissue hypoxia was specific for TGF $\beta$ 1-treated animals and was not seen in the peritoneum of control adenovirus-treated animals (Fig. 1A). Animals not administered pimonidazole were used to demonstrate antibody specificity (Fig. 1C). HIF1 $\alpha$  gene expression was increased in the peritoneal tissue at day 10 after infection with AdTGF $\beta$ 1 compared with control adenovirus-treated animals. Three days after adenovirus infection, animals received the mTOR inhibitor rapamycin by oral gavage. Rapamycin did not have an effect on HIF1 $\alpha$  gene expression (Fig. 1D). Measurable increase in HIF1 $\alpha$  protein expression was delayed until day 21 (Fig. 1E and F). HIF1 $\alpha$  protein expression was significantly reduced in animals treated with rapamycin.

# Rapamycin inhibited TGFβ1-induced submesothelial angiogenesis

Control adenovirus infection did not induce any substantial histologic changes to the peritoneal membrane (Fig. 2A). Rapamycin treatment had no effect on the peritoneum in control adenovirus-treated animals (Fig. 2B). TGF $\beta$ -induced significant submesothelial thickening and angiogenesis as seen in Figure 2C. These changes are quantified in Figure 2E and F. Rapamycin had no effect on peritoneal histology 10 days after adenovirus infection, but 21 days after infection, rapamycin treatment significantly reduced the number of blood vessels and the submesothelial thickening induced by TGF $\beta$ 1. These results were confirmed in a second, separate experiment (data not shown).

# Rapamycin treatment did not affect fibrogenic gene expression

Type 1 collagen gene expression was significantly elevated in animals treated with AdTGF $\beta$ 1 (Fig. 3A) 10 days after adenovirus infection. Interestingly, treatment with rapamycin in animals infected with AdTGF $\beta$ 1 further increased collagen gene expression. Changes in VEGF gene expression after infection with AdTGF $\beta$ 1 were not significant (Fig. 3B). Gene expression of the zinc finger regulatory proteins SNAI1 (Snail) and SNAI2 (Slug) was increased in the AdTGF $\beta$ 1-treated animals (Fig. 3C). Rapamycin inhibited SNAI2 but not SNAI1 gene expression (Fig. 3D).

# Rapamycin had no effects on TGF $\beta$ 1-induced changes in mesothelial cell culture, but significantly blocked the response to hypoxia

We studied induction of HIF1 $\alpha$  after 24 hrs of exposure to recombinant TGF $\beta$ 1. TGF $\beta$ 1 did not significantly induce HIF1 $\alpha$  protein or gene expression at this time point (Fig. 4A–C) and HIF1 $\alpha$  protein and gene expression was not significantly affected by rapamycin. TGF $\beta$ 1 significantly increased VEGF gene expression in mesothelial cell culture and rapamycin had no effect on this (Fig. 4D).

As previously demonstrated [26], rapamycin significantly reduced HIF1 $\alpha$  and VEGF expression in mesothelial cells exposed to 1% oxygen for 12 or 24 hrs (Fig. 5). Hypoxia significantly up-



Fig. 1 (A–C) Sections of mouse peritoneal tissue stained for pimonidazole 10 days after injection (N = 3/group). (A) Mouse treated with AdDL showed no submesothelial thickening and no hypoxic response. (B) Peritoneum of AdTGF $\beta$ 1-treated mouse revealed increased hypoxia in the submesothelium. (C) Control section from a mouse receiving AdTGF $\beta$ 1 but no pimonidazole. All sections were taken at 200× magnification. (D) HIF1 $\alpha$  gene expression was increased by AdTGF $\beta$ 1 10 days after infection. (E) HIF1 $\alpha$  protein is up-regulated in peritoneal tissue from AdTGF $\beta$ 1-treated animals at 21 days after adenoviral infection. HIF1 $\alpha$  was significantly down-regulated by rapamycin treatment at both 10 and 21 days after adenovirus infection. (F) Representative blot is shown (N = 4 animals/group for HIF1 $\alpha$  analysis).

regulated HIF1 $\alpha$  protein expression. Rapamycin significantly decreased HIF1 $\alpha$  protein expression at both 12 and 24 hrs (Fig. 5A and B). By two-way ANOVA, hypoxia significantly increased VEGF expression after 24 hrs of hypoxia. Rapamycin significantly decreased hypoxia-induced VEGF expression in a dose-dependent manner (Fig. 5C).

# Rapamycin exacerbated TGFβ1-induced fibrogenic changes

In primary mesothelial cells, we found that TGF $\beta$ 1 significantly induced gene expression of  $\alpha$ SMA, PAI-1 and type 1 collagen

(Fig. 6). Rapamycin did not have an independent effect on fibrogenic gene expression, except for expression of type 1 collagen where rapamycin augmented the effect of TGF $\beta$ 1 (Fig. 6C).

# Overexpression of HIF1 $\alpha$ induces angiogenesis and increased submesothelial thickening in the peritoneum

The peritoneum exposed to AdHIF1 $\alpha$  (full-length HIF1 $\alpha$ ) was similar to AdHIF1 $\alpha$  (mut) (active HIF1 $\alpha$ ) 10 days after infection (Fig. 7B and C). Compared to control adenovirus-treated animals



**Fig. 2** (**A**–**D**) Masson's trichrome stained sections of the anterior abdominal wall in animals treated with (**A**) control adenovirus (DL), (**B**) AdDL + rapamycin, (**C**) AdTGF $\beta$ 1 (TGF) or (**D**) AdTGF $\beta$ 1 + rapamycin. Sections from animals 21 days after adenovirus infection are shown. Original magnification was 200×. (**E**) Submesothelial thickening and (**F**) blood vessel density in the superficial 200  $\mu$ m layer are quantified (*N* = 5 animals/group; results were confirmed in a separate experiment).

(Fig. 7A) there was increased submesothelial thickening with angiogenesis and cellular proliferation. By day 21, the changes in the peritoneum infected with AdHIF1 $\alpha$  had generally resolved (Fig. 7D). In contrast, AdHIF1 $\alpha$  (mut) induced a progressive submesothelial thickening and angiogenesis (Fig. 7E). The differences in response to AdHIF1 $\alpha$  and AdHIF1 $\alpha$  (mut) is likely a result of a difference in persistence in expression of HIF1 $\alpha$  in the omental tissue of these mice (Fig. 8A and B). AdHIF1 $\alpha$  (mut) led to prolonged expression that was measurable to 21 days after infection.

# $HIF1\alpha$ -induced expression of fibrogenic genes but not VEGF is Smad3 dependent

As the changes after AdHIF1 $\alpha$  (mut) appeared to be maximal 21 days after infection, we chose this time point to study animals deficient in the key TGF $\beta$  signalling protein Smad3 (Smad3<sup>-/-</sup>) using littermate Smad3<sup>+/+</sup> mice as control. Histologically, there were subtle differences between Smad3<sup>+/+</sup> and Smad3<sup>-/-</sup> mice 21 days after infection with AdHIF1 $\alpha$ (mut) (shown in Fig. 7F and G and quantified in Fig. 8C and D). These changes were not statistically significant when quantified; however, there was a trend to decreased submesothelial thickening and angiogenesis, suggesting that part of the HIF1 $\alpha$  response may be Smad3 dependent.

RNA was extracted from the anterior abdominal wall of C57BI/6, Smad3<sup>+/+</sup> or Smad3<sup>-/-</sup> mice treated with AdDL, AdHIF1 $\alpha$  or AdHIF1 $\alpha$  (mut) and assayed using quantitative real time PCR (Fig. 9). The expression of type 1 collagen (Fig. 9A),  $\alpha$ SMA (Fig. 9B) and SNAI1 (Fig. 9C) were all up-regulated after AdHIF1 $\alpha$  (mut) but not AdHIF1 $\alpha$  infection. The gene expression of type1 collagen and SNAI1 were both Smad3 dependent, with expression significantly reduced in Smad3<sup>-/-</sup> mice. In contrast, VEGF expression was similar in Smad3<sup>-/-</sup> and Smad3<sup>+/+</sup> mice treated with AdHIF1 $\alpha$  (mut).

#### Discussion

Our results indicate that the expanding extracellular matrix induced by TGF $\beta$  in the peritoneum is hypoxic, and this hypoxia potentiates and prolongs fibrogenesis and angiogenesis. We interpret our results to suggest that after overexpression of TGF $\beta$ 1, there is a period of TGF $\beta$ 1-driven fibrosis and angiogenesis. This is followed by a hypoxia-mediated prolongation of this initial wound healing response. The later hypoxia-driven response is inhibited by rapamycin, but rapamycin has no effect on the earlier TGF $\beta$ 1-mediated events (Fig. 10). This is supported by cell culture



Fig. 3 mRNA was extracted from the peritoneal surface of the anterior abdominal wall 10 days after adenovirus infection. Gene expression was assayed using quantitative real time PCR and normalized to GAPDH expression. The expression of (A) type 1 collagen, (B) VEGF, (C) SNA11 and (D) SNA12 genes were assessed in animals treated with AdDL (DL), AdDL + rapamycin, AdTGF $\beta$ 1 (TGF $\beta$ 1) or AdTGF $\beta$ 1 + rapamycin (N = 5/group).

experiments where rapamycin blocks hypoxia-induced VEGF, but has no effect on TGF $\beta$ 1-induced VEGF.

The interaction between hypoxia and fibrosis is complex. In lung and renal fibrosis, the prevailing paradigm suggests that vascular rarefaction is a primary event, with subsequent hypoxiadriven fibrosis [18, 35, 41]. Peritoneal tissues are comparatively less vascularized than the lung or kidney, so the expanding fibrotic tissue requires blood vessel growth to sustain this; when the expanding submesothelium outstrips its vascular supply, a hypoxic angiogenic response ensues.

The tight regulation of HIF1 $\alpha$  has been described in detail [16]. In our studies, we identified that TGF $\beta$ 1-induced peritoneal fibrosis led to both hypoxia, and increased expression of HIF1 $\alpha$ . Interestingly, in our results, TGF $\beta$ 1 had no effect on HIF1 $\alpha$  gene and protein expression in mesothelial cell culture (Fig. 4A and B). Several previous studies have identified a transient TGF $\beta$ 1mediated induction of HIF1 $\alpha$  either through protein stabilization [42] or interaction between HIF1 $\alpha$  and Smad signalling proteins [43]. These effects were seen transiently at an earlier time point (6 hrs). We looked at a later time point (24 hrs) as we were interested in the interaction between rapamycin and TGF $\beta$ 1-induced fibrogenic gene expression (Fig. 6).

The angiogenic effects of HIF1 $\alpha$  are well known [37]. HIF1 $\alpha$  has also been demonstrated to have fibrogenic effects directly through up-regulation of fibrogenic factors [44] and EMT [15]. In our experiments, we used an adenovirus expressing the full-length HIF1 $\alpha$  or a mutated HIF1 $\alpha$  with the ODDD excised [37]. AdHIF1 $\alpha$  (mut) led to prolonged expression of HIF1 $\alpha$  and persisting histologic changes (Figs 7 and 8). Specifically, HIF1 $\alpha$  overexpression led to prolonged angiogenesis and fibrosis measured by increased submesothelial thickening, blood vessel density and increased expression of fibrogenic-associated genes (Fig. 9).



**Fig. 4** Primary rat mesothelial cells were cultured and exposed to recombinant TGF $\beta$ 1 (2.5 ng/ml) and varying concentrations of rapamycin. Cells were harvested for protein or mRNA. Supernatant was taken for VEGF ELISA. (**A**) Western blot for HIF1 $\alpha$  referenced to  $\beta$ -actin. (**B**) Blots were quantified with a representative blot shown. (**C**) HIF1 $\alpha$  gene expression by quantitative real time PCR referenced to GAPDH (for both protein and gene expression of HIF1 $\alpha$ , P = NS, TGF $\beta$  versus no TGF $\beta$  and P = NS for the effect of rapamycin). (**D**) Cell culture supernatant was analysed for VEGF by ELISA. TGF $\beta$ 1-induced VEGF expression (P < 0.001 by two-way ANOVA) and again rapamycin had no significant effect on this (N = 4 samples for each group at each time point).



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**Fig. 7** Masson's trichrome stained sections from the anterior abdominal wall from C57Bl/6 mice 10 days after injection with (**A**) control adenovirus (AdDL), (**B**) adenovirus expressing full-length HIF1 $\alpha$  (AdHIF1 $\alpha$ ), (**C**) adenovirus expressing mutated, constitutively active HIF1 $\alpha$  [AdHIF1 $\alpha$  (mut)]. Anterior abdominal wall 21 days after injection with (**D**) AdHIF1 $\alpha$ , (**E**) AdHIF1 $\alpha$  (mut). In order to assess the role of TGF $\beta$  signalling in HIF1 $\alpha$ -induced peritoneal changes, AdHIF1 $\alpha$  (mut) was infected into the peritoneum of (**F**) Smad3<sup>+/+</sup> or (**G**) Smad3<sup>-/-</sup> mice and tissue taken 21 days after infection.



**Fig. 8** (A) Expression of HIF1 $\alpha$  in omental tissue from animals treated with AdHIF1 $\alpha$  or AdHIF1 $\alpha$  (mut) (n = 4 samples/group). (B) Representative blot shown. (C, D) Quantification of changes seen in the peritoneum of C57BI/6 mice treated with AdHIF1 $\alpha$  or AdHIF1 $\alpha$  (mut). (E, F) Smad3<sup>+/+</sup> or Smad3<sup>-/-</sup> mice were infected with AdHIF1 $\alpha$  (mut). Twenty-one days later, there were no significant differences in submesothelial thickness (E) or vascular density (F) (N = 5-8 animals/group).

The mechanism of HIF1 $\alpha$ -induced fibrosis is not clear. Some studies have identified a necessity for TGF $\beta$  signalling [20, 21, 45], whereas other studies demonstrate a TGF $\beta$ -independent path for HIF1 $\alpha$ -induced fibrosis [17, 46]. In order to test this in our model, we infected animals lacking the Smad3 signalling molecule with AdHIF1 $\alpha$  (mut). We have previously shown that these mice are resistant to TGF $\beta$ 1-induced fibrosis and angiogenesis, but respond normally to VEGF [47]. Gene expression of the fibrosis-associated genes type 1 collagen and SNA11 was significantly reduced in Smad3<sup>-/-</sup> mice infected with AdHIF1 $\alpha$ (mut) (Fig. 9). VEGF gene expression was not significantly different between Smad3<sup>-/-</sup> and Smad3<sup>+/+</sup> mice treated with AdHIF1 $\alpha$  (mut), suggesting that the fibrogenic effects, but not the angiogenic effects, of HIF1 $\alpha$  are Smad3 dependent.

Rapamycin is a bacterial macrolide that binds with high affinity to mTOR. It is used clinically in renal transplantation for its immunosuppressant properties [27]. mTOR activates p70 S6 kinase which is involved in regulating RNA translation and protein expression and therefore has an effect on cellular growth and proliferation [48]. Rapamycin has been demonstrated to block HIF1 $\alpha$  gene expression and thus has anti-angiogenic properties in the setting of malignancy [26]. Rapamycin has also been shown to be an effective anti-fibrotic agent in 2 models of renal injury [49, 50] and a model of systemic sclerosis [31]. Everolimus has been shown to prevent progression of peritoneal fibrosis in a chemical irritant model of peritoneal injury [32]. mTOR inhibition has been demonstrated to prevent EMT as a mechanism of anti-fibrotic activity. In mesothelial cell culture, rapamycin up-regulated E-Cadherin as an epithelial protective mechanism [33]. This was confirmed in an animal model of renal fibrosis [29]. Lamouille and colleagues did not find that rapamycin had a direct EMT inhibitory effect in cell culture [51] and we found that rapamycin had an inhibitory effect on mesothelial EMT, but only in animals lacking Smad3 signalling [47].

Our results suggest that rapamycin has a significant antiangiogenic effect when angiogenesis is driven by hypoxic response. Rapamycin did not have a significant anti-fibrotic effect, and in mesothelial cell culture, rapamycin tended to accentuate TGF $\beta$ 1-induced up-regulation of pro-fibrotic cytokines. This appears to be contrary to other published *in vitro* data [52, 53]. In



**Fig. 9** RNA was extracted from C57BI/6 mice treated with control adenovirus (AdDL), AdHIF1 $\alpha$  or AdHIF1 $\alpha$  (mut). Collagen 1A2 (**A**),  $\alpha$ SMA (**B**), SNAI1 (**C**) and VEGF (**D**) are all up-regulated by HIF1 $\alpha$  (mut) adenovirus. (**E**) mRNA from Smad3<sup>+/+</sup> or Smad3<sup>-/-</sup> mice 21 days after treatment with AdHIF1 $\alpha$  (mut) was assayed using quantitative real time PCR and normalized to GAPDH. Collagen and SNAI1 were significantly up-regulated in Smad3<sup>+/+</sup> but not Smad3<sup>-/-</sup> mice. VEGF was not differentially regulated (N = 5-8 mice/group were analysed).

Fig. 10 Hypothesized interaction between direct effects of TGF $\beta$  and secondary effects of hypoxia. TGF $\beta$  has direct effects on peritoneal tissues including EMT, cellular proliferation, fibrosis and angiogenesis. The TGF $\beta$ -induced expanding extracellular matrix is hypoxic, and this drives a similar but delayed response. The hypoxia-induced effects are blocked by rapamycin, whereas rapamycin has no effect of the TGF $\beta$ -induced effects (denoted by x). The profibrotic effects of hypoxia are Smad3 dependent. *In vitro*, rapamycin up-regulates type 1 collagen in TGF $\beta$ -treated mesothelial cells (dashed arrow).



extensive *in vivo* work, rapamycin appears to be 'anti-fibrotic' [54]. However, it is difficult to dissociate anti-angiogenic and immunomodulatory properties in these *in vivo* experiments. These other effects of rapamycin may lead to a secondary anti-fibrotic response. Previous cell culture work has generally been carried out in fibroblasts [52] or mesangial cells [55]. Our work was carried out in a primary epithelial cell type which may explain the differential response we observed. Furthermore, *in vitro* it is necessary to separate the anti-proliferative effects from the anti-fibrotic effects which may confound these results [55].

Rapamycin did block TGF $\beta$ 1-induced expression of SNAI2 (Fig. 3D) and this is a potential mechanism whereby rapamycin may inhibit mesothelial EMT and indirectly influence peritoneal fibrosis. Our results concerning the effects of hypoxia and rapamycin on peritoneal EMT are preliminary, and further experiments to detail this interaction are required.

In summary, TGF<sub>β</sub>1-induced peritoneal fibrosis leads to a secondary hypoxic response that amplifies the initial fibrosis and angiogenesis and this secondary response is sensitive to mTOR inhibition. Our findings suggest that HIF1 $\alpha$  is responsible for some of these hypoxia-associated responses.

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#### **Conflict of interest**

This research is funded by Wyeth Canada. The authors have no other conflicts with respect to this manuscript.

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