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TGF- β in the AKI to CKD Transition

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Abstract

TGF- β is a key profibrotic growth factor that is activated in acute kidney injury and associated with cellular responses that lead to the development of chronic kidney disease. The persistently injured, de-differentiated tubular epithelial cell is an important mediator of the transition from AKI to CKD. TGF- β signaling may perpetuate proximal tubule injury through de-differentiation, cell cycle arrest, and increased susceptibility to apoptosis. In addition, TGF- β signaling promotes macrophage chemotaxis, endothelial injury, and myofibroblast differentiation after acute kidney injury. Future studies that block TGF- β signaling after cessation of AKI are needed to better define its role in the progression of acute to chronic renal injury.

Keywords

acute renal injury; chronic renal disease; epithelial cells

Introduction

The incidence of acute kidney injury (AKI) and the prevalence of chronic kidney disease (CKD) continue to rise with almost 15% of Americans having CKD. Several studies have shown that AKI is an independent risk factor for both the development and progression of CKD[1]. The likelihood of AKI transition to CKD depends upon both the severity and persistence of AKI. Although renal function often improves after the initial injury, rodent AKI models demonstrate the development of structural changes which may mediate this AKI to CKD transition. The mitochondria-dense proximal tubule is the primary cell targeted by AKI. Failure of the proximal tubule to undergo repair and recovery leads to persistently de-differentiated epithelia which play an important pathophysiologic role in the development of CKD. These chronically injured proximal tubules are potent producers of profiborotic and

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pro-inflammatory growth factors which mediate deleterious responses in surrounding cellular compartments such as the microvasculature and interstitium. Transforming growth factor- β (TGF- β) is strongly induced in injured proximal tubule cells, and pre-clinical studies suggest that autocrine and paracrine TGF- β signaling may mediate the progression from AKI to CKD (Figure 1).

TGF-β and the Injured Proximal Tubule

In AKI, many proximal tubule cells are lost through cell death, and the remaining viable cells undergo de-differentiation and, ideally, proliferate and re-differentiate to reconstitute the tubule epithelia. However, in severe or repeated injury, some de-differentiated epithelial cells become growth arrested, fail to re-differentiate, exhibit altered mitochondrial function and metabolism, and produce large amounts of growth factors like TGF- β [2]. The three TGF- β isoforms (-1, -2, -3) all bind to the type II TGF- β receptor, a serine/threonine kinase. Upon ligand binding, the type II receptor heterodimerizes with the TGF- β type I receptor (T β RI) which activates both Smad-dependent and -independent pathways (e.g. MAPK, GTPases) that alter gene expression. TGF- β is secreted as an inactive homodimer, but many activators of TGF- β (matrix metalloproteases, integrin $\alpha\nu\beta6$, thrombospondin-1) are upregulated in renal injury leading to greater TGF-ß activity with increased injury. TGF- β activity can reflect the severity of initial injury, but this cytokine may also play a role in maintaining these injured proximal tubules in a de-differentiated state. TGF-B potently induces epithelial de-differentiation by Snail-dependent loss of E-cadherin in vitro[3]. TGF- β 's ability to reduce epithelial protein expression and increase mesenchymal markers (e.g. α -SMA and vimentin) is well documented. Although transient epithelial de-differentiation may be necessary for proliferation and repair, persistent de-differentiation is associated with detrimental responses such as increased matrix and cytokine production. Despite strong in vitro data supporting TGF- β -dependent de-differentiation, the in vivo data has been somewhat mixed with some studies showing that systemic blockade of TGF-ß signaling improves tubule differentiation early[4] but others showing few tubular effects after ischemia/reperfusion[5].

TGF- β also mediates strongly cytostatic effects on epithelia which may hinder proliferation that is necessary to repair the injured tubule. The persistence of G2/M-arrested proximal tubules after AKI is associated with a profibrotic epithelial phenotype and the progression to CKD[2]. Although G2/M-arrested epithelial cells secrete large amounts of TGF- β , it is unclear whether TGF- β itself induces arrest at the detrimental G2/M or at G1. Studies have shown TGF- β -dependent arrest at both checkpoints in vitro with the difference likely related to the dose of TGF- β used with lower concentrations associated with G1 arrest (0.5– 2ng/mL) and higher amounts (5–10ng/mL) with G2/M[6]. How these levels of TGF- β in vitro correlate with bioactive TGF- β in the injured kidney remains unknown.

Increased proximal tubular cell death is associated with tubular atrophy and development of CKD, and TGF- β has been shown by our group and others to sensitize epithelia to apoptosis[7]. Whether this TGF- β -induced apoptosis worsens the initial response to injury or also contributes to ongoing tubular cell loss after the initial injury needs further investigation. Overexpression of TGF- β 1 by renal tubular epithelia increased autophagy,

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severe tubular degeneration and peritubular fibrosis[8]. However, the levels of TGF- β 1 in this transgenic mouse were likely much higher than in pathophysiologic states, raising the question of whether TGF- β -dependent autophagy occurs after AKI and contributes to the CKD transition. Autophagy has protective roles in AKI, but persistent autophagy has been linked with augmented tubular atrophy and interstitial fibrosis[9]. In summary, TGF- β signaling in tubular cells promotes epithelial de-differentiation, cell cycle arrest, and reduced survival in vitro and is associated with these responses in vivo. These epithelial effects likely play a role in the AKI to CKD transition, but additional studies that inhibit TGF- β signaling after the acute phase of injury are warranted to assess the role of TGF- β specifically in AKI to CKD progression.

TGF-β and Vascular Endothelia

Loss of the renal microvasculature is recognized as another important mediator of AKI to CKD progression. Rodent studies show that severe AKI causes microvascular rarefaction and peritubular capillary loss associated with increased TGF- β signaling[10]. A systemic T β RI antibody administered at the time of I/R and for 5 weeks afterwards showed an attenuated loss in the renal microvasculature without a change in the acute tubular injury[5]. A direct role of TGF- β signaling in the vascular endothelium was demonstrated when mice haploinsufficient for T β RII had less fibrosis and preserved microvasculature after folic acid administration[11]. These data suggest that TGF- β signaling in the vascular endothelia for the vascular endothelia for tributes to peritubular capillary rarefaction, potentiating hypoxia and the development of fibrosis.

TGF-β and Interstitial Cells

The tubulointerstitium contains myofibroblasts, the predominant producers of extracellular matrix, and inflammatory cells, and both cell types are important mediators of the AKI to CKD progression. TGF- β stimulates fibroblasts and pericytes to develop into myofibroblasts and is a potent inducer of collagen production. This topic has been extensively reviewed previously and is beyond the scope of this mini-review[12]. Many different inflammatory cells contribute to AKI to CKD progression, but macrophage infiltration is strongly associated with CKD progression on human biopsies[13]. Macrophages were thought to promote fibrosis through the production of growth factors like TGF- β , but macrophage-specific deletion of TGF- β 1 did not attenuate fibrosis after AKI[14]. By contrast, T β RII deletion in macrophages did reduce fibrosis after AKI[15]. These conditional knockout mice had less macrophage infiltration with TGF- β signaling in macrophages acting as a potent chemoattractant. Thus, increased TGF- β production in the injured kidney leads to greater macrophage infiltration which may promote fibrosis through other factors.

TGF-β and AKI to CKD: Unanswered Question

There is convincing data that TGF- β mediates effects on the tubular, endothelial, and interstitial compartments which can potentially lead to the development of CKD after AKI. However, most of the data comes from either genetic or pharmacologic approaches in which TGF- β activity was inhibited before or shortly after injury. Delaying TGF- β inhibition by at

least 2 days after injury would better elucidate the role of this important growth factor in the AKI to CKD transition.

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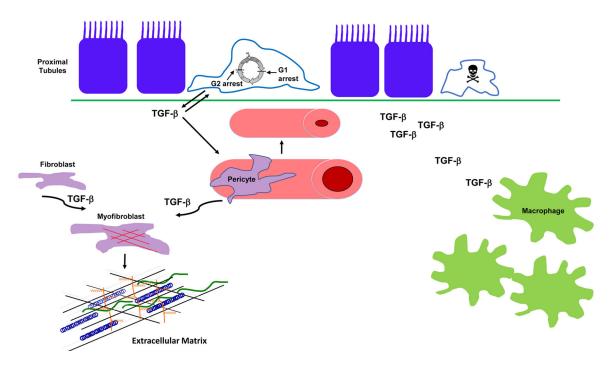


Figure 1.

TGF- β signaling mediates effects on different renal compartments that may contribute to the development of CKD after AKI. TGF- β is produced by the chronically de-differentiated proximal tubule cell after renal injury. TGF- β promotes epithelial de-differentiation, cell cycle arrest, and sensitizes epithelial cells to apoptosis during injury. TGF- β signaling can shorten peritubular capillaries and reduce their patency which may promote CKD progression through worsening hypoxia. TGF- β signaling stimulates fibroblasts and pericytes to transform into myofibroblasts, potent producers of extracellular matrix, the hallmark of fibrosis. Finally, TGF- β acts as a potent chemoattractant for macrophages and may promote further injury by augmenting macrophage infiltration. One caveat is that most of these TGF- β -dependent actions were determined by genetic or pharmacologic inhibition of TGF- β signaling either before or shortly after acute injury.