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# Induction of renal fibrotic genes by TGF- $\beta$ 1 requires EGFR activation, p53 and reactive oxygen species

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- 24 SMAD signaling

#### ABSTRACT

While transforming growth factor- $\beta$  (TGF- $\beta$ 1)-induced SMAD2/3 signaling is a critical event in the progression of 25chronic kidney disease, the role of non-SMAD mechanisms in the orchestration of fibrotic gene changes remains 26 largely unexplored. TGF-β1/SMAD3 pathway activation in renal fibrosis (induced by ureteral ligation) correlated 27 with epidermal growth factor receptor (EGFR Y845) and p53 Ser15 phosphorylation and induction of disease 28 causative target genes plasminogen activator inhibitor-1 (PAI-1) and connective tissue growth factor (CTGF) 29 prompting an investigation of mechanistic involvement of EGFR and tumor suppressor p53 in profibrotic signal- 30 ing. TGF-\(\beta\)1, PAI-1, CTGF, p53 and EGFR were co-expressed in the obstructed kidney localizing predominantly to 31 the tubular and interstitial compartments. Indeed, TGF-B1 activated EGFR and p53 as well as SMAD2/3. Genetic 32 deficiency of either EGFR or p53 or functional blockade with AG1478 or Pifithrin-a, respectively, effectively 33 inhibited PAI-1 and CTGF induction and morphological transformation of renal fibroblasts as did SMAD3 knockdown or pretreatment with the SMAD3 inhibitor SIS3. Reactive oxygen species (ROS)-dependent mechanisms 35 initiated by TGF-B1 were critical for EGFRY845 and p53Ser15 phosphorylation and target gene expression. The 36 p22<sup>Phox</sup> subunit of NADPH oxidase was also elevated in the fibrotic kidney with an expression pattern similar 37 to p53 and EGFR. EGF stimulation alone initiated, albeit delayed, c-terminal SMAD3 phosphorylation (that 38 required the TGF-β1 receptor) and rapid ERK2 activation both of which are necessary for PAI-1 and CTGF induction in renal fibroblasts. These data highlight the extensive cross-talk among SMAD2/3, EGFR and p53 pathways 40 essential for expression of TGF-\beta1-induced fibrotic target genes.

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#### 1. Introduction

Excess deposition of ECM, leading to the progressive decline in renal function, is a common pathologic hallmark of CKD [1,2]. TGF-β1 is a potent inducer of ECM synthesis and fibrosis regardless of etiology (diabetes, hypertension, ischemic injury and obstructive uropathy) [1–6]. PAI-1 and CTGF, two prominent downstream targets of TGF-β1, are major causative factors in CKD etiology and disease progression

Abbreviations: ALK, activin-like kinase;  $\alpha$ -SMA,  $\alpha$ -smooth muscle actin; BMP, bone morphogenic protein; CDK, chronic kidney disease; CTGF, connective tissue growth factor; DAB, 3,3'-diaminobenzidine; DPI, diphenyleneiodonium chloride; ECM, extracellular matrix; EGF, epidermal growth factor; EGFR, epidermal growth factor receptor; ERK, extracellular signal-regulated kinases; HB-EGF, heparin-binding EGF; IHC, immunohistochemistry; MEFs, (mouse embryo fibroblasts); MEK, mitogen-activated protein kinase kinase; NAC, N-acetyl cysteine; NOX, NADPH oxidase; PAI-1, plasminogen activator inhibitor-1; ROS, reactive oxygen species; TGF- $\beta$ 1, transforming growth factor- $\beta$ 1; UUO, unilateral ureteral obstruction.

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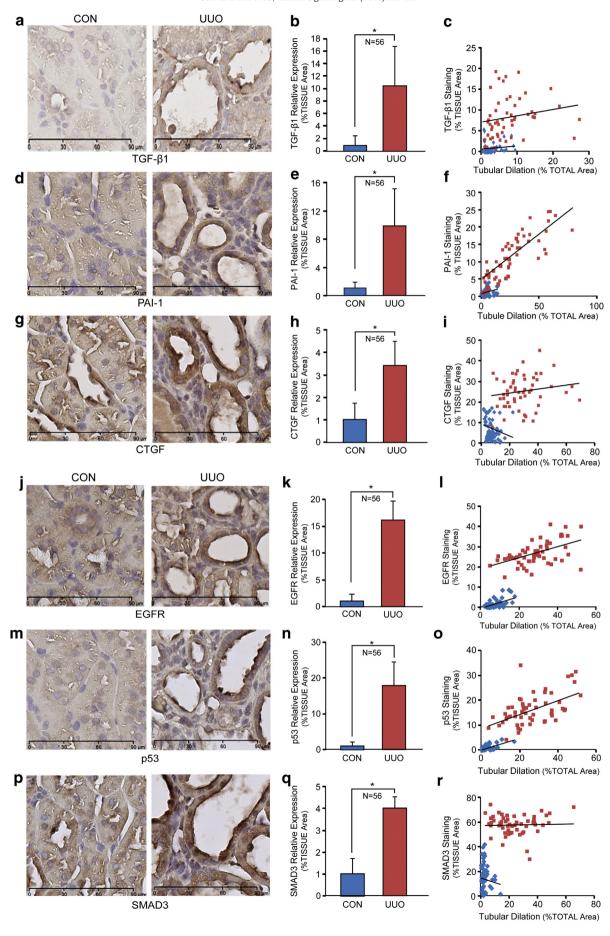
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[7–11]. PAI-1 is the major negative regulator of the plasmin-  $^{54}$  dependent pericellular proteolytic cascade while CTGF facilitates  $^{55}$  TGF- $^{64}$ 1 binding to its receptor (TGF- $^{64}$ R) and inhibits BMP-7/receptor  $^{56}$  interactions suppressing, thereby, the negative effect of BMP-7 on  $^{57}$  TGF- $^{64}$ 1/SMAD3 signaling [12,13]. Transgenic overexpression of TGF- $^{64}$ 1 sin the tubular epithelium is sufficient to induce kidney fibrosis [14] and  $^{59}$  the attenuation of disease severity in SMAD3 $^{-/-}$  mice or upon administration of TGF- $^{64}$ 1 receptor inhibitors further highlighted the pathophysiologic importance of the TGF- $^{64}$ 1/SMAD3 axis in the kidney [15,16].

Previous studies implicated the tumor suppressor p53 and the EGFR 63 in the progression of renal disease [17–20]. The mechanistic involvement 64 of such non-canonical mechanisms in the TGF- $\beta$ 1-initiated profibrotic 65 program, including the role(s) of p53 and EGFR in TGF- $\beta$ 1-dependent 66 expression of fibrosis-related genes, however, remains to be clarified. 67 This paper provides evidence that the EGFR and p53 are activated in 68 the UUO model of renal fibrosis, a largely TGF- $\beta$ 1-driven disease, and 69 required for PAl-1, CTGF and fibronectin expression in response to 70 TGF- $\beta$ 1. Generation of ROS, a causative factor in tissue fibrosis [21,22], 71 furthermore, is required for EGFR and p53 activation, consistent with 72

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