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Angiotensin II Activation of mTOR Results in Tubulointerstitial Fibrosis through Loss of N-Cadherin

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Key Words

Angiotensin II • mTOR • N-Cadherin • Proximal tubule • Tubulointerstitial fibrosis

Abstract

Background/Aims: Angiotensin (Ang) II contributes to tubulointerstitial fibrosis. Recent data highlight mammalian target of rapamycin (mTOR)/S6 kinase 1 (S6K1) signaling in tubulointerstitial fibrosis; however, the mechanisms remain unclear. Thereby, we investigated the role of Ang II on mTOR/ S6K1-dependent proximal tubule (PT) injury, remodeling, and fibrosis. Methods: We utilized young transgenic Ren2 rats (R2-T) and Sprague-Dawley rats (SD-T) treated with the Ang type 1 receptor (AT₁R) blocker telmisartan (2 mg \cdot kg⁻¹ \cdot day-1) or vehicle (R2-C; SD-C) for 3 weeks to examine PT structure and function. Results: Ren2 rats displayed increased systolic blood pressure, proteinuria and increased PT oxidant stress and remodeling. There were parallel increases in kidney injury molecule-1 and reductions in neprilysin and megalin with associated ultrastructural findings of decreased clathrin-coated pits, endosomes, and vacuoles.

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Accessible online at: www.karger.com/ajn Ren2 rats displayed increased Serine²⁴⁴⁸ phosphorylation of mTOR and downstream S6K1, in concert with ultrastructural basement membrane thickening, tubulointerstitial fibrosis and loss of the adhesion molecule N-cadherin. Telmisartan treatment attenuated proteinuria as well as the biochemical and tubulointerstitial structural abnormalities seen in the Ren2 rats. **Conclusions:** Our observations suggest that Ang II activation of the AT₁R contributes to PT brush border injury and remodeling, in part, due to enhanced mTOR/S6K1 signaling which promotes tubulointerstitial fibrosis through loss of N-cadherin. Copyright © 2011 S. Karger AG, Basel

Introduction

In the development of chronic kidney disease, progressive tubulointerstitial fibrosis contributes to declining kidney function [1, 2]. There is accumulating evi-

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dence that activation of the kidney renin-angiotensin system (RAS) and associated increases in oxidative stress are important instigators for proximal tubule cell (PTC) injury, structural remodeling, and tubulointerstitial fibrosis that contribute to progressive kidney disease [3-6]. Indeed, angiotensin (Ang) II signaling through the Ang type 1 receptor (AT_1R) induces tubulointerstitial fibrosis by downregulating E-cadherin (an epithelial phenotype marker) while upregulating alpha-smooth muscle actin and other mesenchymal markers in epithelial-mesynchymal transition (EMT) [7, 8]. Blockade of Ang II signaling through the AT₁R attenuates the progression of kidney disease in preclinical as well as clinical studies [9, 10]. However, the role of the renal RAS in eliciting these early changes and the reversibility of the process with AT₁R blockade remain unclear.

Recent data support a role for AT₁R signaling in promotion of fibrosis in coronary smooth muscle cells, cardiac tissue, and skeletal muscle through activation of the mammalian target of rapamycin (mTOR)/S6 kinase 1 (S6K1) signaling pathway [11–13]. mTOR is a serine (Ser)/ threonine (Thr) kinase that regulates transcription and protein synthesis, cell growth, proliferation, and cell phenotype transition. In studies involving cardiomyocytes and mesangial cells, blockade of the AT₁R may lead to reductions in mTOR signaling by targeting pathways that lead to reductions in NADPH oxidase subunits [13-15]. Furthermore, Ang II actions on the AT₁R contribute to PTC injury through NADPH oxidase generation of oxidative stress [16, 17]. Collectively, these data suggest a role for Ang II in injury to the proximal tubule (PT) and fibrosis through similar mechanisms that engage the mTOR/S6K1 pathway.

Data from models of diabetic and polycystic kidney disease indicate that targeting reductions in mTOR activity attenuates the progression of tubulointerstitial fibrosis [18-20]. The improvements in fibrosis following mTOR inhibition with rapamycin have been shown to be related to reductions in phosphorylation of S6K1. Recent evidence in carcinogenic models further suggests that increased mTOR/S6K1 signaling promotes loss of adherens junction markers and increased fibrosis [18, 21]. The catenin-cadherin complex is integral in cell-cell adhesion, and loss of cadherins is a critical first step in development of tubulointerstitial fibrosis and EMT [22-24]. Recent data support the importance of loss of PT-specific N-cadherin in promotion of tubulointerstitial fibrosis [23]. However, our understanding of agonists that elicit loss of adhesion is limited in tubulointerstitial fibrosis and progressive kidney disease.

Accordingly, we hypothesized that Ang II contributes to PTC injury through oxidative stress, the mTOR/S6K1 pathway, and associated loss of N-cadherin thereby promoting tubulointerstitial fibrosis. To ascertain the impact of Ang II signaling through the AT₁R on PT injury and promotion of tubulointerstitial fibrosis, we examined structural and functional properties of the PT in relation to oxidant stress, mTOR/S6K1, and the PT-specific adhesion molecule N-cadherin in the transgenic (mRen2)27 (Ren2) rats, a model that manifests elevated tissue Ang II and PTC injury related proteinuria [16, 25] treated with the AT₁R blocker telmisartan.

Animals and Methods

Animals and Treatments

All animal procedures were approved in advance by the Harry S. Truman Veterans Memorial Hospital Subcommittee for Animal Safety, as well as by the University of Missouri IACUC, and animals were cared for in accordance with NIH guidelines. Fiveweek-old male heterozygous transgenic Ren2 (n = 12) and Sprague-Dawley (SD) littermates (n = 12) were housed under standard laboratory conditions. Rats were randomly distributed into the following groups: SD control (SD-C) n = 6, telmisartantreated SD (SD-T) n = 6, Ren2 control (R2-C) n = 6, and telmisartan-treated Ren2 (R2-T) n = 6. Beginning at 7 weeks of age, treated rats received daily telmisartan (2 mg/kg) or vehicle solution in drinking water for a total of 21 days [26].

Metabolic Parameters

Systolic Blood Pressure. Systolic blood pressures (SBPs) were obtained on a subset of rats previously reported [26]. At 6 weeks of age, rats were anesthetized and instrumented with a radio telemetric transmitter (TA11PA-C40; Data Sciences, St. Paul, Minn., USA) for blood pressure monitoring. Rats recovered a minimum of 7 days before data collection. Recordings of SBP were obtained at 0, 1, 2, and 3 weeks post-treatment initiation. For each week, data were recorded continuously for 5 min at 15-minute intervals (sampling rate: 1,000 Hz) for 3 light and 3 dark 12-hour cycles.

Urine Measures. Both creatinine and protein concentrations in urine were analyzed on an automated clinical chemistry analyzer (Olympus AU680) using commercial assays [16, 25]. Creatinine was determined using an automated Jaffe reaction assay and urine protein using an automated colorimetric assay.

Transmission Electron Microscopy

Renal cortical tissue was thinly sliced and placed immediately in primary transmission electron microscopy (TEM) fixative as previously described [16, 25]. Sections (85-nm) were stained with 5% uranyl acetate, and Sato's triple-lead stain and a TEM (model JEM 1400; Joel, LTD., Tokyo, Japan) were utilized. To maintain uniformity, we examined S-1 segments of the PT (n = 4 per group) with identifiable microvilli that were immediately adjacent to the glomeruli.

Immunohistochemistry

Kidney cortical tissue was harvested and prepared as previously described [16]. Briefly, 4- μ m sections were incubated with primary antibodies overnight at room temperature. Then slides were washed and incubated with 1:300 secondary antibodies for 4 h, and the images were captured with a bi-photon confocal laser scanning microscope. Images were captured by LSM imaging system and signal intensities quantified by MetaVue as average gray scale intensities.

NADPH Oxidase Activity. NADPH oxidase activity was determined in kidney cortical tissue as previously described [16, 25]. Briefly, NADPH oxidase activity was determined by measuring the conversion of radical detector using spectrophotometry (450 nm).

3-Nitrotyrosine. Briefly, $4-\mu m$ kidney sections were incubated with 1:150 primary rabbit polyclonal anti-nitrotyrosine antibody overnight (Chemicon, Temecula, Calif., USA) as previously described [16, 25]. Sections were then washed and incubated with secondary antibodies, biotinylated linked, and Streptavidin-HRP for 30 min each. After several rinses with Tris-buffered saline Tween-20 (TBST), diaminobenzidine was applied for 8 min, sections rinsed several times with distilled water, and counterstained with hematoxylin for 80 s, dehydrated, and mounted with a permount. The slides were inspected under a bright-field (Nikon 50i) microscope, the ×40 images captured with a Cool Snapcf camera, and signal intensities quantified by MetaVue.

Western Blot Analysis

Sodium dodecyl sulfate polyacrylamide gel electrophoresis (SDS-PAGE) was performed on kidney homogenates. Samples (40 μ g/lane) were separated and transferred to polyvinylidene difluoride membranes. Blots were blocked with 5% BSA for 3 h and then washed briefly in TBST and incubated overnight at 4°C with the following primary antibodies: S6K1, kidney injury molecule-1 (KIM-1), N-cadherin, and neprilysin (1:1,000 dilution; Cell Signaling Technology, Inc.). After rinsing, blots were incubated with horseradish peroxidase-conjugated secondary antibodies (1:10,000, Jackson Immunoresearch) for 1 h at room temperature. The bands were visualized by chemiluminescence, and images were recorded using a Bio-Rad image analysis system. The quantified proteins were normalized to the density of total protein for each sample as determined by amido black, performed using Quantity One software.

Light Microscopic Analysis of Tubulointerstitial Fibrosis

Five-microns thick paraffin sections were mounted on glass slides and stained with Verhoeff-Van Gieson (VVG) stain, which stains collagen fibers pink, to evaluate interstitial fibrosis, as previously described [16, 25]. The relative amount of collagen within 10 representative regions of interest was determined with the aid of MetaVue Software and an average value was recorded for each kidney sample and expressed as arbitrary units. Samples from 5 rats from each of the 6 treatment groups were analyzed.

Statistical Analysis

This investigation was powered based on prior sensitivity and variability measurements of non-log transformed proteinuria to achieve a significance of p < 0.05 with a power of 0.8 [16, 25]. All values are expressed as mean \pm standard error. Statistical analyses were performed in SPSS 13.0 (SPSS Inc., Chicago, Ill., USA) using ANOVA with Fisher's LSD as appropriate.

Results

Ang II Contributes to Elevations in SBP and Proteinuria in the Transgenic Ren2 Rat

As reported elsewhere [26], there were increases in SBP in the Ren2 rats compared to age-matched SD controls, findings were prevented following 3 weeks of treatment with the AT₁R blocker telmisartan in the Ren2 rats. There were parallel increases in non-log transformed proteinuria in the Ren2 rats (4.83 \pm 0.64 mg/mg urine protein/creatinine) compared to SD controls (1.68 \pm 0.29 mg/mg; p < 0.05), and improvement was observed with telmisartan treatment (2.77 \pm 0.18 mg/mg; p < 0.05).

Ang II Contributes to PTC Oxidative Stress through Activation of the AT_1R in the Transgenic Ren2 Rat

Ang II activation of the AT₁R has been implicated in development and progression of kidney disease through increases in oxidant stress and pro-inflammatory mechanisms [27-29]. In this regard, semi-quantitative immunohistochemical analysis showed increased density of PT AT₁R in the Ren2 rats compared to SD controls. Telmisartan treatment resulted in reduced expression of AT₁R in Ren2-treated rats (fig. 1a). In parallel with the increases in AT₁R receptor expression, there were increases in 3-nitrotyrosine (3-NT), a marker for peroxynitrite (ONOO⁻) formation in Ren2 rats [16, 25]. ONOO⁻ is a highly reactive oxidant species that can be formed endogenously by the interaction of nitric oxide (NO) and superoxide anion (O_2^{-}) , and this product reacts readily with tyrosine residues of proteins to form 3-NT. Renal cortical PT 3-NT content was higher in the Ren2 rats compared to SD controls, findings improved with telmisartan treatment in the Ren2 rats (fig. 1b).

Enhanced Ang II Generation of Reactive Oxygen Species in the PT Is Mediated through AT₁R Activation of NADPH Oxidase in the Transgenic Ren2 Rat

The NADPH oxidase enzyme complex is an important regulator of Ang II-mediated generation of oxidant stress in the kidney [17, 29]. Consistent with this notion, there were increases in cortical tissue NADPH oxidase activity (fig. 2a) and PT intensity of subunits Rac1 and p47^{phox} in the Ren2 rats compared to SD controls. These collective findings improved with telmisartan treatment in the Ren2 rats (fig. 2b).

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Fig. 1. Angiotensin II contributes to PTC oxidative stress through activation of the AT_1R . **a** Representative images of semi-quantitative immunohistochemical analysis and localization of the AT_1R to the PT with corresponding measures to the right. **b** Representative images of PT 3-NT content, a marker of

peroxynitrite formation (ONOO⁻), with corresponding measures to the right. * p < 0.05 when compared to age-matched Sprague-Dawley controls (SD-C); [†] p < 0.05 when telmisartan-treated R2 rats (R2-T) are compared to age-matched Ren2 controls (R2-C). Scale bar = 50 μ m.

Ang II Contributes to PT Brush Border Injury in the Transgenic Ren2 Rat

Recent work has highlighted that Ang II contributes directly to PT injury [16, 17, 28]. PTC injury is characterized by release of enzymes such as KIM-1, a marker of brush border injury [30, 31], and neprilysin (e.g. neutral endopeptidase 24.11), an enzyme localized to the brush border and responsible for degrading Ang II [32]. Both are abnormally released from the endosomal brush border region from lysosomes [33]. KIM-1 is upregulated during PT injury, a finding observed in the Ren2 rats compared to SD controls, and improvement was observed with telmisartan treatment in the Ren2 rats (fig. 3a). Neprilysin is a brush border enzyme which cleaves and downregulates during PT injury. Here, neprilysin decreased in the Ren2 rats compared to SD controls and showed an improved trend with telmisartan treatment in the Ren2 rats (fig. 3b).

PTC brush border injury contributes to reductions in endosomal protein reabsorption. In the Ren2 rats, there



Fig. 2. Angiotensin II activation of PTC NADPH oxidase. **a** Total NADPH oxidase activity. **b** Depicts representative images of semi-quantitative immunohistochemistry analysis and localization of NADPH oxidase subunits Rac1 and p47^{phox} to the PT with

corresponding measures of intensity below. * p < 0.05 when compared to age-matched Sprague-Dawley controls (SD-C); [†] p < 0.05 when telmisartan-treated R2 rats (R2-T) are compared to age-matched Ren2 controls (R2-C). Scale bar = 50 μ m.

were reductions in PT-specific megalin by semi-quantitative immunohistochemical analysis compared to SD controls, a finding that showed improvement with telmisartan treatment (fig. 3c). These findings were corroborated on ultrastructural analysis of the apical PT wherein there were reductions in clathrin-coated pits, endosomes, and apical vacuoles (arrows) within the endosomal regions of the Ren2 rats compared to SD controls. Similarly, improvement was observed in the telmisartantreated Ren2 rats (fig. 3d).

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Ang II Contributes to Activation of Renal mTOR Signaling in the Transgenic Ren2 Rat

Recent data highlight an important role for targeting mTOR signaling in tubulointerstitial fibrosis and proteinuria in rodent models of diabetic and polycystic kidney disease [18–20]. Consistent with this notion, there were increases in mTOR and Ser²⁴⁴⁸ phosphorylation of mTOR (p-mTOR) in the Ren2 rats compared to SD controls, findings improved with telmisartan treatment (fig. 4a, b). Ser²⁴⁴⁸ phosphorylation serves as a marker of

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Fig. 3. Angiotensin II contributes to PTC brush border injury. Western blot analysis: **a** KIM-1, a type 1 membrane protein localized to PT epithelial cells expressed at high levels in human and rodent kidneys after injury, and **b** neprilysin (e.g. neutral endopeptidase), a PT brush border enzyme responsible for processing filtered peptides and cleaved during PT injury. **c** Representative images from semi-quantitative immunohistochemistry analysis of PT-specific megalin with corresponding measures of intensities to the right. Scale bar = $50 \ \mu$ m. * p < 0.05 when compared to

age-matched Sprague-Dawley controls (SD-C); [†] p < 0.05 when telmisartan-treated R2 rats (R2-T) are compared to age-matched Ren2 controls (R2-C). **d** Ultrastructural analysis of the PTC with TEM. Representative images depict the endosomal region of the Ren2 control (R2-C) model (middle panel), that demonstrates decreased clathrin-coated pits, endosomes, and apical vacuoles (arrows) compared to the SD-C models (left panel): note that telmisartan treatment in the R2 (R2-T; right panel) restores the alterations compared to the R2-C.

activation and engagement of the main downstream effector of the mTOR pathway S6K1. In this context, there were concurrent increases in total S6K1 in the Ren2 rats compared to SD controls with an improved trend with telmisartan treatment in the Ren2 rats (fig. 4c).

Ang II Contributes to Loss of Adhesion, PT Basolateral Remodeling, and Tubulointerstitial Fibrosis in the Transgenic Ren2 Rat

Activation of mTOR/S6K1 contributes to promotion of tubulointerstitial fibrosis and a process characterized by loss of adherens junctions [18–21, 34]. In the Ren2 rats, there were reductions in the PT-specific N-cadherin compared to SD controls, findings improved with telmisartan treatment in the Ren2 rats (fig. 5a). Consistent with loss of adhesion as a promoter of fibrosis, the loss of PT N-cadherin occurred contemporaneous with increases in tubulointerstitial fibrosis in the Ren2 rats compared to SD controls, again improved with telmisartan treatment in the Ren2 rats (fig. 5b).

PT epithelial cell promotion of tubulointerstitial fibrosis may occur through EMT with remodeling of the basilar region and transition of the PT to a fibroblast phenotype [1, 2, 8]. Early stage EMT is structurally characterized by loss of basal polarity and basement membrane thickening on ultrastructural analysis. In the Ren2 model, consistent with reductions in endosomal clathrincoated pits, endosomes, and vacuoles (fig. 5c), there were reductions in basilar lysosomes (fig. 5c, top panel) with parallel loss in basal polarity (fig. 5c, bottom panel). Loss of polarity was manifested by ultrastructural remodeling consisting of mitochondrial fragmentation characterized by loss of elongation and spherical enlargement with mitochondrial matrix abnormalities. Further, there was basement membrane thickening and elongated canalicular plasma membrane infoldings characterized by electron-dense finger-like protrusions along the canaliculi in Ren2 renal tissue, collective findings were not observed in the SD or the telmisartan-treated Ren2 rats.

Discussion

Results from this investigation support a role for Ang II in eliciting PTC injury and mTOR-associated loss of PTC adhesion and tubulointerstitial fibrosis. Our current observations extend evidence in smooth muscle cells, cardiac tissue, and skeletal muscle cells that Ang II engages mTOR/ S6K1 through a PI3-K/Akt-dependent pathway [8, 11–13]. Herein, we report that in the transgenic Ren2 model, a

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model of increased tissue Ang II and proteinuria, there were increases in renal tissue levels of NADPH oxidase and dependent increases in reactive oxygen species (ROS). Furthermore, it also displayed PT brush border injury with increases in KIM-1 and reductions in neprilysin with parallel increases in p-mTOR, mTOR and S6K1. Contemporaneous with these findings, there were associated reductions in the PT-specific adhesion molecule N-cadherin and ultrastructural findings of PT remodeling consistent with EMT and tubulointerstitial fibrosis. Collectively, our findings were largely improved with AT₁R blockade, supporting a role for AT₁R-mediated signaling through mTOR/S6K1 in PT injury and tubulointerstitial fibrosis.

Blockade of the AT_1R reduces NADPH oxidase activity and the Nox subunit Rac1, a small GTP-ase [10, 14, 15]. Rac1 is a critical G-protein that not only contributes to activation of the enzyme complex NADPH oxidase and generation of oxidative stress in PT cells, but also functions as an important regulator of endocytosis of albumin in the PT [17, 35]. Our finding that AT_1R blockade reduced subunits Rac1, $p47^{phox}$ and NADPH oxidase-dependent generation of 3-NT in the Ren2 further supports a role for Ang II engagement in NADPH oxidase generation of oxidant stress in the PT.

Recent data also highlight a potential role for Ang II in regulating the mTOR pathway and in PT albumin endocytosis that is mediated, in part, via regulation of megalin [11-13, 35, 36]. Megalin is involved in albumin reabsorption through a retrieval mechanism wherein albumin is transported through an endosomal/lysosomal degradation pathway and degraded into its constituent polypeptides and amino acids [37, 38]. In this context, our data extend findings from PT culture models in recent studies [17, 35] into an in vivo model (e.g. the Ren2), where Ang II-dependent signaling through the AT₁R reduces megalin expression that occurs in conjunction with ultrastructural findings of reduced endosomal lysosomes and increases in proteinuria [25, 35, 37–39]. Although the exact mechanism remains unclear, our data would suggest that this reduction in megalin expression may be mediated, in part, through an AT₁R-dependent signaling pathway.

In this investigation, renal cortical KIM-1 levels were increased in Ren2 rats and normalized with AT₁R blockade. Although KIM-1 was originally described to be induced in the post-ischemic kidney, tubular KIM-1 induction has also been reported recently in models of proteinuric and polycystic kidney disease [28, 30]. Recent data suggest that KIM-1 expression is increased in the Ren2 rats and attenuated following treatment with AT₁R blockade or mitogen-activated protein kinase inhibition [28]. This is a

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Fig. 4. Angiotensin II contributes to activation of PTC mTOR signaling. **a**, **b** Representative images from semi-quantitative immunohistochemistry analysis and localization of PT mTOR and Ser²⁴⁴⁸ p-mTOR with corresponding intensities below. Scale

bar = 50 μ m. **c** Western blot analysis of S6K1 the main effector peptide of mTOR. * p < 0.05 when compared to age-matched Sprague-Dawley controls (SD-C); [†] p < 0.05 when telmisartan-treated R2 rats (R2-T) are compared to age-matched Ren2 controls (R2-C).

particularly important finding in that mitogen-activated protein kinase is known to converge with mTOR signaling through phosphorylation of S6K1. Our data further highlight a role for Ang II in PT injury as the observed improvement in KIM-1 following AT₁R blockade was temporally related to increases in neprilysin (e.g. neutral endopeptidase), a brush border enzyme responsible for processing peptides, such as Ang II, that is reduced during injury to

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the PT [33]. Our collective findings suggest that increased KIM-1 and reduced neprilysin expression represent markers of brush border injury and targets for Ang II-dependent generation of ROS and altered mTOR/S6K1 signaling in PT injury.

Targeting reductions in mTOR/S6K1 activity with rapamycin treatment improves tubulointerstitial fibrosis and proteinuria in rodent models of diabetic nephropa-



Fig. 5. Angiotensin II contributes to loss of PT N-cadherin, basolateral remodeling and tubulointerstitial fibrosis. **a** Western blot analysis of the PT-specific adhesion molecule N-cadherin. **b** Verhoeff-Van Gieson (VVG) stain for elastin and collagen with measures of tubulointerstitial fibrosis to the right. Scale bar 50 μ m. * p < 0.05 when compared to age-matched Sprague-Dawley controls (SD-C); [†] p < 0.05 when telmisartan-treated R2 rats (R2-T) are compared to age-matched Ren2 controls (R2-C). **c** Representative images from ultrastructural analysis of TEM for the S-1

region of the PT. Top panel depicts lysosomes in the basal region of the PT wherein there is loss of electron dense lysosome (arrows) in the R2-C model (middle panel) compared to agematched SD-C, restored with telmisartan treatment in the Ren2. Bottom panel depicts elongated canalicular plasma membrane infoldings, loss of basal polarity, elongated mitochondria and basement membrane thickening in the basal region of the R2-C compared to SD-C, findings improved in the telmisartan-treated R2 model. thy and polycystic kidney disease [18–20]. The activity of mTOR appears to be highly regulated under conditions of high nutrient/energy levels, hypoxia, or various hormones such as Ang II, as suggested by our study, which allow mTOR kinase activity to be turned on promoting cell phenotype transition and fibrosis. Ser²⁴⁴⁸ p-mTOR serves as a marker for downstream activation of S6K1 as well as mTOR inhibition. While our study did not include a direct inhibitor of Ser²⁴⁴⁸ p-mTOR such as rapamycin, our finding that blockade of the AT₁R in this transgenic RAS model inhibits total and Ser²⁴⁴⁸ phosphorylated mTOR further support a role for Ang II in mTOR/S6K1 signaling in the kidney.

Our observation that AT₁R blockade in the Ren2 rat also led to improvements in tubulointerstitial fibrosis temporally related to improvements in p-mTOR further support a role for Ang II in fibrosis. Ang II actions on the AT₁R have been shown to regulate collagen synthesis and extracellular matrix protein synthesis as well as through generation of ROS [29, 40, 41]. Under steady state conditions, PTCs are attached to each other and to the basement membrane through specialized junctional complexes (adherens junctions) that are susceptible to ROS and include molecules such as cadherin. Recent data suggest that the cadherin present in the kidney [23, 24] and specific to the PT is N-cadherin [23]. N-cadherin in the PT has been shown to bind cytoskeletal components that provide a structural foundation for adherens junctions. Of note, cadherins not only function as static structural components of adherens junctions but also play a role in cell-signaling pathways [42] suggesting that cadherins may be a target of Ang II and mTOR signaling. Our finding that AT₁R blockade improves the PT-specific N-cadherin in the Ren2, support a role for Ang II activation of mTOR/S6K1 and loss of N-cadherin in association with EMT and tubulointerstitial fibrosis.

Recent evidence suggests EMT as a potential initial mechanism for tubulointerstitial fibrosis in human and other species [1, 2, 8]. Our findings that loss of N-cad-

herin occurred contemporaneous with ultrastructural remodeling of the basilar region of the PT in the Ren2 rats is consistent with structural changes of EMT and fibrosis. The finding that these changes were improved with AT₁R blockade supports a potential role for Ang II. Although our findings are ultrastructural in nature and not related to the alterations in EMT transcriptome. Ultrastructural changes of early-stage EMT are characterized by loss of basal polarity and basement membrane thickening on ultrastructural analysis [2, 25, 42]. Our observations in the Ren2 kidney of reductions in basilar lysosomes with parallel loss in basal polarity, mitochondrial fragmentation, basement membrane thickening and canalicular infoldings of the plasma membrane with improvements following AT₁R blockade support a direct role for Ang II actions on loss of N-cadherin and promotion of structural changes consistent with EMT and tubulointerstitial fibrosis.

Thereby, this preclinical investigation highlights a novel mechanism in eliciting tubulointerstitial fibrosis. The clinical/translational importance of our study resides in the identification of an AT₁R-mediated mechanism at an early juncture in the development of tubulointerstitial fibrosis, a process the dictates progressive kidney disease.

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Disclosure Statement

The authors have no conflicts of interest to disclose.

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