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The cell membrane repair protein MG53 modulates transcription factor NF-κ**B signaling to control kidney fibrosis**

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Abstract

Kidney fibrosis is associated with the progression of acute kidney injury to chronic kidney disease. MG53, a cell membrane repair protein, has been shown to protect against injury to kidney epithelial cells and acute kidney injury. Here, we evaluated the role of MG53 in modulation of kidney fibrosis in aging mice and in mice with unilateral ureteral obstruction (UUO) a known model of progressive kidney fibrosis. Mice with ablation of MG53 developed more interstitial fibrosis with age than MG53-intact mice of the same age. Similarly, in the absence of MG53, kidney fibrosis was exaggerated compared to mice with intact MG53 in the obstructed kidney compared to the contralateral unobstructed kidney or the kidneys of sham operated mice. The ureteral obstructed kidneys from MG53 deficient mice also showed significantly more inflammation than ureteral obstructed kidneys from MG53 intact mice. In vitro experiments demonstrated that MG53 could enter the nuclei of proximal tubular epithelial cells and directly interact with the p65 component of transcription factor NF-κB, providing a possible explanation of enhanced inflammation in the absence of MG53. To test this, enhanced MG53 expression through engineered cells or direct recombinant protein delivery was given to mice subject to UUO. This reduced NF-κB activation, inflammation and attenuated kidney fibrosis. Thus, MG53 may have a therapeutic role in treating chronic kidney inflammation and thereby provide protection against fibrosis that leads to the chronic kidney disease phenotype.

SUPPLEMENTARY MATERIAL

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AUTHORS CONTRIBUTION

H. Li, P. Duann, P. Lin, J. Ma and B.H. Rovin conceived/designed the studies. H. Li, P. Duann, P. Lin, Z. Li and X. Zhou conducted experiments, acquired data and data analysis. P. Duann, P. Lin and B.H. Rovin interpreted the results. P. Lin and B.H. Rovin drafted, revised and approved the manuscript. All authors had read and approved the final version.

DISCLOSURES

J. Ma has an equity interest in TRIM-edicine, Inc., which develops rhMG53 for treatment of human diseases. Patents on the use of MG53 are held by Rutgers University and The Ohio State University. All other authors declare no competing interests.

Supplementary information is available on Kidney International's website.

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Keywords

TRIM72; inflammation; myokine; cytokine; chronic kidney disease

INTRODUCTION

Kidney dysfunction, which can be classified as acute kidney injury (AKI) or chronic kidney disease (CKD), has grown into an epidemic in older populations. In 2017, the prevalence of CKD reached 14.5% of people aged 65 and over, resulting in Medicare expenditures of over \$84 billion for treatment.^{1, 2} AKI and CKD are more common in older individuals, mainly due to increased susceptibility to injury plus a diminished ability to repair the aging kidney. Clinical studies have shown that CKD is a major risk factor for AKI and conversely, episodes of AKI can accelerate the progression of CKD toward end-stage kidney disease (ESKD). Currently, the treatment of AKI is mainly supportive, and the treatment of established CKD is focused on slowing progression by through inhibition of the reninangiotensin-aldosterone system, and now possibly through sodium-glucose co-transporter-2 (SGLT2) inhibition.^{3, 4}

Depending on the severity of injury, proximal tubular cells, the major target of acute injury, may undergo changes in cell cycle progression,^{5, 6} metabolism,⁷ secretion of pro-inflammatory and profibrotic cytokines, or partial epithelial-mesenchymal transition (EMT) , which will determine whether remodeling/repair will be successful, or maladaptive and result in kidney fibrosis, a hallmark of $CKD^{9, 10}$ This remodeling generally occurs on a background of renal inflammation, diminished vascular supply and production of extracellular matrix (ECM) proteins, and includes loss of epithelial cells and accumulation of collagens, α-smooth muscle actin (SMA) and fibronectin, and has a high correlation with deterioration of kidney function.

Chronic inflammation leads to fibrotic remodeling that may also underlie the transition from AKI to CKD.^{11–13} Cumulative research suggests the involvement of the NF- κ B

transcription factor in the pathogenesis of renal inflammation caused by infection, injury, or transplantation.^{11,12, 14} Activation of the canonical NF- κ B pathway starts with activation of IκB kinase (IKK) which leads to phosphorylation and degradation of IκBα and the nuclear translocation of NF- κ B heterodimers.¹⁵ Activation of NF- κ B signaling in kidney epithelial cells and infiltrating immune cells can be elicited by pathophysiological triggers such as exposure to lipopolysaccharides (LPS) or ischemia-reperfusion injury (IRI).16 Molecular profiling studies reveal *nfkb1* as a major driver of renal fibrosis.¹⁷ In addition to renal tubular cells, innate immune cells such as macrophages and dendritic cells also contribute to renal injury, inflammation and fibrotic remodeling.13, 18–20

Growing evidence suggests that muscle-derived secretory factors, myokines, modulate systemic physiology via tissue crosstalk to influence the progression of kidney diseases.²¹ MG53 (also named TRIM72) is a muscle-enriched tripartite motif (TRIM) protein with a critical function in cell membrane repair.^{22, 23} TRIM family proteins have diverse functions ranging from regulation of immune signaling to tissue repair.²⁴ MG53-mediated membrane repair is involved in mitigating acute injury to skeletal muscle, 2^5 kidney, 2^6 heart, 2^7 lung, 2^8 brain,29, 30 and skin.31 Our previous studies identified a low level of MG53 expression in the kidney, which is a key component in AKI protection, and mice deficient in MG53 $(mg53^{-/-})$ were more susceptible to stress-induced AKI²⁶. We also recently demonstrated that sustained elevation of MG53 in the circulation enhances tissue injury-repair and regeneration.32 The relative roles of kidney-specific and extrinsic circulating MG53 in modulating renal fibrosis during progressive CKD are still unknown.

Recently, an emerging role of MG53 in modulating tissue protection through inhibition of inflammation, especially via modulation of NF-κB signaling has evolved. As examples, MG53 attenuates LPS-induced neurotoxicity and neuroinflammation by inhibiting the TLR4/NF-κB pathway33 and cardiac MG53 regulates KChIP2 expression to control electrophysiological remodeling during cardio-hypertrophy.34 In addition, we demonstrated a dual function of MG53 in preventing a maladaptive hyper-inflammatory response during a sub-lethal influenza A virus (IAV) infection, characterized by excessive IFNβ production, via suppression of IRF3/NF-κB activation, and inhibition of aberrant intracellular Ca^{2+} signaling in macrophages.³⁵ Furthermore, upon a lethal dose of IAV infection, MG53 was shown to prevent mortality and lung damage, not by directly affecting viral titers *per se*, but by mitigating cytokine storm (IFNβ, IL-6, and IL-1β) through negative regulation of the NLRP3 inflammasome and prevention of pyroptosis of the lung.³⁶ Moreover, high dose chronic treatment of exogenous MG53 was linked to suppression of NF-κB-mediated inflammation in the aged heart.³⁷ These studies suggest MG53 can modulate aberrant NF-κB signaling during inflammation, but whether this occurs in the context of kidney inflammation has not been studied.

We postulated that MG53 regulates kidney inflammation by modulating the transcriptional activity of NF-κB, and in so doing can attenuate the kidney fibrosis that occurs as a consequence of chronic inflammation. This study was undertaken to address this hypothesis.

METHODS

Animals Studies

All experiments with animals adhered to the NIH Guide for the Care and Use of Laboratory Animals, and followed protocols approved by the Ohio State University IACUC. Agematched *wild type* (wt) or $mg53^{-/-}$ non-littermates of the 129S1/SvImJ strain were used²² to establish the role of MG53 in modulation of age-dependent and unilateral ureteric obstruction (UUO)-induced kidney fibrosis. C57B6/J mice (9–10 weeks) were purchased from Jackson Labs.

For kidney injury studies, mice (9–12 weeks) were subjected to UUO to induce kidney fibrosis. Mice were anesthetized via isoflurane $(1.5 - 2.0\%)$ to ensure a deep plane of anesthesia. The UUO procedure was performed by ligating the left ureter with 5–0 surgical silk twice according to published protocols.³⁸ Sham mice only had an abdominal skin incision. To evaluate the effect of rhMG53 on obstructive kidney injury, one group of UUO mice received rhMG53 (2 mg/kg) via tail vein injection beginning immediately after UUO surgery (day 0) and then daily for 7 days, with two additional doses on days 9 and 11. A control group of UUO mice received saline injections according to the same schedule. Mice were sacrificed on day 12 post UUO surgery, kidneys were perfused *in situ* with PBS, and tissue samples from the UUO kidney and contralateral kidney (CLK) were collected for histology and western blot analysis.

In separate studies, wt and $mg53^{-/-}$ mice underwent UUO or sham surgery and were sacrificed day 7 post-operation. MG53 was given to mice using a cell-based therapy approach.39 RAW 264.7 macrophages were infected with doxycycline (DOX)-dependent Ad-tPA-MG53-mcherry viral particles for 24 hr. The infected cells (at a concentration of 2 \times 10 δ /100 ul saline per mouse) were tail-vein-injected into mice immediately after surgery (day 0 injection). Mice received four injections of Ad-tPA-MG53 transduced RAW cells on days 0, 1, 3 and 6. Immediately after the day-0 cell injection, mice were randomized into a group that received and a group that did not receive DOX (2 mg/ml in 5% sucrose solution) in their drinking water for 7 days and then sacrificed. Tissue samples from the UUO kidney and contralateral kidney (CLK) were collected for histology, RNA and protein analyses.

Statistical Analyses

Data are expressed as means \pm SD. Comparison within groups was made by Student's t-test when comparing two experimental groups and by one-way analysis of variance (ANOVA) for more than two groups (Graphpad Prism 8.2), followed by ad-hoc either Bonferroni's or Holm-Sidak's multiple comparison test. A value of $P < 0.05$ was considered statistically significant.

Supplementary Methods

Full methods including histologic evaluation, serum creatinine (Scr) measurements, plasmid constructs, cell culture and primary tubular epithelial cell isolation, adenovirus preparation and cell infection, NF-κB p65 nuclear translocation assay, immunohistochemistry and confocal images, NF-κB Luciferase Reporter Assay, cytokines ELISA, tissue fractionation,

immunoblotting, co-immunoprecipitation (IP), quantitative real-time PCR (qRT-PCR), rhMG53 production are in the Supplementary Methods.

RESULTS

Mice with ablation of MG53 develop age-dependent renal fibrosis

We compared the effect of aging on kidney function and fibrosis in $mg53^{-/-}$ and wild type (wt) mice. We confirmed our previous observation that while Scr levels of young mice (2-month) with and without MG53 were not significantly different, older (16–20 month) $mg53^{-/-}$ mice exhibited a significantly higher Scr than age-matched wild-type controls (Figure 1a). Based on trichrome staining, we found that $mg53^{-/-}$ mice had more renal fibrosis compared with the wt mice, starting at the age of 5-months and continuing through the age of 20 months (Figure 1b, 1c). Immunohistochemical staining was used to evaluate the impact of MG53 ablation on infiltration of leukocytes, monocytes and macrophages in 2-month and 5-month old mouse kidneys (Figure 1d, e). Kidneys from young wild-type and $mg53^{-/-}$ mice had similar levels of intra-renal leukocytes, but significantly more inflammatory cells were found in $mg53^{-/-}$ kidneys than *wt* kidneys in 5-month old mice.

Moreover, immunohistochemistry for α-SMA and fibronectin, two extracellular matrix proteins commonly found in areas of kidney fibrosis, showed enhanced (3–8 fold) staining in the glomeruli and tubulointerstitium of kidneys from 10-month old $mg53^{-/-}$ mice compared to 10-month old wt mice (Supplementary Figure S1).

UUO induces aggravated renal inflammation and fibrosis in *mg53−/−* **mice**

Kidney MG53 levels detected by immunoblotting significantly increased 7-days after UUO (Figure 2a). The $mg53^{-/-}$ UUO kidneys displayed exaggerated fibrosis, with 30% additional trichrome-stained area, than wt UUO kidneys (Figure 2b). Fibronectin was also significantly increased in UUO kidneys (Figure 2c). Kidney infiltration by leukocytes and macrophages was significantly increased after UUO in wt and $mg53^{-/-}$ mice, but was further increased in mg53 deficient animals (Figure 2d and 2e).

MG53 modulates NF-κ**B activation and p65 nuclear localization**

The activity of NF-κB (p65) was assessed in the UUO model. The amount of total p65 (t-p65) was significantly increased in UUO kidneys from wt and $mg53^{-/-}$ mice, but NF- κ B activation, assessed as phosphorylation of p65, was greater (16-fold) in the mg53−/− UUO kidney than the wt UUO kidney (2-fold) compared to sham-operated animals (Figure 3a). We analyzed the relative RNA levels of TGF-β1, PAI-1 and Col1a1 in the obstructed kidney.40 We detected low levels of gene expression in sham animals, and similar induction (TGF- β 1~10 fold; PAI-1 ~40 fold; Col1a1 ~40 fold) in UUO kidney from wt and $mg53^{-/-}$ (Figure 3b). To examine the spatial distribution of MG53 in the kidney cortex under normal conditions, renal tissue fractions were analyzed. We regularly obtained 10 fold more total cytosol protein than the total extracted nuclear protein. The t-p65 levels were comparable in the cytosolic fraction between *wt* and $mg53^{-/-}$ mice, but was greater in the nuclear fraction of $mg53^{-/-}$ mice (Figure 3c). MG53 was detected in the cytosol fraction (with tubulin as

cytosol marker). Unexpectedly, a significant amount of MG53 was detected at the nuclear fraction (with Histone H3 as nuclear fraction marker).

To confirm this finding, we also examined MG53 localization in skeletal muscle fractions (Supplementary Figure S2). We reproduced our previous studies that MG53 localized to cytosol and sarcolemmal membrane (Na-K-ATPase as marker of membrane fraction),22, 23, 25 but also detected enriched nuclear MG53 from skeletal muscle.

MG53 interacts with p65 to control TNFα**-induced transcriptional activity**

To confirm our tissue fractionation results, we conducted the following immunohistochemical studies. In cultured wt proximal tubular epithelial cells (PTEC), p65 was found mainly in the cytosol, but localized to nuclei in the absence of MG53 (Figure 4a). When these cells were treated with LPS (5μg) for 8 hours, pro-inflammatory cytokines secretion was significantly greater in $mg53^{-/-}$ cells than wt cells (Figure 4b).

As these data suggested MG53 may interact with p65 to regulate inflammatory cytokine expression, a direct interaction between p65 and MG53 was assessed by co-IP assays using HEK293 cells co-transfected with Flag-p65 and HA-MG53. We demonstrated weak interaction (Figure 4c). Co-transfection of GFP-p65 and RFP-MG53 plasmids and confocal analysis confirmed colocalization of p65 and MG53 (Figure 4d). We then quantified whether MG53 affects NF-κB (p65) transcriptional activity after TNFα stimulation by measuring the luciferase activity from a stable NF-κB/293-GFP-Luc transcriptional reporter cell line that was transiently transfected with either pHM6 vector or HA-MG53. The reporter cells contain a transduced lentivirus expression Firefly Luciferase Reporter driven by a minimal cytomegalovirus (mCMV) promoter in conjunction with four copies of consensus NF-κB transcriptional response elements (κB site) upstream. In the absence of TNFα there was very low intrinsic luciferase activity detected in the reporter cells expressing MG53 proteins. MG53 suppressed TNFα-elicited NF-κB (p65) transcriptional activity by about 40%. (Figure 4e).

Overexpression of MG53 prevented NF-κ**B p65 nuclear translocation upon TNF**α **treatment**

NF-κB p65 translocated to nucleus upon TNFα treatment in kidney PTE⁴¹. To explore whether treatment with MG53 could atteuate NF-κB p65 nuclear translocation upon TNFα stimulation, we overexpressed MG53 in HKC-8 cells either via DOX-dependent Ad-tPA-MG53 viral transduction or through treatment with recombinant human MG53 (rhMG53). Using an MG53-specific mAb, after DOX-dependent induction MG53 was detected by immunoblotting. MG53 was present at a very high level in Ad-tPA-MG53 transduced HKC-8 cells, equivalent to 1 ng rhMG53/μg cellular lysate. In contrast, HKC-8 took up and expressed a much lesser amount of rhMG53 (Figure 5a). Uptake of fluorescent-labeled rhMG53 by HKC-8 was confirmed by confocal microscopy (Supplementary Figure S3). After TNFα treatment time to peak p65 activation was 15 – 30 min (Figure 5b). Under basal conditions, NF-κB p65 resided mainly in the cytosol, and overexpression of MG53 did not change this (Figures 5c and d). However, after TNFα stimulation, p65 translocated to the nuclei in HKC-8 cells not over-expressing MG53. In comparison, nuclear translocation

of p65 fell by 33% (cells given rhMG53) to 50% (transduced cells given DOX) in cells overexpressing MG53 (Figures 5c and d).

Doxycycline-inducible expression and secretion of MG53 from RAW 264.7 cells ameliorates UUO-induced kidney fibrosis in mice

To understand whether treatment with MG53 could attenuate fibrosis, MG53 was increased in the UUO model via engineered RAW 264.7 macrophages (RAW). We engineered RAW via Dox-dependent induction of Ad-tPA-MG53 expression. DOX-dependent induction of tPA-MG53 expression was confirmed by confocal immuostaining of RAW with an MG53 specific mAb (Supplementary Figure S4a), immunoblotting RAW cell lysates after viral transduction (Supplementary Figure S4b), and immunoblotting concentrated culture medium of transduced RAW (Supplementary Figure S4c).

Adoptive transfer of engineered RAW cells into the UUO mouse model was done with 1 \times 10⁶ cells administered through tail vein injection immediately after UUO-surgery. Mice were randomly divided into a group that received normal drinking water (−DOX) and a group that received DOX (2 mg/ml) in drinking water. UUO mice received RAW every other day for a total of 4 injections. Kidneys were harvested 7-days post-surgery.

Mice infused with engineered RAW followed by DOX induction displayed significantly reduced (by 30%) kidney fibrosis compared to those who received no cells, or those infused with engineered RAW cells but without Dox induction (66% fibrotic area) (Figure 6a and 6b). Only a small RAW/Ad-tPA-MG53 population survived the surveillance of reticuloendothelial system by the end of the experiments. IHC staining confirmed that Doxtreatment induced RAW-mediated MG53 expression in the kidney (Figure 6c). Interestingly, UUO kidneys from RAW infused mice and DOX-induced MG53 exhibited a lower p-65/tp65 ratio (by about 40%) than those who either did not receive RAW, or those given RAW without DOX (Figure 6d and 6e). The reduced p-p65 level was confirmed with p-p65 (S536) IHC staining of UUO kidneys (Figure 6f). We think circulating MG53 delivered by Raw/AdtPA-MG53/+DOX and uptake by kidney proximal tubule cells could participate in reducing p-p65 (S536) and decrease UUO kidney inflammation. Compared to UUO-only kidneys, TGF-β1, PAI-1 and Col1a1 expression were increased significantly in UUO kidneys from mice infused with RAW without DOX (no MG53 induction). Elevation of these pro-fibrotic genes could be attributed to induction of a strong allograft reaction from infused cells into immunocompetent mice. However, UUO kidneys from mice infused with RAW and given DOX (MG53 induction) expressed significantly lower $(60 - 72\%$ reduction) PAI-1, and relatively lower levels of TGF-β1 and Col1a1 than mice without Dox treatment (Figure 6g).

Systemic administration of rhMG53 mitigates NF-κ**B activation and fibrosis development in mice subjected to UUO**

We tested whether rhMG53 displayed a similar anti-fibrotic effect as the engineered RAW cell therapy described above. Our prior pharmacokinetic data indicated the short, circulating half-life (about 90 minutes in rodents) of rhMG53.⁴² Mice tolerated chronic administration of 2–6 mg/kg rhMG53^{26, 37} well, so the administration strategy shown in Figure 7a was used. UUO mice were randomly given either vehicle (saline) or rhMG53 (2 mg/kg)

daily for the first 7 days post-surgery, then every other day until they were euthanized on day 12. Sham and CLK kidneys exhibited roughly equivalent t-p65 and fibronectin expression levels, with the p-p65 levels slightly increased in CLK kidneys compared to sham kidneys. After UUO, renal t-p65 increased. Compared to saline-treated animals, the p-65/t-p65 ratio in UUO kidneys of rhMG53-treated animals was reduced by about 37%, while IκBα increased by about 27% (Fig. 7b, c). Additionally, fibronectin fell by about 33% (Fig. 7b, c). IHC confirmed rhMG53 was present in UUO kidneys after treatment (Figure 7d).Total fibrosis was reduced in rhMG53-treated UUO kidneys (Figure 7e). However, RNA expression of TGF-β1, PAI-1 and Col1a1 was similar in mice treated with either saline or rhMG53 (Figure 7f).

DISCUSSION

MG53 is a muscle-enriched TRIM protein initially identified as a prominent component of plasma membrane repair machinery.22, 26, 42, 43 However, the actions of MG53 are not confined to skeletal muscle, and MG53 can be released into the circulation as a myokine.32 We have previously shown MG53 can control inflammation during tissue damage in remote organs.27, 29, 33, 35, 36, 37 The present investigation extends the scope of MG53-mediated organ protection to the kidney. We conclude that MG53 can attenuate intra-renal inflammation and kidney fibrosis by blocking cytokine-induced activation of the pro-inflammatory and pro-fibrotic transcription factor NF-κB. This conclusion is based on the following observations: i) Genetic deletion of MG53 in normal mice resulted in a decline in kidney function as the mice aged that was accompanied by an increase in kidney fibrosis and infiltration of the kidney by leukocytes and macrophages. ii) In a murine UUO kidney injury model, the obstructed kidneys of MG53-deficient mice exhibited more fibrosis and inflammation than MG53-replete mice. iii) TNFα and interleukin-6 production by lipopolysaccharide-treated PTECs was attenuated in the presence of MG53. iv) Treatment with, or over-expression of MG53 in a PTEC cell-line diminished TNFα-induced NF-κB activation by blocking the nuclear translocation of the p65 component of NF-κB. MG53 was found to bind directly to p65, albeit weakly. v) Phosphorylation of p65, a necessary step in activation of NF-κB was decreased in the obstructed kidneys of UUO mice treated with recombinant MG53 or induced to over-express MG53 in macrophages. Furthermore, the inhibitor of NF-κB activation, IκBα, was increased in the obstructed kidneys of UUO mice injected with recombinant MG53.

Genetic deletion of MG53 in normal mice was permissive to the adverse effects of aging on the kidneys. This suggests that the presence of endogenous MG53 provides a baseline level of protection during aging. Similarly, loss of endogenous MG53 made acute kidney injury due to obstruction worse. The UUO kidney accumulated MG53, possibly reflecting remote myokine activity from skeletal muscle. It remains unclear how muscle MG53 may be recruited to the kidney during injury, however a damaged kidney often develops hypoxia and undergoes oxidative stress, $44, 45$ signals that could serve to recruit MG53. If so, this "musclekidney-immune cell" axis could be leveraged to attenuate kidney damage.⁴⁶ Alternatively, kidney cells do express low levels of MG53 at baseline, 26 and during acute or chronic injury MG53 production could be upregulated.

NF-κB is a prominent transcriptional driver of inflammation and fibrosis in kidney injury and aging.^{14, 17, 47, 48} We were prompted to investigate whether the effects of MG53 could be mediated via NF-κB becasue the presence of MG53 in the nuclear compartment of kidney homogenates suggested the potential for transcriptional interference. Nuclear localization of MG53 is not without precedent. In a transgenic model overexpressing myocardial MG53, nuclear MG53 regulated the expression of peroxisome proliferationactivated receptor alpha (PPARα) at the transcriptional level and affected cardiac lipid metabolism.49 Moreover, we recently demonstrated rhMG53 suppressed NF-κB activation to mitigate age-related heart failure through reducing pro-inflammatory cytokines, apoptosis and oxidative stress in aged cardiomyocytes.³⁷

The molecular mechanism of how MG53 modulates p65 nuclear/cytoplasmic mobilization remains unclear. MG53 contains no nuclear localization signal (NLS), but it contains 3 putative N-terminal leucine-rich nuclear export signals (NES), a motif recognized by nuclear export protein exportin (CRM1).⁵⁰ The direct interaction between MG53 and p65 was rather weak as the only or main disruptive effect of MG53 on NF-κB activation. Although MG53 appears to affect the dynamic nucleo-cytoplasmic shuttling of NF-κB critical components, the exact mechanisms remain to be determined.

The critical outcome of MG53-mediated renoprotection appears to be to decrease fibrotic remodeling through attenuation of NF-κB-mediated inflammation. Kidney fibrosis is a complex process involving pathologic deposition of various ECM proteins.51 MG53 seemed to have clear effects on the ECM protein fibronectin. Fibronectin increased in UUO kidneys deficient in MG53 and decreased in UUO mice treated with MG53. The expression of other ECM proteins in the presence and absence of MG53 is more difficult to understand. We previously demonstrated that rhMG53 negatively regulates TGF-β1/Smad signaling and suppresses the synthesis of ECM proteins in an in vitro study.31 In vivo, TGF-β1, PAI-1, and Col1a1 transcripts did not increase in UUO kidneys from $mg53^{-/-}$ deficient animals. These transcripts did decrease in UUO animals induced to over express MG53 through administration of virally-induced mouse macrophages, but when recombinant MG53 was given, no effect was seen. This may be simply a case of sufficient dosing, as the intrarenal MG53 levels achieved were far greater in animals given the macrophage vector as opposed to recombinant MG53. Additionally, protein levels were not measured, and post-transcripitonal events could modify the ECM in UUO. Further research is needed to elucidate the emerging role of MG53 in modulating specific ECM proteins.

In conclusion, this study has demonstrated that a myokine may help regulate baseline inflammation in a remote organ like the kidney, and in the setting of kidney injury may help to attenuate and control inflammation. Figure 8 presents a working model of the role of MG53 in renoprotection. During DAMP (TNFα) and PAMP (LPS) stimulation and ischemic stress, the NF- κ B signalosome is activated in the kidney. MG53 protects the kidney by inhibiting the phosphorylation and activation of NF-κB signaling, stabilizing IκBα, and reducing p65 nuclear mobilization, blocking activation of pro-inflammatory programs. Although endogenous MG53 may mediate these effects, we suggest more robust protection may be afforded using MG53 therapeutically, which is feasible based on the ability of

exogenous MG53 to reduce injury in UUO. Further investigation is warranted to understand how this myokine may be clinically exploited.

Supplementary Material

Refer to Web version on PubMed Central for supplementary material.

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Translational Statement

Chronic inflammation leads to fibrotic remodeling that may also underlie the transition from AKI to CKD. Activation of the pro-inflammatory transcription factor NF-κB is involved in the pathogenesis of kidney inflammation. Here, we provide evidence that MG53, a previously identified cell membrane repair protein, directly interacts with the NF-κB and reduces its transcriptional activity. Concomitantly, exogenously administered MG53 can decrease fibrotic remodeling of the inflamed kidney. These findings point to a protective interplay between MG53 and NF-κB to attenuate the development of inflammation-mediated kidney fibrosis. Pharmacologic administration of MG53 might be a promising approach for the treatment of progressive kidney fibrosis.

Figure 1. The effect of the absence of MG53 on kidney function and histology in aged mice. (a) Compared to young (2-month) mice, Scr concentrations from aged mice (16–20 month) are significantly higher in the absence of MG53. (b) Masson's trichrome stain for renal fibrosis in kidneys from WT and $mg53^{-/-}$ mice of different ages (2-20 month). Scale bars, 50 μm. (c) Quantification of renal fibrosis in aging WT and $mg53^{-/-}$ mice (n=6). Data are presented as mean \pm SD. Data were compared by two-way ANOVA test, followed by Bonferroni's correction for multiple comparison. ns, not significant; * $P < 0.05$, ** $P < 0.01$. (d and e) Immunostaining of kidney sections from WT (left) or $mg53^{-/-}$ (right) (2 or 5 months of age) with anti-CD11b (d) or anti-F4/80 (e). Representative image of cortex and outer medulla are shown. Scale bars, 100 μm. (e). Quantification of renal inflammation in WT and $mg55^{-/-}$ mice (n=4). Values are given as means \pm SD, Student's t-test. ** P<0.01.

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Figure 2. Loss of MG53 exacerbates renal fibrosis and immune cell accumulation after UUO. (a) Kidney MG53 expression increases 7 days post-UUO. Total kidney lysates (50 μg) from Sham or UUO kidneys were examined by Western blotting with a custom made anti-MG53 antibody. rhMG53 (0.3 ng) was loaded in lane 1 as a positive control. (b) Representative pairwise trichrome stains showed Sham (left) and UUO (right) kidney from WT (top) or $mg53^{-/-}$ (bottom) mice and quantified with Masson's Trichrome fibrosis analysis (n=6– 8). Bar = 50 µm. Values are given as means \pm SD, Student's t-test. *** P < 0.001, *** P <0.0001. (c) Western blotting and quantification for fibronectin (FN) from WT and $mg53^{-/-}$ contralateral (CLK) and UUO kidney lysates. Data are presented as mean \pm SD. (d and e) Immunostaining of the CLK or obstructed kidney after UUO in WT or $mg53^{-/-}$ mice with either anti-CD11b (d) or anti-F4/80 (e). Representative image of cortex and outer medulla are shown. Scale bars, 100 μ m. n= 6 *WT* and $mg55^{-/-}$ mice. Data are presented as means \pm S.D, ** $P < 0.01$, *** $P < 0.001$, **** $P \le 0.0001$ (One way ANOVA followed by Bonferroni's test).

Figure 3. NF-κ**B signaling is activated in the obstructed kidney of** *mg53***−/− mice.**

(a) Western blot of phospho (p)-p65 and total (t)-p65 levels from kidney cortex lysates prepared from sham, contralateral (CLK) and UUO kidneys. GAPDH was used as loading control. ($N = 6$ for each group). Analysis of the p-p65 to t-p65 ratio. Values are given as means \pm SD. Student's t-test. ns, not significant; **** P <0.0001 (b) Quantitative RT-PCR showed the relative abundance of TGF-β1, PAI-1 and Col1a1 mRNA in different groups as indicated. Values represent the means \pm SD. ns, not significant; (c) Western blot analysis for t-p65 and MG53 from the cytosol and nuclear fractions (40 μg) of kidney cortex. Histone H3 was used as a nuclear marker and tubulin as a cytosol marker. The relative p65 N/C ratio was calculated with nuclear p65 (normalized to Histone H3) to cytosol p65 (normalized to tubulin) and factored in 10 to represent the 10 fold more of total cytosol protein extracted than that from the total nuclear protein. (n=6 per group) Values are given as means \pm SD, Student's t-test. $*P<0.01$.

Figure 4. MG53 interacts with p65 to control TNFα**-induced transcriptional activity.** (a) Primary proximal tubule epithelial cells (PTEC) were isolated from WT and $mg53^{-/-}$ kidneys and stained for total p65 (t-p65, left panels), nuclei (DAPI, middle panels) and merged images (right panels). Representative images shown from $n = 3$ experiments. Scale bar = 25 μ m. (b) PTEC secretion of TNF α (left) or IL-6 (right) 8 hours post LPS (5 μ g/ml) stimulation. Values are the means \pm SD of 3 independent experiments, One way ANOVA test followed by Bonferroni's test. *** $P < 0.001$, *** $P < 0.0001$. (c) Coimmunoprecpitation assay using HEK293 cells co-transfected with Flag-p65 and HA-MG53 plasmids as indicated. Total cell proteins (TCP, 2%) were loaded as input. Lysates were immune-precipitated with either anti-mouse IgG control, or anti-HA magnetic beads or anti-Flag agarose beads. The immunoprecipitated proteins were loaded on gels and blots were probed with anti-p65 (left) or anti-MG53 (right). (d) HKC-8 cells were co-transfected with GFP-p65 and RFP-MG53 and analyzed with confocal microscope. Scale bar = $10 \mu m$. (e) Quantification of MG53 effects on NF-κB/(p65) transcriptional activity. An NF-κB/(p65) luciferase reporter activity was measured with a bioluminescence luciferase assay under control conditions or after 4 hours, 8 hours of TNFα stimulation. Values are represented means \pm SD from n=5 independent experiments from different days, ** P < 0.01, *** P < 0.001 (Two way ANOVA followed by Bonferroni's test).

Figure 5. MG53 prevents p65 nuclear translocation upon TNFα **treatment.**

(a) Western blot showing MG53 overexpression either by Ad-tPA-MG53 viral transduction (upper panel) or rhMG53 uptake (lower panel) in HKC-8 cells. Various quantities of rhMG53 were loaded as standards. Exposure time was adjusted as needed. (b) time-course of TNFα treatment and expression of p-p65 and t-p65. (c and d) Confocal analysis the effect of HKC-8 cells expressing Ad-tPA-MG53 (c) or uptake rhMG53 (d) on nuclear translocation of NF-κB p65 upon TNFα treatment. Values represent the means ± SD from 3 independent experiments. **** $P < 0.0001$ (One way ANOVA followed by Turkey's multiple comparison test).

Figure 6. MG53 treatment of UUO mice blocks kidney fibrosis.

(a) Representative pairwise trichrome stains showed CLK (left) and UUO (right) kidney from UUO alone (UUO, left, without RAW cells xenotransplantation), −DOX (middle, UUO mice injected with virally transduced RAW cells, no DOX in drinking water), and + DOX (right, UUO mice injected with viral transduced RAW cells, with DOX in drinking water). Scale bars, 5 mm. (b) Quantification of Masson's Trichrome fibrosis analysis from (a). Average of 2 independent experiments, $N = 5-6$ from each group. Data are the means \pm SD. **P < 0.01. (One way ANOVA followed by Bonferroni's test). (c) Immunofluorescence to show kidney expression of MG53 (anti-MG53, green) from xenotransplantation of virally-transduced RAW cells labeled with F4/80 (red fluorescence). Scale bars, 25 μm. (d) Western blot analysis of p-p65 and t-p65 from renal cortical lysates from sham and UUO mice, 7 days post-UUO, with or without RAW (Ad-tPA-MG53) xenotransplantation. Tubulin served as the loading control. (e) Relative p-p65/t-p65 ratios were analyzed. Values represent the means \pm SD from 3 independent experiments. N= 9–10 mice per group. ns, not significant, **P < 0.01; ***P < 0.001 (One way ANOVA followed by Bonferroni's test). (f) Immunohistochemistry of CLK and UUO kidneys stained with p-p65 (S536) antibody. Scale bars = 100μ m. (g) Quantitative RT-PCR showing the relative abundance of TGF- β 1, PAI-1 and Col1a1 mRNA in the indicated groups. Values represent the means \pm SD. ns, not significant; * $P < 0.05$, ** $P < 0.01$. *** $P < 0.001$.

Figure 7. rhMG53 modulates p65 signaling to control renal fibrosis.

(a) rhMG53 administration scheme. (b) Kidney cortex lysates were prepared on day 12 post-surgery from sham or UUO mice that had received either saline or rhMG53 treatment. Lysates were immunoblotted for p-p65, t-p65, t-IκBα and fibronectin (FN). GAPDH was used as a loading control. (c) Quantitative analyses of the blots. All values represent means \pm SD from 3 independent experiments. N= 6–10 mice per group. ns, not significant, *P < 0.05 , *** $P < 0.001$, **** $P < 0.0001$. (One way ANOVA followed by Holm-Sidak's test). (d) Immunofluorescence of anti-MG53 staining (red) on kidney samples from sham (left), UUO + saline (center) or UUO with rhMG53 treatment (right). DAPI (blue) staining. Scale bars, 25 μm. (e) Hematoxylin and eosin and trichrome images and analysis of the kidneys as described above. Scale bar =100 μ m. Values represent the means \pm SD from 3 independent experiments. N= 4 mice per group. *** $P < 0.001$ (One way ANOVA followed by Bonferroni's test). (f) Quantitative RT-PCR showed the relative abundance of TGF-β1, PAI-1 and Col1a1 mRNA in different groups as indicated. Values represent the means \pm SD. ns, not significant.

Figure 8. Schematic illustrating the proposed function of MG53 in the regulation of the NF-κ**B signalosome.**

In response to tissue stress (UUO), DAMP (TNFα) or PAMP (LPS) stimuli, the canonical NF-κB pathway is induced via activation of the IKK complex, which leads to degradation of IκBα and the nuclear translocation of NF-κB. MG53 is proposed to modulate and suppress NF-κB transcriptional activity by directly binding to p65 and preventing its translocation to the nucleus. Additionally, MG53 might stabilize IκBa in the cytosol to prevent NF-κB activation.