Brief report

Type 1 angiotensin receptors on macrophages ameliorate IL-1 receptor–mediated kidney fibrosis

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In a wide array of kidney diseases, type 1 angiotensin (AT1) receptors are present on the immune cells that infiltrate the renal interstitium. Here, we examined the actions of AT1 receptors on macrophages in progressive renal fibrosis and found that macrophage–specific AT1 receptor deficiency exacerbates kidney fibrosis induced by unilateral ureteral obstruction (UUO). Macrophages isolated from obstructed kidneys of mice lacking AT1 receptors solely on macrophages had heightened expression of proinflammatory M1 cytokines, including IL-1. Evaluation of isolated AT1 receptor–deficient macrophages confirmed the propensity of these cells to produce exaggerated levels of M1 cytokines, which led to more severe renal epithelial cell damage via IL-1 receptor activation in coculture compared with WT macrophages. A murine kidney crosstransplantation concomitant with UUO model revealed that augmentation of renal fibrosis instigated by AT1 receptor–deficient macrophages is mediated by IL-1 receptor stimulation in the kidney. This study indicates that a key role of AT1 receptors on macrophages is to protect the kidney from fibrosis by limiting activation of IL-1 receptors in the kidney.

Results and Discussion

AT1A receptor deficiency on macrophages exacerbates kidney fibrosis. To examine infiltration of LysM-expressing macrophages in the kidney during renal fibrosis, we subjected LysM-Cre+ mT/mG reporter mice to UUO. By day 7 of UUO, robust accumulation of GFP+ macrophages was detected in the interstitial areas of obstructed kidneys, but not the contralateral, unobstructed kidneys (Figure 1A), confirming that LysM-Cre–mediated gene excision affects macrophages infiltrating the diseased kidney. To evaluate the actions of AT1 receptors on macrophages that infiltrate the kidney undergoing fibrosis, we harvested kidneys from mice with macrophage–specific deletion of Agtr1a (LysM-Cre+ Agtr1aflox/fox, herein referred to as Macro KO) and WT mice 7 days following UUO. Obstructed kidneys from Macro KO mice contained 64%
more interstitial fibrosis (Figure 1B and Supplemental Figure 5), 80% more collagen-producing myofibroblasts (Supplemental Figure 6), and 80% more collagen I (Col I) protein (Figure 1C) than in WT mice. Moreover, at day 7 after UUO, mRNA expressions of Col I, PAI-1, and TGF-β1 were enhanced by 129%, 47%, and 40%, respectively, in Macro KO kidneys compared with WTs (Figure 1D). Additional experiments further confirmed our results (Supplemental Paragraph 2 and Supplemental Figures 7 and 8). These findings contrast starkly with studies in which global AT1A receptor deficiency ameliorates kidney fibrosis (14), but align with reports of exaggerated tissue damage in AT1A receptor–deficient BM chimeras (15, 16). Collectively, our data indicate that activation of AT1 receptors on infiltrating macrophages protects the kidney from progressive fibrosis.

The heterogeneity of macrophage polarization plays a critical role in the pathogenesis of kidney damage and fibrosis (6). In simplified terms, the balance between proinflammatory M1 and reparative M2 macrophages determines the local composition of cytokine reservoirs that orchestrate disruptions in renal architecture following an insult. Therefore, on days 3 and 7 after UUO, we profiled the renal expression of M1 and M2 markers, including M1 cytokines IL-1β and TNF-α, which have been implicated in the progression of renal fibrosis (Figure 1E). Obstructed Macro KO kidneys had significantly higher expression of M1 markers, including IL-1β, IL-1R1, TNF-α, IL-12p40, and CCL2, compared with obstructed WT kidneys 3 days after UUO, whereas expression of M2 markers Arg-1, FIZZ-1, YM-1, and IL-1R2 was virtually identical between cohorts. A similar pattern was noted at day 7 after UUO (Figure 1E), suggesting that activation of AT1 receptors on macrophages represses M1 cytokine expression during kidney fibrosis.

**AT1A receptors on macrophages limit their expression of M1 cytokines.** We next examined the effects of macrophage AT1 receptor activation on the phenotype of infiltrating macrophages at day 7 after UUO by histologic staining (Supplemental Paragraph 3 and Supplemental Figures 7 and 8). We observed that obstructed Macro KO kidneys had significantly higher expression of M1 markers, including IL-1β, IL-1R1, TNF-α, IL-12p40, and CCL2, compared with obstructed WT kidneys 3 days after UUO, whereas expression of M2 markers Arg-1, FIZZ-1, YM-1, and IL-1R2 was virtually identical between cohorts. A similar pattern was noted at day 7 after UUO (Figure 1E), suggesting that activation of AT1 receptors on macrophages represses M1 cytokine expression during kidney fibrosis.

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array (TLDA) card–based microarray was performed to survey gene expression for inflammatory mediators and signaling molecules in the Ly6C<sup>hi</sup>-activated macrophages (Figure 2A and Supplemental Table 1). Among the upregulated genes, IL-1β but not TNF-α showed enhanced expression in Macro KO kidneys at both days 3 and 7 of UUO (Figure 1E). Moreover, by real-time PCR, Ly6C<sup>hi</sup> and Ly6C<sup>lo</sup> macrophages from the Macro KO kidneys had higher IL-1β expression levels than found in WT controls (Figure 2B). These data indicate that AT1A receptor activation constrains infiltrating macrophage activation by suppressing key inflammatory cytokines including IL-1 during kidney fibrosis.

To explore in vitro the role of the macrophage AT1A receptor in regulating macrophage activation, we cultured Macro KO and WT peritoneal macrophages and subjected them to M1- or M2-polarizing stimuli or vehicle (M0). M1 stimulation increased mRNA expression of M1 markers in both groups, confirming
M1 polarization (Figure 2C). However, M1 macrophages lacking the AT1A receptor showed exaggerated expression levels for each marker. These findings together with M2 polarization studies (Supplemental Paragraph 4 and Supplemental Figures 12 and 13) suggest that AT1A receptor activation during M1 differentiation constrains M1 cytokine production without altering the macrophage’s susceptibility to M2 differentiation.

Enhanced IL-1 production by AT1 receptor–deficient macrophages drives renal cell injury in coculture. Proinflammatory cytokines including IL-1β have been implicated in renal tubular cell (RTC) damage (18). We therefore hypothesized that exaggerated production of inflammatory cytokines by AT1A receptor–deficient macrophages could alter gene expression programs for injury within the kidney tubular cell. Testing this hypothesis in vitro, we measured expres-
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Sion of Col I and kidney injury marker neutrophil gelatinase–associated lipocalin (NGAL) in RTCs following coculture with M0- or M1-conditioned macrophages from our experimental animals (Figure 2D). At 6 hours, M1 macrophages from Macro WT or KO mice, but not M0 macrophages, upregulated Col I and Ngal mRNA expression in RTCs. Thus, cytokines from activated macrophages can mediate damage to kidney tubular cells without direct contact. Moreover, compared with WT M1 macrophages, M1 macrophages lacking AT1A receptors induced an even more profound increase in RTX expression of Col I and NGAL (Figure 2D). To explore whether AT1 receptor–deficient macrophages mediate this exaggerated tubular cell damage through enhanced IL-1 production acting on the renal cell IL-1 receptor (IL-1R), we repeated our coculture experiments with Il1r1-deficient (KO) RTCs (Figure 2D).

In Il1r1 KO RTCs, coinoculation with M1-conditioned macrophages from WT or Macro KO animals induced severely blunted and equivalent increases in Col I and NGAL expression. Finally, recombinant IL-1β induced robust expression of Col I and NGAL in WT but not Il1r1 KO RTCs (Supplemental Figure 14). Thus, activation of the IL-1R on kidney cells by IL-1 secreted from activated macrophages triggers RTC damage. We therefore posited that AT1 receptor activation on infiltrating macrophages limits their IL-1 production, which in turn suppresses gene expression programs for injury in kidney parenchymal cells by preventing renal IL-1R stimulation.

The kidney cross-transplant UUO model. Testing this possibility in vivo requires a system that can separate the contribution of AT1 receptor activation on macrophages from the downstream effects of renal IL-1R activation during UUO-mediated renal fibrosis. We therefore developed a kidney transplantation–ureteral obstruction (KT-UO) model (Supplemental Paragraph 5 and Supplemental Figure 15). To examine the infiltration of macrophages from the kidney recipient into the donor KT-UO kidney, we transplanted a nonfluorescent WT kidney into a LysM-Cre+ mT/mG recipient mouse in which green GFP signals present in the KT-UO kidney represented macrophages from the kidney recipient, and red fluorescent protein (RFP) signals represented other BM-derived cells from the recipient not of the myeloid lineage. Seven days following KT-UO, robust GFP signals in the KT-UO kidney (Supplemental Figure 16) mirrored the levels of fibrosis across the groups. Thus, enhanced activation of the kidney IL-1R accounts for the exaggerated renal fibrosis that accrues from AT1 receptor deficiency on macrophages. As seen earlier in the nontransplanted groups, renal macrophage infiltration in the Macro KO Il1r1 WT group was numerically but not significantly higher than in the WT group (Supplemental Figure 17). Levels of renal macrophage accumulation in the 2 groups lacking IL-1Rs on the KT-UO kidney were similar to each other, but significantly reduced compared with their Macro KO Il1r1 WT counterparts lacking AT1A receptors on macrophages, but expressing IL-1Rs in the KT-UO kidney, and down to levels seen in the Macro WT Il1r1 KOs (Figure 3C). Col I protein content in the KT-UO kidney (Supplemental Figure 16) mirrored the levels of fibrosis across the groups. Thus, enhanced activation of the kidney IL-1R accounts for the exaggerated renal fibrosis that accrues from AT1 receptor deficiency on macrophages. As seen earlier in the nontransplanted groups, renal macrophage infiltration in the Macro KO Il1r1 WT group was numerically but not significantly higher than in the WT group (Supplemental Figure 17). Levels of renal macrophage accumulation in the 2 groups lacking IL-1Rs on the KT-UO kidney were similar to each other, but significantly reduced compared with their Macro KO Il1r1 WT counterparts lacking AT1A receptors on macrophages, but expressing IL-1Rs in the KT-UO kidney, and down to levels seen in the Macro WT Il1r1 KOs (Figure 3C). Col I protein content in the KT-UO kidney (Supplemental Figure 16) mirrored the levels of fibrosis across the groups. Thus, enhanced activation of the kidney IL-1R accounts for the exaggerated renal fibrosis that accrues from AT1 receptor deficiency on macrophages. As seen earlier in the nontransplanted groups, renal macrophage infiltration in the Macro KO Il1r1 WT group was numerically but not significantly higher than in the WT group (Supplemental Figure 17). Levels of renal macrophage accumulation in the 2 groups lacking IL-1Rs on the KT-UO kidney were similar to each other, but significantly reduced compared with their Macro KO Il1r1 WT counterparts lacking AT1A receptors on macrophages, but expressing IL-1Rs in the KT-UO kidney, and down to levels seen in the Macro WT Il1r1 KOs (Figure 3C).

Summary. These experiments establish a mechanism through which AT1 receptors on macrophages function to protect the kidney from progressive fibrosis. Activation of AT1 receptors on infiltrating macrophages suppresses their release of the proinflammatory cytokine IL-1 and thereby prevents stimulation of the kidney IL-1R. We also establish a new KT-UO model that, in conjunction with conditional gene targeting, can serve as a tool for elucidating interactions between hematopoietic and kidney parenchymal cells. The ability of AT1 receptor activation on macrophages to mitigate kidney damage has broad implications for the design of potent therapies to overcome the shortcomings of global angiotensin receptor blockade related not only to previously recognized side effects but also to patently detrimental effects of blocking angiotensin receptors on hematopoietic cells.

Methods

Detailed Supplemental Methods are available online.

Statistics. Data are expressed as mean ± SEM and analyzed by 2-tailed unpaired t test or ANOVA according to statistical methods described in detail in the Supplemental Methods. *P < 0.05 was considered significant.

Study approval. Mice were bred and maintained in the Association for Assessment and Accreditation of Laboratory Animal Care–accredited animal facilities at the Durham Veterans’ Affairs Medical Center (DVAMC) per NIH guidelines. The animal studies were approved by the DVAMC Institutional Animal Care and Use Committee and conducted in accordance with the NIH Guide for the Care and Use of Laboratory Animals.
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