Sisters in arms: myeloid and tubular epithelial cells shape renal innate immunity

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Hato T, El-Achkar TM, Dagher PC. Sisters in arms: myeloid and tubular epithelial cells shape renal innate immunity. *Am J Physiol Renal Physiol* 304: F1243–F1251, 2013. First published March 20, 2013; doi:10.1152/ajprenal.00101.2013.—The importance of innate immunity for survival is underscored by its presence at almost every level of the evolutionary tree of life. The task of "danger" recognition by the innate immune system is carried out by a broad class of pattern recognition receptors. These receptors are expressed in both hematopoietic and nonhematopoietic cells such as renal epithelial cells. Upon activation, pattern recognition and phagocytosis. In this review, we highlight evidence that renal epithelial cells are endowed with such defensive capabilities and as such fully participate in renal innate immune responses.

epithelial cells; innate immunity; pattern recognition receptor;; Toll-like receptor; endotoxin

THE FUNDAMENTAL ROLE OF THE innate immune system is to initiate a quick response immediately after detecting "danger signals" in the setting of infection (nonself) or tissue injury (self). Swiftness is key, because, for example, the doubling time of a single bacterium would allow it to produce millions of progeny within a day if not kept in check. Innate immune cells carry several families of receptors, collectively called pattern recognition receptors (PRRs), which recognize conserved features of pathogens. Some PRRs, such as the mannose receptor and complement system, bind microbes and facilitate phagocytosis. Other PRRs, such as Toll- and NOD-like receptors, induce a wide array of proinflammatory and reactive cytokines in response to danger signals (65).

Most mammalian species have 10-13 types of Toll-like receptors (TLRs) (9). TLRs are responsible for triggering innate immune responses to many forms of pathogens. TLRs are heavily expressed in myeloid cells where they have been extensively investigated. However, some TLRs, such as TLR4, are also expressed and functional in other cell types, including renal epithelial cells (124, 125, 140). Renal tubular TLRs participate in the inflammatory response characteristic of many forms of acute kidney injury. In addition, during injury, tubules can also exhibit phagocytic function (55). Finally, renal epithelial cells express major histocompatibility complex (MHC) class II protein, costimulatory molecules, and produce a plethora of inflammatory and chemotactic cytokines (10). Accordingly, it is increasingly appreciated that epithelial cells and traditional innate immune cells can exhibit remarkable similarities in behavior and function. In this review, we will

primarily focus on the less explored "innate immune cells", i.e., renal epithelial cells.

Epithelial Cells Are Not Alone

The kidney is a complex organ, consisting of at least 12 functionally different epithelial cell types (1). Epithelial cells are surrounded by a dense network of immune cells. In particular, macrophages and dendritic cells, collectively called mononuclear phagocytes, are the most predominant immune cells in the kidney. The visually stunning mononuclear phagocytic network was beautifully described by Soos et al. (129) in 2006. This remarkable network may come as a surprise given the fact that, unlike the gut, the kidney is a nonlymphoid organ and lacks microbial exposure under normal conditions.

It is now increasingly recognized that mononuclear phagocytes have markedly diverse functions: from traditional phagocytic function to versatile, trophic roles (23, 42, 75, 79, 91, 94, 128). Under normal conditions, the mononuclear phagocytic system is believed to play an important role in maintaining the integrity of the tissue microenvironment. In fact, mononuclear phagocytes (CSF1R⁺) are abundantly present even in early embryonic kidneys. Interestingly, when the embryonic kidney was cultured with CSF1 and endotoxin, significant growth in the branch tips and nephrons was observed, presumably because CSF1 and endotoxin induced mononuclear phagocytes to stimulate nephron growth (105).

Like for many other organs, the conventional classification of dendritic cells and macrophages in the kidney remains controversial because of overlap in function and surface markers (37, 54, 96). Multidimensional data such as multicolor flow cytometry and microarray studies continue to reveal the complexity of immune cells in various organs. As detailed in the Immunological Genome Project, it is now evident that each organ has unique sets of immune cell makeup (36, 89). The

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modern classification of immune cell subpopulations now considers subset-specific transcription factors and ontogenies (72, 116). The interested reader is referred to Refs. 15, 38, and 48 (general) and 23, 78, 96, 108, and 132 (kidney) for further details.

Epithelial Cells and PRRs

Much progress has been made in understanding the origins of microbe sensing and innate immune responses. Among diverse PRR families, TLRs are the most extensively studied receptors over the past decade. TLR4 is a functional homomer and requires coreceptors CD14 and MD2 for tighter binding with its ligand, lipopolysaccharide (LPS). TLR2 is heteromeric and complexes with TLRs 1 or 6. Crystallographic analysis showed detailed interactions between the TLR4:MD2 and LPS complex. Similarly, TLR2/TLR1 and its ligand lipopeptide, as well as TLR3 and its ligand poly (I:C), have been crystallized (60, 80, 100). TLRs are strategically located in different cellular compartments, allowing them to sense distinctive pathogen-associated molecular patterns and assemble downstream signaling cascades (62). Some TLRs are exclusively expressed in myeloid cells whereas others are relatively ubiquitous and can be expressed by renal epithelial cells.

Renal expression of TLRs has been studied and confirmed by many investigators. Nevertheless, some uncertainty remains regarding the precise distribution of TLRs in the kidney. This is partly because TLRs are such potent receptors that the expression levels are naturally low at the levels of mRNA and protein. In monocytes, it is estimated that TLR4 is present at 1,300 molecules, whereas CD14 is expressed at 115,000 molecules/cell (135). In nonmyeloid cells, TLR4 expression is likely much lower. Furthermore, due to the inherently complex kidney architecture, one needs to combine technically intricate microdissection, in situ hybridization, and immunostaining to adequately characterize TLRs expression and distribution among various renal cell populations. In that regard, immunostaining remains very challenging because of lack of firm antibodies in this class (41, 71, 123, 140). Nevertheless, collective evidence strongly supports that a number of TLRs are indeed expressed in renal epithelial cells. We and others have previously reviewed the expression of TLRs in the kidney (13, 25, 39, 126).

TLRs in Kidney Injury

Many investigators have reported that tubular expression of TLR2 and TLR4 is increased by experimental ischemia-reperfusion injury in rats and mice (66, 113, 140). For example, TLRs are believed to be activated by various damage-associated molecular patterns, such as the high-morbidity group box 1 (3, 4, 58, 82, 112). Importantly, Wu et al. (141) examined bone marrow chimeric mice between TLR4 knockout and wild-type mice. Chimeric mice lacking renal TLR4 had significantly less tubular damage and azotemia compared with mice lacking hematopoietic TLR4, indicating that intrinsic TLR4 in the kidney is instrumental in mediating tubular damage (141). Pulskens et al. (104) also demonstrated the importance of intrinsic renal TLR4 after ischemic injury. Similarly, Leemans et al. (76) examined bone marrow chimeric mice between TLR2 knockout and wild-type mice and found that intrinsic renal TLR2 has a central role in the unfolding of the injury process.

In models of urinary tract infections, TLR4, TLR5, and TLR11 have been shown to play protective roles (12). When these receptors are defective or absent, clearance of the infection is hindered. It has also been shown that urinary tract epithelial TLR4 and hematopoietic TLR4 are both crucial in mounting a proper inflammatory response to infected bladder mucosa or even pyelonephritis (101, 117). A role for renal TLR4 was also proposed in more chronic models of injury such as obstructive uropathy (103).

In human kidney transplantation, Kruger et al. (69) found that TLR4 expression in proximal and distal tubules is increased in deceased donor kidneys compared with living donor kidneys. Furthermore, the authors determined donor TLR4 genotypes in a cohort of 276 subjects and found 30 subjects with loss-of-function single nucelotide polymorphisms (SNPs), Asp299Gly and Thr399Ile. These two loss-of-function SNPs diminish receptor binding of endotoxin but do not affect TLR4 gene or protein expression (5, 106). Compared with kidneys with wild-type alleles, kidneys with a TLR4 loss-of-function allele had fewer proinflammatory cytokines, and the rate of immediate graft function was higher. It remains to be determined whether this acute protection in TLR4-mutant receivers translates into long-term protection. In summary, compelling evidence indicates that renal epithelial TLRs are central to the regulation of tissue immunity and inflammation.

The Danger Model in the Kidney: S1 Proximal Tubules as the First Line of Defense

To best illustrate the role of renal TLR4 in innate immunity, we next discuss in some detail an animal model of endotoxemia. As opposed to cecal ligation and puncture, ischemia, or transplant models, endotoxin injury models can circumvent concerns such as simultaneous activation of multiple TLRs induced by often uncharacterized damage-associated molecular patterns or polymicrobial infections (20). As such, they are more useful models to characterize specific cellular and molecular pathways of injury.

Endotoxin, released from bacteria in various molecular sizes, can be filtered by nephrons and interact with TLR4 expressed on the proximal tubules. We have recently shown in vivo that systemically administered endotoxin is indeed filtered and taken up by proximal tubules, resulting in tubular oxidative stress (63). Importantly, endotoxin-induced tubular toxicity has an absolute requirement for tubular TLR4. Conversely, TLR4-expressing hematopoietic cells are not essential or sufficient for endotoxin-induced tubular oxidative stress. Note that circulating hematopoietic cells are the primary source of systemic cytokines (11). Taken together, the direct endotoxin-tubular interaction is an important pathway leading to acute kidney injury in endotoxemia.

We also found that filtered endotoxin is internalized predominantly by S1 proximal tubules where TLR4 appears to be expressed the most (63). Two more observations support a role for S1 as the first line of defense in the kidney against endotoxemia. First, S1 endotoxin uptake can be upregulated by endotoxin preexposure, indicating that it is a receptor-mediated process rather than nonspecific endocytosis. Second, and interestingly, this S1-endotoxin interaction does not result in any

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apparent immediate injury to S1 (Fig. 1). This is due in part to the upregulation of cytoprotective molecules such as heme oxygenase-1 and sirtuin-1 in S1 tubules (43, 47, 49, 95). The lack of injury (e.g., oxidative stress) to S1 segments, despite their direct interaction with endotoxin, underscores their high potential for autoprotection. Such a phenomenon has been reported in mononuclear phagocytes after TLR4-mediated exposure to endotoxin (114). Like mononuclear phagocytes, the S1 autoprotection mechanism seems to be dependent, in part, on upregulation of cytoprotective molecules with antioxidant properties. In this model of endotoxemia, the S1 segment acts as the "sensor" of endotoxin in the filtrate and as such autoprotects itself while simultaneously signaling to neighboring segments such as S2 and S3. We note that while S1 autoprotects itself, there is widespread oxidative stress in S2 and S3 (Fig. 1). Whether this represents unavoidable collateral damage or is actually part of a broad signaling cascade is unknown. This function of S1 segments is remarkably similar to that of Kupffer cells in the liver, which also signal the presence of endotoxin to neighboring hepatocytes (110). In summary, S1 segments may play a sentinel role similar to innate immune cells by sensing danger signals and signaling to neighboring cells.

Teleologically, the role of S1 tubules as sentinels for immunity is appealing. The kidney is a highly vascular organ which filters hundreds of liters of blood per day, and a significant part of the filtrate is reabsorbed by proximal tubules. S1 cells, with their upstream location, are strategically poised to screen the filtrate and "watch" for alarm signals coming from the circulation. In particular, it is increasingly appreciated that endotoxemia is a rather common event, occurring daily at subclinical levels during routine breaches to mucosal integrity in various locations (84, 102, 119). An attractive possibility is that, through this pathway, S1 could act as a "sink" for the uptake and degradation of filtered endotoxin. This function is very similar to that proposed for the liver, a major detoxifying center for endotoxemia, particularly that originating from the gut (57, 83). The mechanisms of hepatic endotoxin removal have been reviewed elsewhere (16, 85, 110). These detoxifying functions of the liver, and possibly the kidney, could represent one aspect of the widely recognized ability of many organisms to develop endotoxin tolerance (6).

We also note that TLR4 signaling pathways in the tubules, both in their molecular details and ultimate functions, may not be identical to those present in hematopoietic cells. For example, SIGIRR, a TLR4-inhibitory molecule, while expressed in tubular cells, does not seem to inhibit TLR signaling (73). Feulner et al. (32) reported that murine proximal tubules produce and secrete acyloxyacyl hydrolase into the urinary lumen. Acyloxyacyl hydrolase is an endotoxin-detoxifying enzyme, which could act to minimize the downstream effects of endotoxin that evaded proximal reabsorption. Watts et al.

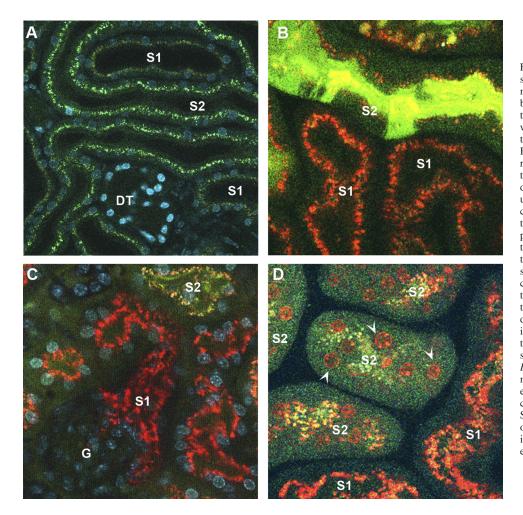


Fig. 1. Endotoxin-induced tubular oxidative stress. A: live 2-photon microscopy of the mouse kidney. S1 and S2 proximal tubules can be discerned by their autofluorescence signatures. S1 exhibits brown autofluorescence, whereas S2 exhibits bright green punctate autofluorescence. Nuclei were stained blue with Hoechst. Distal tubules emit minimal autofluorescence (DT). B: endotoxin-induced oxidative stress was measured with carboxy-2',7'dichlorodihydrofluorescein diacetate (green) using 2-photon intravital imaging. Systemically administered endotoxin (red) was filtered and internalized predominantly by S1 proximal tubules and yet prominent oxidative stress was observed in S2 proximal tubules. Endotoxin uptake observed in S2 is secondary to fluid-phase endocytosis. C: because glomeruli are located at depths beyond the reach of 2-photon microscopy, kidney tissues were freshly dissected to image the deeper segments. This further confirms the internalization of endotoxin by S1 proximal tubules and tubular oxidative stress in downstream S2 and S3 segments. G, glomerulus. D: endotoxin-induced oxidative stress was measured live with dihydroethidium, which emits nuclear fluorescence in the presence of cytosolic superoxide (orange; arrowheads). Similar to B, S2 proximal tubules exhibited oxidative stress whereas S1 tubules exhibited no oxidative stress despite their greater endotoxin uptake.

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(138) demonstrated that endotoxin inhibits HCO_3^- absorption in medullary thick ascending limbs of the loop of Henle (TAL) through a TLR4-dependent pathway. Such downstream effects of endotoxin are likely very varied and will require better characterization in in vivo models.

In this section, we highlighted the roles of renal tubular TLRs in tissue inflammation and immunity. This epithelial cell-centric view could well apply to other organs (33, 74, 87, 127). Polly Matzinger (86), who proposed the danger model, states:

The danger model says that it is a tissue that controls whether you turn on an immune response, by sending alarm signals. It is also a tissue that induces tolerance by allowing its antigens to be presented without alarm signals. Perhaps, therefore, it could also be the tissue that determines the class of immunity.

The Danger Model in the Kidney: Thick Ascending Limbs Regulate Innate Immunity

If S1 proximal tubules are the first line of defense against endotoxemia, TAL, on the other hand, may be essential regulators of the innate immune response within the kidney. TAL tubules span across all areas of the kidney except the inner medulla (28). Consequently, TAL cells are contiguous to most cell types (epithelial and hematopoietic) within the kidney. With this distribution, they are strategically positioned to sense and react to changes (physiological and pathological) in these microenvironments, and possibly mediate various forms of horizontal cross talk (27, 29, 40). Furthermore, TAL in the highly susceptible outer stripe are resistant to acute kidney injury compared with neighboring proximal tubules (18, 27). Therefore, it is plausible to consider that tubules such as S1 and TAL, with an essential modulatory role during activation of innate immunity, must be more resilient to injury.

A unique feature of TAL is the production of Tamm-Horsfall protein (THP; also known as uromodulin). THP is a heavily glycosylated protein that is uniquely produced in the kidney by TAL (29, 107, 120). While predominantly targeted to the apex of the TAL by a GPI anchor signal (107), THP is also released basolaterally by an unknown mechanism (26). Although the functions of THP were elusive for many decades, there has been a recent surge in our understanding of the important role of this glycoprotein in various kidney diseases (29). Interestingly, THP appears to function as an essential effector produced by TAL during kidney injury to modulate innate immunity. In fact, the immunomodulatory functions of THP were a subject of controversy (29). Initially, THP was shown to have anti-inflammatory properties, by suppressing T cells in vitro (93) and binding renal cytokines and lymphokines (IL-1 and TNF) (50). However, a number of subsequent studies also performed in vitro suggested a proinflammatory role of THP, specifically in activating neutrophils (53, 67, 139) and monocytes (130, 144). In addition, Saemann and colleagues (115) demonstrated that THP activates myeloid dendritic cells via TLR4 to acquire a fully mature phenotype. With the availability of THP knockout mice, we provided strong in vivo evidence confirming that the role of THP is indeed antiinflammatory and protective during kidney injury (26, 27, 30). In fact, the presence of THP, produced in TAL, inhibits the production of cytokines and chemokines such as CXCL2 (27) and CCL2 (26) in injured neighboring proximal tubules. Therefore, THP mediates a regulatory cross talk between TAL and proximal tubules, aimed at suppressing tubular activation of innate immunity and promoting recovery (29). This is thought to occur, in part, through basolateral THP released in the interstitium and interacting with the basolateral domain of proximal tubules, where its putative receptor was localized (26, 27). In addition, systemic levels of THP increase during recovery from acute kidney injury (26), suggesting a broader role for THP such as mediating cross talk between the kidney and other organs. Interestingly, recent data also showed that THP regulates the levels of circulating cytokines by acting as a urinary cytokine trap (81). Therefore, through the production of THP, TAL tubules directly modulate innate immunity by regulating tubular epithelial production of cytokines/chemokines and their systemic levels. However, the extent of the interaction of THP with the renal phagocytic system remains uncertain. Dong et al. (21) suggested that THP may be part of the renal antigens presented by dendritic cells after injury caused by LPS injection. Whatever the extent of the interaction between THP and the renal phagocytic system, the outcome, based on the in vivo data from THP knockout mice (27, 30), must be to limit injury and promote repair.

Finally, the role of TAL in renal defense comes full circle through the functions of urinary THP in defense against bacterial colonization of the bladder mucosa. In fact, THP knockout mice are more susceptible to bladder colonization by uropathogenic *Escherichia coli* (7, 90). This occurs because of the binding of THP to *E. coli*, which prevents the interaction of these pathogens with uroplakins on the surface of urothelial cells (99). Therefore, THP serves as a decoy for pathogenic bacteria in the bladder and limits their interaction with cell surface receptors.

In summary, TAL tubules regulate innate immunity by shaping the evolving response to danger signals during kidney injury and by defending the urinary tract from pathogens. This complex task is accomplished through the production of THP, a unique kidney-specific glycoprotein.

Epithelial Cell-Immune Cell Interactions in the Kidney

Many mononuclear phagocyte markers are elevated in the kidney after acute tubular injury and even in chronic diseases such as polycystic kidney disease (44, 92, 147). It was also shown that many of these proteins are upregulated not only in mononuclear phagocytes but also in epithelial cells. Renal epithelial cells secrete chemokines in response to direct stimulation with TLR ligands (133). MHC I and II are highly expressed on proximal tubules after transplant and other stimuli (8, 34, 142). Tubular injury also increases tubular expression of costimulatory molecules (77, 97, 136). There are even some data to suggest that proximal tubules could present antigen to T cells (17, 45, 59, 70, 142).

The generation and activation of mononuclear phagocytes is dependent on CSF1R and its ligand CSF1 (46). Interestingly, CSF1R and its ligand CSF1 are upregulated in the tubules after ischemia-reperfusion injury and transplant (88, 146). Tubular recovery is CSF1R and CSF1 dependent and requires the presence of mononuclear phagocytes (2). Proximal tubules also express GM-CSF (19), a molecule which induces differentiation of monocytes into phagocytes (46). Finally, kidney injury molecule-1 (KIM-1) was shown to be highly expressed in

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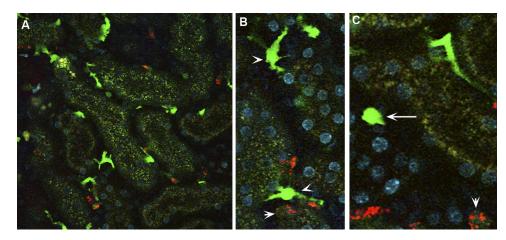


Fig. 2. Mononuclear phagocytic network in the kidney. A-C: CX₃CR1-enhanced green fluorescence protein (EGFP) mouse was injected with rhodamine-labeled (poly I:C) (red), a TLR3 ligand, and the kidney was imaged live. The CX₃CR1-EGFP kidney shows stationary dendritic cells with high EGFP fluorescence (CX₃CR1^{high} dendritic; arrowhead). Mobile spheroidal cells with either high or intermediate EGFP fluorescence are denoted as CX₃CR1^{high} spheroidal (long arrow) and CX₃CR1^{int} spheroidal (short arrow). Poly (I:C) localized to CX3CR1^{int} spherical cells. CX₃CR1^{high} dendritic cells and CX₃CR1^{high} spheroidal cells did not take up poly (I:C). See also Supplemental Movie (supplementary material for this article is accessible on the journal website).

injured proximal tubules. Ichimura et al. (55) demonstrated that KIM-1 is in fact a phosphatidylserine receptor and as such can function as a scavenger receptor. Therefore, during tubular injury, proximal tubules are transformed into "semiprofessional phagocytes." Of note, KIM-1 can be upregulated anywhere from S1 to S3 proximal tubules, depending on the type of injury (145).

We have reviewed similarities between epithelial cells and innate immune cells. However, one important difference remains between the two cell types: mobility. Renal epithelial cells do not typically translocate. Therefore, epithelial cells alone will not be able to accomplish higher levels of immune activities (such as remote information transfer) unless they are supported by immune cells (56, 122). Ultimately, epithelial cells and immune cells are both essential in shaping renal immunity.

Figure 2 and a Supplemental Movie show CX_3CR1^+ myeloid cells in the live kidney (all supplementary material for this article are accessible on the journal website). The chemokine receptor CX_3CR1 is widely expressed in mononuclear phagocytes, and CX_3CR1 has been central to define its lineage and subsets (121, 143). The CX_3CR1^+ renal mononuclear phagocytes are remarkably heterogeneous in shape, signal intensity, and motion, likely reflecting their functional diversity.

Traditionally, immune cells are thought to exacerbate tubular injury through inflammatory cytokines. However, it is increasingly recognized that certain subsets of immune cells play protective roles via immune cell-epithelial cell interactions. Many groups have reported intriguing epithelial cellimmune cell cross talk. Lee et al. (75) demonstrated that M1 macrophages switch to a M2 phenotype when cocultured with proximal tubular cells. Wang et al. (137) showed that proximal tubules stimulated by endotoxin inhibit macrophage activation. Others also showed that proximal tubules modulate mononuclear phagocyte function, maturation, and differentiation (64, 68). The interactions between renal epithelial cells and mononuclear phagocytic cells are not restricted to proximal tubules. Collecting duct epithelial cells also influence macrophage phenotypes (35). To investigate the role of myeloid cells in vivo, clodronate and CD11b- or CD11c-diphtheria toxin transgenic mice are often used. Although the outcomes may vary depending on the timing of depletion and models used (22, 24, 51, 52, 61, 75, 118), beneficial roles of mononuclear phagocytes have been demonstrated by multiple groups (98). In a model of cisplatin nephrotoxicity, CD11c depletion of diphtheria toxin transgenic mice resulted in more severe injury, suggesting that renal CD11c⁺ mononuclear phagocytes mediate protection in this model (131). Recently, Ferenbach et al. (31) showed that clodronate does not deplete alternative M2 macrophages and gives rise to less severe renal ischemia-reperfusion injury. Taken together, compelling evidence indicates that reciprocal interactions between mononuclear phagocytes and renal epithelial cells are instrumental in maintaining the integrity of the tissue environment. In other organs, even stronger evidence exists that epithelial-immune cell interactions shape overall organ immunity (14, 109, 127, 134).

Concluding Remarks

The kidney is a nonlymphoid, sterile organ, and yet renal epithelial cells are surrounded by an extensive mononuclear phagocytic network. These mononuclear phagocytes are pivotal in maintaining the tissue environment in health and disease. There exists considerable cross talk between mononuclear phagocytes and epithelial cells. In fact, renal epithelial cells share many phenotypic and functional characteristics with mononuclear phagocytes. Because renal epithelial cells are positioned at the interface between the internal milieu and external environment, it comes as no surprise that they can serve as primary guardians of the kidney and the body as a whole. As an example, we featured renal epithelial TLR4, which is strategically located on the tubules so it can respond to both systemic infection and local injury. Although innate immune cells activated by injured renal epithelial cells are commonly viewed as amplifiers of injury, protective roles of innate immune cells are increasingly appreciated. Investigations of immunity at the whole organ level will likely reveal more facets to the functions of renal epithelial and myeloid cells. Some of these functions will be unique to one cell type but others are likely shared by these sisters in arms.

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DISCLOSURES

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AUTHOR CONTRIBUTIONS

Author contributions: T.H. and P.C.D. provided conception and design of research: T.H. performed experiments: T.H. and P.C.D. analyzed data: T.H. and P.C.D. interpreted results of experiments; T.H. prepared figures; T.H. drafted manuscript; T.H., T.M.E.-A., and P.C.D. edited and revised manuscript; T.H. and P.C.D. approved final version of manuscript.

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