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Calcium oxalate crystals induce renal inflammation by NLRP3-mediated IL-1β secretion

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Nephrocalcinosis, acute calcium oxalate (CaOx) nephropathy, and renal stone disease can lead to inflammation and subsequent renal failure, but the underlying pathological mechanisms remain elusive. Other crystalllopathies, such as gout, atherosclerosis, and asbestosis, trigger inflammation and tissue remodeling by inducing IL-1β secretion, leading us to hypothesize that CaOx crystals may induce inflammation in a similar manner. In mice, intrarenal CaOx deposition induced tubular damage, cytokine expression, neutrophil recruitment, and renal failure. We found that CaOx crystals activated murine renal DCs to secrete IL-1β through a pathway that included NLRP3, ASC, and caspase-1. Despite a similar amount of crystal deposits, intrarenal inflammation, tubular damage, and renal dysfunction were abrogated in mice deficient in MyD88; NLRP3, ASC, and caspase-1; IL-1R; or IL-18. Nephropathy was attenuated by DC depletion, ATP depletion, or therapeutic IL-1β antagonism. These data demonstrated that CaOx crystals trigger IL-1β-dependent innate immunity via the NLRP3/ASC/caspase-1 axis in intrarenal mononuclear phagocytes and directly damage tubular cells, leading to the release of the NLRP3 agonist ATP. Furthermore, these results suggest that IL-1β blockade may prevent renal damage in nephrocalcinosis.

Introduction

Kidney stone disease affects about 12% of men and 5% of women during their lifetimes in the United States (1). Unlike symptomatic urolithiasis, intrarenal nephrocalcinosis is often asymptomatic, but can lead to significant kidney injury and renal failure (1–3). Calcium oxalate (CaOx) stones account for the vast majority of calculi in primary and secondary forms of hyperoxaluria (4, 5). In addition, acute oxalosis (e.g., in ethylene glycol poisoning) can induce acute kidney injury as a result of widespread intrarenal CaOx deposition (6). Histopathology of human CaOx nephropathy as well as microarray analysis of rodent CaOx nephropathy document a substantial contribution of interstitial inflammation to tissue remodeling, but the molecular and cellular mechanisms of CaOx-induced renal inflammation remain largely unknown (7–11).

Inflammation and tissue remodeling in various crystalllopathies, such as gout (caused by monosodium urate crystals), pseudogout (caused by calcium pyrophosphate crystals; ref. 12), atherosclerosis (caused by cholesterol crystals; ref. 13) silicosis (caused by silica crystals; ref. 14) and asbestosis (caused by asbestos; ref. 14), involve a unique intracellular signaling pathway for the secretion of the proinflammatory cytokine IL-1β: the NLRP3 inflammasome/apoptosis-associated speck-like protein containing CARD/caspase-1 (NLRP3/ASC/caspase-1) axis (15). The NLRP3 inflammasome is a large cytoplasmatic protein complex able to integrate multiple extracellular and intracellular danger signals into ASC binding and activation of caspase-1 (also known as pro–IL-1β-convertase enzyme; ref. 15). In contrast to other NF-kB-dependent cytokines that are readily secreted, the inactive pro-form of IL-1β requires cleavage (e.g., by caspase-1) as an additional enzymatic step before the bioactive form of IL-1β can be secreted from the cell (15).

Here, we hypothesized that CaOx crystals have a similar agonistic potential on NLRP3 and trigger IL-1β secretion and that this process contributes to CaOx-induced tissue inflammation, for example, in nephrocalcinosis and/or CaOx nephropathy.

Results

CaOx crystals need NLRP3/ASC/caspase-1 to induce IL-1β in DCs. We hypothesized that CaOx crystals have the same potential to activate DCs as do crystals of uric acid, calcium pyrophosphate, or cholesterol (12, 13). CaOx crystals were of small size when used in vitro together with bone marrow–derived DCs (BMDCs; Supplemental Figure 1; supplemental material available online with this article; doi:10.1172/JCI63679DS1). CaOx crystal exposure activated murine BMDCs to release IL-1β in a dose-dependent manner, but only when cells were prestimulated with LPS, TNF, IL-1, or histones (Figure 1A and Supplemental Figure 2). Western blotting of DC supernatants revealed that LPS prestimulation was a prerequisite for induction of pro–IL-1β expression (Figure 1A). LPS also induced pro–caspase-1 in BMDCs, and CaOx crystals induced caspase-1 activation similar to that by the NLRP3 agonist ATP (Figure 1A). To address a potential NLRP3 agonistic effect of CaOx crystals, BMDCs from Nlrp3−/−, Asc−/−, and Casp1−/− mice were also prepared; LPS/CaOx crystals were unable to induce IL-1β secretion in these cells (Figure 1, B and C). Although LPS induced pro–IL-1β and pro–caspase-1 expression, CaOx crystals

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were unable to induce caspase-1 activation in Nlrp3–/–, Asc–/–, and Casp1–/– BMDCs, as was the case with LPS/ATP stimulation (Figure 1, B and C). Additionally, the caspase inhibitor Z-VAD-FMK had the same effect on LPS/CaOx crystal–induced IL-1β secretion in BMDCs (Figure 1D). Thus, CaOx crystals induced IL-1β secretion in DCs via the NLRP3/ASC/caspase-1 axis.

CaOx crystal–induced NLRP3 activation depends on phagocytosis and potassium efflux. Whether agonists directly interact with NLRP3 remains unknown, but various mechanisms are involved in NLRP3 activation by extracellular danger signals: particle uptake into the cell by phagocytosis, lysosomal cathepsin leakage, potassium efflux from the cell, P2X7 channel activation, and intracellular oxidative stress (15). We therefore performed a series of experiments to determine which of the aforementioned factors contributes to CaOx crystal–induced NLRP3 activation. Cytochalasin D, a phagocytosis-blocking agent that inhibited actin polymerization and crystal uptake into intracellular endosomes (Figure 2A), significantly reduced LPS/CaOx crystal–induced IL-1β secretion in BMDCs, whereas the cathepsin inhibitor CA074Me and the antioxidant N-acetyl cysteine had no effect (Figure 2B). Furthermore, blocking potassium efflux by a high concentration of potassium in the extracellular fluid had the same effect (Figure 2C), which suggests that potassium efflux is required for CaOx-induced NLRP3 activation. Comparison of LPS/CaOx crystal–induced IL-1β secretion between WT and P2x7–/– BMDCs did not support a contribution of P2X7 in this context (Figure 2D). Thus, CaOx crystals induced NLRP3-mediated IL-1β secretion in DCs via phagocytic uptake of CaOx crystals into intracellular endosomes as well as via inducing potassium efflux from the cell, a process known to activate NLRP3 (16).

CaOx crystals induce IL-1β secretion in renal DCs, but not in renal parenchymal cells. To determine whether CaOx crystals have a similar potential to activate renal DCs or renal parenchymal cells, we stimulated primary renal CD45+CD11c+ DCs, ECs, mesangial cells (MCs), and tubular epithelial cells (TECs) with LPS/CaOx crystals as above. LPS/CaOx crystals induced IL-1β secretion in renal DCs in a manner similar to that of LPS/ATP, whereas none of the renal parenchymal cell types responded to the stimulation (Figure 2E). Western blotting confirmed this finding and also showed that LPS stimulation only induced pro–IL-1β expression in renal DCs (Figure 2F). Hence, CaOx crystals induced secretion of mature IL-1β only in renal DCs, not in renal parenchymal cells.

CaOx crystals induce ATP release from dying TECs. Consistent with previous reports (10), CaOx crystal exposure killed TECs (as demonstrated by lactate dehydrogenase [LDH] release) in a dose-dependent manner, independent of Nlrp3 genotype (Supplemental Figure 3). Dying cells release intracellular damage-associated molecular pattern molecules (DAMPs) that have the potential to activate pattern recognition receptors of other cells (17). ATP is one such intracellular DAMP, reported to drive sterile inflamma-
Figure 2
Mechanisms of CaOx crystal–induced IL-1β secretion. (A) BMDCs were primed with LPS (1 μg/ml) and stimulated with CaOx crystals in the presence or absence of cytochalasin D. TEM showed that CaOx crystals appeared outside the cell (asterisks) and inside intracellular endosomes (arrowheads), whereas cytochalasin D treatment inhibited their uptake in DCs. (B) Incubation with cytochalasin D, CA074Me, and N-acetyl cysteine (NAC) affected IL-1β secretion, measured by ELISA in cell culture supernatants after 6 hours. (C) Similar experiments were performed using 75 mM KCl in cell culture medium to block potassium efflux from the DCs. NaCl of an identical molarity was used as a molarity control. High extracellular potassium prevented CaOx-induced IL-1β secretion. (D) DCs were isolated from WT and P2x7−/− mice. ATP-induced, but not CaOx-induced, IL-1β secretion depended on the presence of P2X7. Data are means ± SD from 3 independent experiments. (E) Western blotting. LPS priming induced pro–IL-1β expression only in DCs. (F) Primary TECs were isolated from WT mice and cultured with increasing concentrations of CaOx. ATP release was quantified in cell culture supernatant upon 18 hours of CaOx crystal exposure with and without increasing concentrations of apyrase. (G) Coculture of LPS-primed DCs with CaOx (100 μg/ml) damaged TECs with or without apyrase (1 U/ml), IL-1β secretion was measured by ELISA after 24 hours. Data are means ± SD from 3 independent experiments. *P < 0.05, **P < 0.01, ***P < 0.001 vs. medium (B, C, and E); vs. WT (D); vs. 1,000 μg/ml CaOx alone (G); or as indicated by brackets.

Figure 2
Mechanisms of CaOx crystal–induced IL-1β secretion. (A) BMDCs were primed with LPS (1 μg/ml) and stimulated with CaOx crystals in the presence or absence of cytochalasin D. TEM showed that CaOx crystals appeared outside the cell (asterisks) and inside intracellular endosomes (arrowheads), whereas cytochalasin D treatment inhibited their uptake in DCs. (B) Incubation with cytochalasin D, CA074Me, and N-acetyl cysteine (NAC) affected IL-1β secretion, measured by ELISA in cell culture supernatants after 6 hours. (C) Similar experiments were performed using 75 mM KCl in cell culture medium to block potassium efflux from the DCs. NaCl of an identical molarity was used as a molarity control. High extracellular potassium prevented CaOx-induced IL-1β secretion. (D) DCs were isolated from WT and P2x7−/− mice. ATP-induced, but not CaOx-induced, IL-1β secretion depended on the presence of P2X7. Data are means ± SD from 3 independent experiments. (E) Western blotting. LPS priming induced pro–IL-1β expression only in DCs. (F) Primary TECs were isolated from WT mice and cultured with increasing concentrations of CaOx. ATP release was quantified in cell culture supernatant upon 18 hours of CaOx crystal exposure with and without increasing concentrations of apyrase. (G) Coculture of LPS-primed DCs with CaOx (100 μg/ml) damaged TECs with or without apyrase (1 U/ml), IL-1β secretion was measured by ELISA after 24 hours. Data are means ± SD from 3 independent experiments. *P < 0.05, **P < 0.01, ***P < 0.001 vs. medium (B, C, and E); vs. WT (D); vs. 1,000 μg/ml CaOx alone (G); or as indicated by brackets.
CaOx nephropathy in C57BL/6 mice. (A) Kidney sections showed CaOx crystals distributed throughout, especially in the tubular lumina (Pizzolato stain; left) or viewed under polarized light (right). Original magnification, ×100 (left); ×200 (right); ×400 (inset). (B) At higher magnification, Pizzolato stain showed smaller (granular) crystals inside TECs as well as in the renal interstitial compartment. Original magnification, ×400 (left), ×1,000 (right). (C) TEM (left) confirmed large CaOx crystal aggregates within the tubular lumen (black arrow) as well as in the interstitial space (white arrow). Freeze-fracture EM (right) identified CaOx crystals in the tubular lumen (black arrow) as well as inside cytoplasmic organelles, some of which appeared to be lysosomes (white arrows). Original magnification, ×3,000. (D) TEM also showed CaOx crystals within the interstitial compartment in close proximity to interstitial cells (left; black arrows). At higher magnification (right), an interstitial cell (white asterisk indicates nucleus) engulfed a CaOx crystal (black arrows denote outer cell membrane); neighboring TECs are denoted by black asterisks. Original magnification, ×7,000 (left); ×25,000 (right).

Oxalate nephropathy involves diffuse crystal deposition, renal inflammation, and failure. Given the potential of CaOx crystals to activate renal DCs via NLRP3/ASC/caspase-1 in both direct and indirect ways, we suspected they make a functional contribution to oxalate nephropathy. Direct DC activation would require CaOx crystals to also reach the renal interstitial compartment, where DCs and other antigen-presenting cells span a dense network around the tubules (20). Murine oxalate nephropathy was characterized by diffuse CaOx crystal deposition throughout the kidney, with focal crystal casts obstructing proximal and distal tubules within the outer medulla as well as within the cortex, which was absent in control mice (Figure 3A and Supplemental Figure 4). Single crystals or small clumps of crystals, similar in appearance to the ones we applied for cell culture (Supplemental Figure 1), not only adhered to the luminal membranes of TECs, but were also found intracellularly within endosomes (Figure 3, B and C, and Supplemental Figure 5A). Diffuse CaOx crystal deposits were also found in the renal interstitial compartment, in close proximity to and within renal interstitial cells (Figure 3, B and D, and Supplemental Figure 5B). In fact, transmission EM (TEM) imaging showed that interstitial cells occasionally engulfed and took up crystals (Figure 3D). Renal CaOx crystal deposition was associated with diffuse neutrophil infiltrates and tubular necrosis mainly at the inner stripe of the outer medulla, as demonstrated by the disintegration of TECs and granular casts in tubular lumen (Figure 4, A and B). The structural alterations of oxalate nephropathy were associated with renal failure, as shown by a transient increase in plasma creatinine and BUN levels, peaking at 24 hours, compared with those in vehicle-injected mice (Figure 4A and Supplemental Figure 6). Intra-renal mRNA expression of the kidney injury markers KIM-1 and L-FABP, as well as that of the proinflammatory cytokines IL-6 and TNF-α, was increased (Figure 4C). Together, these observations indicated that murine oxalate nephropathy was characterized by diffuse CaOx crystal formation inside the tubular and the interstitial compartments, which is associated with tubular injury, renal inflammation, and failure.

Oxalate nephropathy depends on MyD88, IL-1R, and IL-18. Cytokine expression and neutrophil infiltration are suggestive of the involvement of innate immunity in oxalate nephropathy. To test this concept, we induced oxalate nephropathy in Myd88−/− mice, since Myd88 is a central signaling adaptor that translates innate pattern recognition into NF-κB activation and expression of proinflammatory cytokines, including pro-IL-1B and pro-IL-18 (21, 22). Unlike WT mice, Myd88−/− mice were almost entirely protected from acute kidney injury and failure, despite harboring identical CaOx crystal depositions (Figure 5). Il1r−/− mice were almost entirely protected, whereas Il18−/− mice displayed only partial protection (Figure 5). Thus, oxalate nephropathy depended on Myd88-mediated innate immunity, which also involved IL-1R and IL-18.

Oxalate nephropathy is abrogated in Nlrp3−/−, Asc−/−, and Casp1−/− mice. To test the functional contribution of the NLRP3/ASC/caspase-1 axis, we induced oxalate nephropathy in Nlrp3−/−, Asc−/−, or Casp1−/−
mice. Despite renal CaOx deposits identical to those of WT controls, all 3 groups were partially protected from kidney injury and failure, as evidenced by significant reductions in plasma creatinine, BUN, tubular damage, mRNA expression of tubular injury markers KIM-1 and L-FABP, and interstitial neutrophil infiltrates (Figure 6 and Supplemental Figure 7). Hence, NLRP3, ASC, and caspase-1 contributed to renal inflammation, damage, and dysfunction in CaOx nephropathy.

CaOx crystal–induced acute kidney injury depends on intrarenal mononuclear phagocytes. The results of our in vitro studies suggested that CaOx crystal–induced IL-1β secretion may selectively originate from the intrarenal network of mononuclear phagocytes, such as DCs and macrophages. Flow cytometry of renal cell suspension obtained after 24 hours of CaOx exposure revealed activation of CD45+CD11c+ intrarenal myeloid DCs, as evidenced by increased surface expression of CD40 and CD86 as well as by protrusions and dendrite expansion, visualized by scanning EM (Figure 7, A and B, and Supplemental Figure 8A). This was associated with partial protection from acute kidney injury compared with control liposome–treated mice (despite identical amounts of CaOx deposits), as documented by a significant reduction of plasma creatinine level, BUN level, and tubular damage (Figure 7C). Similar results were obtained when cells were selectively depleted by diphtheria toxin in CD11c+ DTRg mice (Figure 7D and Supplemental Figure 8B). Renal neutrophil counts were determined by Ly6G B1.2 immunostaining and quantified per 15 hpf. Original magnification, ×100. (C) Renal mRNA expression was determined by real-time RT-PCR. Data are means ± SEM from 5–6 mice per group. *P < 0.05, ***P < 0.001.

Degradation of extracellular ATP with apyrase protects mice from oxalate nephropathy. Our in vitro studies showed that dying TECs released ATP, an endogenous NLRP3 agonist that triggers IL-1β secretion in DCs (19). Apyrase injection degrades extracellular ATP and can block the NLRP3-agonistic effect of this endogenous danger signal during sterile inflammation (18). We therefore used apyrase injections to test whether extracellular ATP also contributes to renal inflammation and injury during oxalate nephropathy. A single intraperitoneal injection of 20 U apyrase partially reduced tubular injury and neutrophil recruitment during oxalate nephropathy and improved renal excretory function, as documented by a significant reduction of plasma creatinine and BUN levels compared with vehicle-treated control mice (Figure 8A). Systemic IL-6 and MCP-1 levels were also reduced (Supplemental Figure 9A). We concluded that extracellular ATP (e.g., as it is released from dying TECs) served as an additional trigger for renal inflammation, injury, and failure in oxalate nephropathy.

The recombinant IL-1R antagonist anakinra prevents oxalate nephropathy. All of the above observations suggested a functional role of NLRP3/ASC/caspase-1–mediated IL-1β secretion as a previously unrecognized pathomechanism for inflammation and tissue damage in CaOx nephropathy. Thus, therapeutic blockade of IL-1β may be able to interfere with this pathomechanism and protect against renal failure. Anakinra, a recombinant version of the endogenous IL-1 receptor antagonist, is approved by the FDA for the treatment of rheumatoid arthritis (23, 24). Intraperitoneal injection of anakinra dose-dependently reduced tubular injury and neutrophil recruitment and improved renal excretory function during oxalate nephropathy (Figure 8B and Supplemental Figure 9B). We concluded that IL-1β-mediated inflammation and tissue damage in kidney injury induced by CaOx crystals, and thus IL-1β blockade protected from renal failure in oxalate nephropathy.
Discussion
Renal stone disease is widely considered to be a mechanical problem causing intra- or extrarenal urinary outflow obstruction, but crystal nephropathies are also associated with significant intrarenal inflammation that can cause acute kidney injury (3, 11). We hypothesized that the NLRP3 inflammasome — an intracellular pattern recognition platform that translates danger signaling into secretion of mature IL-1β — is involved in this process, because immune recognition of numerous crystals involves the NLRP3/ASC/caspase-1 axis (12–14).

In contrast to widely expressed Toll-like receptors or Rig-like helicases, the NLRP3/ASC/caspase-1 axis is restricted to antigen-presenting cells inside the kidney (25–27). We used clodronate liposome in WT mice or diphtheria toxin in CD11c DTRg mice to demonstrate that CaOx-induced intrarenal IL-1β secretion originated from the intrarenal network of interstitial mononuclear phagocytes, which includes several DC populations (28). CaOx crystals also formed in the interstitial compartment, where they were internalized by interstitial cells, a process demonstrated by us and others using TEM (present study and ref. 11). Our in vitro studies confirmed that phagocytic uptake was required for CaOx-induced IL-1β secretion in DCs, an observation consistent with the mode of NLRP3 activation by monosodium urate and calcium pyrophosphate crystals (12), silica (14), asbestos (14), and cholesterol crystals (13) in other disease contexts. Potassium efflux from the cell, occurring upon cell membrane disruption or pore formation by bacterial toxins, is another trigger of NLRP3 activation in DCs (15, 29). CaOx crystal–induced NLRP3 activation also depended on this mechanism, which was again consistent with previous reports on crystals of asbestos, monosodium urate (14), silica (30), cholesterol (31), and calcium phosphate (32). Intrarenal DCs and macrophages are important regulators of innate immunity and tissue remodeling in the kidney (28, 33). They express all known pattern recognition platforms that translate infectious and noninfectious dangers into local expression of proinflammatory cytokines and chemokines, causing subsequent immune-mediated pathology in the kidney (28, 33, 34). In addition, renal DCs express numerous negative regulators of innate immunity that limit overshooting renal inflammation and inappropriate kidney injury, e.g., in postischemic tubular necrosis (35–37). The present study is the first to our knowledge to demonstrate that acute kidney injury in crystal nephropathy largely depends on the capacity of intrarenal mononuclear phagocytes to trigger renal inflammation upon crystal recognition.

TEM and freeze-fracture EM demonstrated crystal uptake into intracellular compartments of TECs in vivo. TECs express NLRP3 but lack pro–IL-1β induction upon NF-κB priming (26, 38, 39). Therefore, CaOx crystals did not trigger mature IL-1β secretion in TECs. However, TECs internalize CaOx crystals, inducing oxidative stress and tubular cell necrosis (40–42). We confirmed this finding and showed that necrotic tubular cells released ATP, a potent NLRP3 agonist (19). Iyer et al. recently demonstrated that ATP is released upon acute tubular necrosis, which triggers NLRP3 activation and renal inflammation, a process that aggravates postischemic acute kidney injury (43). These results were supported by our present in vitro studies. In addition, the protective effects of apyrase treatment on CaOx nephropathy supported the functional contribution of ATP to renal inflammation and damage. It is noteworthy that apyrase injections also reduced plasma levels of IL-6 and MCP-1, implicating some systemic effects, which could be either causative or secondary for the improvement of the renal phenotype.
IL-1 blockade with appropriate antagonists is an established treatment of IL-1–dependent diseases, such as rheumatoid arthritis, Still’s disease, and rare hereditary autoinflammatory disorders (24). Recently, IL-1 blockade also proved effective in crystal-mediated diseases, such as gout and pseudogout (44, 45). The dramatic therapeutic effects seem to directly relate to the fact that NLRP3 translates crystal recognition into IL-1β secretion as a central element of crystal-induced tissue inflammation and pathology (24). Our data demonstrated a similar therapeutic effect of the recombinant IL-1 receptor antagonist anakinra in CaOx nephropathy in mice, which suggests that IL-1 blockade may also reduce renal inflammation and kidney damage in other crystal nephropathies.

In conclusion, CaOx crystals triggered IL-1β–dependent innate immunity via the NLRP3/ASC/caspase-1 axis in DCs. Thus, intrarenal antigen-presenting cells promoted IL-1–dependent inflammation and kidney injury in renal stone disease. In addition, CaOx crystals indirectly activated NLRP3 by damaging TECs and releasing the NLRP3 agonist ATP. IL-1β blockade with anakinra prevented CaOx kidney injury disease, which could be a novel therapeutic approach to limit kidney damage in crystal nephropathies.

Methods

Animal studies

C57BL/6, NOD/ShiLt, NOD/Casp1–/–, Il1r1–/–, Il18–/–, and Myd88–/– mice were procured from Jackson Laboratories. Nlrp3–/– and Asc–/– mice were provided by J. Tschopp (University of Lausanne, Lausanne, Switzerland) and by V. Dixit (Genentech, San Francisco, California, USA). CD11c DTRg mice were provided by C. Kurts (University of Bonn, Bonn, Germany) and D. Engel (University of Bonn, Bonn, Germany). Mice were housed in groups of 5 in filter top cages with unlimited access to food and water. Cages, food, and water were sterilized by autoclaving before use. 6- to 8-week-old mice received a single intraperitoneal injection of 100 mg/kg sodium oxalate and 3% sodium oxalate in drinking water, and kidneys were harvested after 24 hours. DC depletion was achieved by 2 intravenous injections of liposomal clodronate (200 μl) at 48 and 24 hours before injection of sodium oxalate. Control liposomes were used as vehicle control. In another experiment, CD11c DTRg mice were given a single intravenous injection of 8 ng diphtheria toxin (Sigma-Aldrich) 6 hours before sacrifice. Urine samples were collected after 24 hours, before sacrifice by cervical dislocation and harvest of plasma and kidney tissue (46). Kidneys were kept at –80°C for protein isolation and in RNA later solution at –20°C for RNA isolation. One part of the kidney was also kept in formalin to be embedded in paraffin for histological analysis.

Assessment of renal injury

Kidney sections (2 μm) were stained with periodic acid–Schiff (PAS) reagent. Tubular injury was scored by assessing the percentage of necrotic tubules and presence of tubular casts. Ly6B.2+ neutrophils were identified by immunostaining (clone 7/4; Serotec). Pizzolato staining visualized CaOx crystals, and crystal deposit formation in the kidney was evaluated as described previously (47), using the following point values: 0, no deposits; 1, crystals in papillary tip; 2, crystals in cortical medullary junction; 3, crystals in cortex. When crystals were observed in multiple areas, points were combined. Plasma creatinine and BUN were measured using Urea FS kit (DiaSys Diagnostic Systems) according to the manufacturer’s protocol. Both parameters were corrected for body weight. Neutrophil infiltrates were counted in 15 high-power fields (hpf) per section.

EM

Scanning EM. Kidney tissues were rinsed with PBS and postfixed with 1% aqueous osmium tetroxide for a total of 2.5 hours. Subsequently, tissues were rinsed and dehydrated through a graded series of ethanol to abso-
sion values were normalized using 18s RNA as a housekeeping gene. All primers used for amplification were from Metabion. The expression of KIM-1, IL-6, and TNF-α was analyzed using the following primers: Kim-1 forward, TCAGCTCGGGAATGCACA; Kim-1 reverse, TGGTTGCCTTCGTTGCT; L-FABP forward, AGGCAATAGGTCTGCCCGAGGAC; L-FABP reverse, CCAGTTCGCACTCCTCCCCCA; IL-6 forward, TGATGCACTTGCAGAAAACA; IL-6 reverse, ACCAGAGGAAATTTTCAATAGGC; TNF-α forward, CCACCACGCTCTTCTGTCTAC; TNF-α reverse, AGGGTCTGGGCCATAGAACT.

Flow cytometry
Anti-mouse CD11c, CD40, CD86, and CD45 antibodies (BD Biosciences — Pharmingen) were used to detect activated DC and leukocyte populations in kidneys. F4/80, 7/4, and Ly6G antibodies (Abd Serotec) and CD3, NK1.1, CD11b, and CD103 antibodies (BD Biosciences — Pharmingen) were used to detect macrophage, neutrophil, T cell, natural killer cell, and mononuclear phagocyte populations in kidneys. Respective isotype antibodies were used to demonstrate specific staining of cell populations. Quantification of cell number was achieved using counting beads for FACS (Invitrogen).

Cell culture studies
BMDCs were established as described previously (50). Renal DCs were isolated from whole kidneys by magnetic bead separation (MACS) using CD11c MicroBeads (Miltenyi Biotech). The purity of CD11c+ renal cells was confirmed by FACS analysis using anti-CD11c antibody (BD Biosciences — Pharmingen). Cells were stimulated for 3 hours with 1 μg/ml LPS followed by 100 μg/ml CaOx crystals or stimulated with 5 mM ATP for 6 hours (all Sigma-Aldrich). Cell-free supernatants were analyzed for cytokine secretion by ELISA (BD Biosciences — Pharmingen) and immunoblot analysis. Primary TECs were lute ethanol and critical-point dried using liquid CO2. After mounting on stubs, the specimens were sputter-coated with a gold-palladium alloy for 1.5 minutes. DCs in suspension were rinsed in PBS and centrifuged to concentrate the cells. Then cells were dropped on coated coverslips and allowed to adhere; coverslips were fixed as above. Coverslips were cut into pieces prior to mounting on stubs. Images were viewed with a Hitachi 2600 electron microscope.

TEM. Kidneys were immersed in cold modified Karnovsky fixative containing 3% glutaraldehyde and 1% paraformaldehyde in sodium cacodylate buffer (pH 7.4) processed as previously described (48). In another set of experiments, BMDCs were stimulated with 1 μg/ml LPS and 100 μg/ml CaOx, with or without cytochalasin D pretreatment. After stimulation, cells were fixed with 2% paraformaldehyde with 2% glutaraldehyde in PBS buffer (pH 7.4) and processed. The samples were viewed with a JEOL model 1200EX electron microscope.

Quick-freeze deep-etch EM. 1.5-mm bread slices of nonfixed kidney or kidney sections fixed overnight in 2% glutaraldehyde in NaHCa (100 mM NaCl, 30 mM Hapes, and 2 mM CaCl2, pH 7.2) were used after rinsing in NaHCa. Tissues were then cooled to 4°K with liquid helium; mounted in a Balzers 400 vacuum evaporator, fractured, and etched for 2.5 minutes at −104°C; rotary replicated with approximately 3 nm platinum, as previously described (49); and viewed using a JEOL 1400 microscope with attached AMT digital camera.

RNA preparation and real-time quantitative RT-PCR
Total RNA was isolated from kidneys using a Qiagen RNA extraction kit following the manufacturer’s instructions. After quantification, RNA quality was assessed using agarose gels. From isolated RNA, cDNA was prepared using reverse transcriptase (Superscript II; Invitrogen). Real-time quantitative RT-PCR (TaqMan) was performed using SYBRGreen PCR master mix and analyzed with a Light Cycler 480 (Roche). All gene expression values were normalized using 18s RNA as a housekeeping gene. All primers used for amplification were from Metabion. The expression of KIM-1, IL-6, and TNF-α was analyzed using the following primers: Kim-1 forward, TCAGCTCGGGAATGCACA; Kim-1 reverse, TGGTTGCCTTCGTTGCT; L-FABP forward, AGGCAATAGGTCTGCCCGAGGAC; L-FABP reverse, CCAGTTCGCACTCCTCCCCCA; IL-6 forward, TGATGCACTTGCAGAAAACA; IL-6 reverse, ACCAGAGGAAATTTTCAATAGGC; TNF-α forward, AGGGTCTGGGCCATAGAACT; TNF-α reverse, AGGGTCTGGGCCATAGACT.
isolated from the kidneys and grown on type IV collagen–coated tissue culture plates (Sigma-Aldrich). Primary renal ECs and MCs were isolated as described previously (26). The isolated primary cells were confirmed under light microscope and further characterized by FACS, immunostaining, Western blot, and RT-PCR. Cells were grown to confluency before use. 1 × 10⁶ cells per well were used. Primary TECs were maintained in DMEM/F12 (Sigma-Aldrich) containing 10% FCS (Sigma-Aldrich), 1% penicillin-streptomycin (Sigma-Aldrich), 125 ng/ml prostaglandin E1 (Calbiochem), 25 ng/ml EGF (Sigma-Aldrich), 1.8 μg/ml l-thyroxine (Sigma-Aldrich), 3.38 ng/ml hydrocortisone (Sigma-Aldrich), and 2.5 mg/ml insulin–transferrin–sodium selenite supplement (I-T-SS; Sigma-Aldrich). Primary endothelia and MCs were maintained in DMEM (Gibco, Invitrogen) plus 10% FCS and 1% penicillin-streptomycin, and primary renal DCs were maintained in RPMI (Gibco, Invitrogen) and used immediately. All cells were cultured in an incubator at 37°C, 5% CO₂. Supernatants were analyzed for LDH release using the CytoTox 96 Non-Radioactive Cytotoxicity Assay Kit (Promega) to evaluate cell death, measuring LDH after 18 hours. Results were expressed as percent of positive control. For ATP release, the system bioluminescence detection kit for ATP measurement was used (Promega), following the manufacturer’s instructions.

**Coculture studies**

Briefly, TECs and BMDCs were generated as described above. TECs were treated with or without 100 μg/ml CaOx for 18 hours, after which the supernatants were treated with 1 U/ml apyrase (Sigma-Aldrich). Next, LPS-primed (1 μg/ml) or unprimed BMDCs were cocultured with the TECs at a 4:1 ratio in RPMI 1640 (Gibco, Invitrogen) supplemented with 10% FBS. After 24 hours of coculture, the supernatants were collected, centrifuged, and analyzed for IL-1β concentration by ELISA (BD Biosciences — Pharmingen).

**Immunoblots**

Upon completion of the 6-hour stimulation experiment, supernatants and cells were collected. Supernatants were concentrated using Amicon Ultra centrifugal filters (Millipore Billerica) and resolved on 15% SDS polyacryl-
amide gels. After electrophoresis, proteins were transferred to PVDF membrane. Primary antibodies were goat polyclonal anti–IL-1β (B&F401; R&D Systems) and rabbit polyclonal anti–caspase-1 (sc-514; Santa Cruz); this was followed by incubation with secondary antibody anti-gold or anti-rabbit IgG labeled with HRP. Immunostained bands were detected using a chemiluminescence kit (ECL kit).

Statistics

Data are presented as mean ± SEM. Comparison of groups was performed using ANOVA; post-hoc Bonferroni correction was used for multiple comparisons. A P value less than 0.05 was considered significant.

Study approval

All experimental procedures were approved by the local government Study approval.

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