Altered renal tubular expression of the complement inhibitor Crry permits complement activation after ischemia/reperfusion

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Altered renal tubular expression of the complement inhibitor Crry permits complement activation after ischemia/reperfusion

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Ischemia/reperfusion (I/R) of several organs results in complement activation, but the kidney is unique in that activation after I/R occurs only via the alternative pathway. We hypothesized that selective activation of this pathway after renal I/R could occur either because of a loss of complement inhibition or from increased local synthesis of complement factors. We examined the relationship between renal complement activation after I/R and the levels and localization of intrinsic membrane complement inhibitors. We found that loss of polarity of complement receptor 1–related protein y (Crry) in the tubular epithelium preceded activation of the alternative pathway along the basolateral aspect of the tubular cells. Heterozygous gene-targeted mice that expressed lower amounts of Crry were more sensitive to ischemic injury. Furthermore, inhibition of Crry expressed by proximal tubular epithelial cells in vitro resulted in alternative pathway–mediated injury to the cells. Thus, altered expression of a complement inhibitor within the tubular epithelium appears to be a critical factor permitting activation of the alternative pathway of complement after I/R. Increased C3 mRNA and decreased factor H mRNA were also detected in the outer medulla after I/R, suggesting that altered synthesis of these factors might further contribute to complement activation in this location.

Introduction
Complement activation has been shown to be an important event in the development of ischemic acute renal failure (ARF) in mice. Studies in complement-deficient mice have shown that these mice are protected from renal failure after ischemia/reperfusion (I/R) (1, 2), and that generation of the anaphylatoxin C5a (3) and the membrane attack complex (2) may contribute to the pathogenesis of ischemic ARF. Treatment with agents that inhibit the complement cascade at specific steps during the activation cascade has proven effective at ameliorating ischemic ARF (3, 4). Determining the initiating mechanisms of complement activation after renal I/R is essential to the effective use of complement inhibitors for the treatment and prevention of ischemic ARF.

Complement activation after renal I/R occurs via the alternative pathway (1) and is independent of natural antibody (5). In contrast, I/R injury of other organs such as the heart, intestine, and skeletal muscle appears to involve activation of the classical pathway of complement. Complement activation after intestinal I/R, for example, is the result of natural antibodies that bind to neoantigens exposed within the ischemic bowel (6, 7). Cardiac I/R also appears to involve classical pathway activation by antibodies that recognize newly exposed antigens (8). Renal I/R therefore involves unique mechanisms of complement activation.

The alternative pathway is ordinarily activated at low levels via an enzymatic process, called “tickover,” which results in continuous low-level production of C3b. When C3b binds covalently to an activating surface, such as a bacterial cell wall, it is stabilized such that it can combine with factor B to create the alternative pathway C3 convertase. Such activation of the alternative pathway does not usually cause injury to self cells due to the presence of membrane-bound and fluid-phase complement regulatory proteins (9). These inhibitors are essential to prevent complement-mediated injury to the host, and deficiency of complement inhibitors can lead to spontaneous injury. For example, humans, pigs, and mice deficient in the fluid-phase factor H spontaneously develop membranoproliferative glomerulonephritis (10), the expression of dysfunctional CD46 is associated with development of the hemolytic uremic syndrome (11, 12), and paroxysmal nocturnal hemoglobinuria is associated with deficiencies of decay-accelerating factor (DAF, also known as CD55) and CD59 (13).

Local synthesis of complement components has also emerged as an important cause of complement activation and tissue injury in some models. For instance, in a murine renal transplant model, synthesis of C3 by the kidney was found to contribute to complement activation within the tubulointerstitium and, as opposed to serum C3, to be essential to graft loss (14).

Renal I/R in the mouse (1) and in humans (15) leads to the marked deposition of complement activation products along the tubular basement membrane. Although several inhibitors of complement activation are present within the mouse kidney, only complement receptor 1–related protein y (Crry) is present on mouse tubular epithelial cells (16), the cells primarily injured during I/R. DAF and CD59 expression is limited to the glomeruli and...
renal arteries (17, 18). Deficiency in DAF alone or DAF and CD59 results in vascular complement activation after I/R and worse renal injury (19, 20). However, substantial vascular complement activation does not occur in wild-type mice after I/R (2). Therefore, in the normal host these particular inhibitors must retain their function, preventing further injury. In contrast, the activation of complement along the tubular basement membrane of wild-type mice after I/R suggests that local inhibition by Crry is either hindered or overwhelmed.

Given the importance of membrane-bound Crry for the prevention of alternative pathway complement activation and tubular injury, we hypothesized that altered expression or localization of Crry by tubular epithelial cells might contribute to the activation of complement after I/R, or that local synthesis of complement C3 or factor B in response to I/R could overwhelm tubular-intrinsic Crry. Therefore, we set out to study the effects of renal ischemia upon Crry and the relation of changes in Crry expression, local synthesis of complement components, and complement activation. We found that altered tubular expression of Crry precedes, and appears to permit, complement activation along the basolateral aspect of the tubules. Furthermore, C3 mRNA was increased and factor H mRNA was decreased in the outer medulla after I/R, indicating that altered synthesis of these factors might further contribute to alternative pathway activation at this location. Understanding the mechanisms whereby intrinsic complement inhibitors are unable to control activation may help in the rational use of complement inhibitors as therapeutics for the prevention of ischemic ARF. These studies may also further our understanding of more general inflammatory responses to injury.

**Results**

*Localization of intrinsic complement inhibitors within the kidney.* Frozen sections of kidneys from unmanipulated mice were examined by immunofluorescence to determine the localization of the membrane-bound complement inhibitors. Consistent with previous reports (14), Crry was heavily expressed within the glomeruli as well as the tubules (Figure 1). Crry in the proximal tubules was polarized to the basolateral surface of the epithelial cells. Expression was contained within the basement membrane, and no Crry was detectable within the cytoplasm or brush border. To further evaluate Crry expression by proximal tubular cells, immunogold labeling and evaluation by electron microscopy were performed on rat kidneys (Figure 2). Labeling within both the apical (34 ± 4 particles/field, n = 22 fields) and basal regions (67 ± 8 particles/field, n = 19 fields) was significantly above background labeling with nonimmune serum (12 ± 2 particles/field, n = 10 fields; P < 0.01 versus apical P < 0.001 versus basal labeling with anti-Crry).

### Figure 1

Localization of membrane-bound complement inhibitors within the mouse kidney. Three different membrane-bound complement inhibitors are located within the mouse kidney. (A) Crry is expressed in the glomeruli (arrowhead) and in tubules. (B) Crry in the renal tubules is polarized to the basolateral aspect of the tubules. Inset: Dual staining for Crry (green) and type IV collagen (red) demonstrates that expression of Crry is adjacent to the tubular basement membrane. Staining with secondary antibody alone in unmanipulated kidneys (C) and kidneys subjected to ischemia and 2 hours of reperfusion (D) demonstrate the specificity of staining with the anti-Crry antibody. (E) CD59 and (F) DAF are also expressed in glomeruli (arrowheads). In addition, these inhibitors may be expressed in the small blood vessels (arrows), but not in tubules. Original magnification, x200 (A and C–F); x400 (B).

### Figure 2

Immunogold localization of Crry in PTECs. Immunogold labeling of Crry in kidney sections was performed, and the labeling of PTECs (A) was examined (arrow, brush border; arrowhead, basal cell surface). Gold particles could be seen clustered along the basal aspect of the cells (B), and was present to a lesser degree in the apical brush border (C). Original magnification, x1,200 (A), x18,500 (B and C).
unpaired Student’s t test). Labeling in the region of the basolateral cell surface was significantly higher than that within the apical brush border (P < 0.001, unpaired Student’s t test). As previously reported (17, 18), DAF and CD59 were expressed within the glomeruli and vasculature, but not in the tubules (Figure 1).

**Complement activation is an early event after renal I/R**. Mice were subjected to 24 minutes of bilateral renal ischemia or sham surgery and were sacrificed after 0, 2, 6, 24, and 48 hours of reperfusion. Serum was collected for serum urea nitrogen (SUN) measurements, and kidneys were evaluated for tubular injury and complement C3 deposition. Consistent with previous reports, sham-treated animals demonstrated small areas of C3 deposition along the tubular basement membrane of approximately 10–15% of the tubules in the cortex and outer stripe of the outer medulla (1, 21).

When tubules within the outer medulla were examined by fluorescence microscopy, a significant increase in the percentage of tubules showing complement deposition along the tubular basement membrane was seen by 6 hours of reperfusion (Figure 3A), similar to results previously reported by other groups (22). The degree of tubules showing complement deposition peaked at 24 hours of reperfusion, involving 46–75% of the tubules, and returned to baseline by 48 hours of reperfusion. Western blot analysis of total kidney lysates demonstrated a clear increase in complement activation products after ischemia and 24 hours of reperfusion (Figure 3D).

Although some epithelial cell swelling and brush border injury were evident within 6 hours of reperfusion (Figure 3B), the acute tubular necrosis (ATN) grade did not significantly increase until after 24 hours of reperfusion and returned to baseline by 48 hours of reperfusion (Figure 3B). Thus, complement activation is an early event that precedes severe morphological injury after I/R. Although SUN levels did not significantly increase until after 24 hours of reperfusion (Figure 3C), this was expected because serum markers do not reflect changes in renal function until a new steady state is reached.

**Crry levels in the whole kidney do not significantly change after I/R**. Crry is present in multiple isoforms in the kidney (23). Analysis of protein from immortalized proximal tubular epithelial cells (PTECs) by Western blot revealed that most of the Crry in proximal tubular cells was of the 53-kDa isoform (Figure 4A). Protein from rat kidneys demonstrated that 53-kDa Crry was present in basolateral membrane–enriched fractions but not in brush border–enriched fractions (Figure 4B). Blotting for type IIa Na+/Pi cotransporter, which is known to be expressed on the apical brush border, was stronger on the brush border–enriched fraction. As further evidence that it is the 53-kDa isoform that is expressed by PTECs, the
occasional focal C3 deposition along the tubular basement mem-
"...
other 2 Crry isoforms were not detected in the basolateral mem-
brane–enriched fraction. When protein was isolated from whole-
kidney sections, little difference in overall levels of Crry was dis-
cerned (Figure 4C), likely due to the fact that Crry is expressed in
glomeruli and in tubules throughout the kidney. However, a trend
toward decreased expression of the 53-kDa isoform was seen after
6 hours of reperfusion. Semi-quantitative PCR was performed on
tissue isolated from the outer medulla (Figure 4D). A trend toward
increased mRNA for Crry was seen after 6 hours of reperfusion.

Altered Crry localization precedes complement activation. As discussed
above, Crry is the only complement inhibitor expressed on the sur-
face of renal tubules. At baseline, Crry was polarized to the baso-
lar aspect of the renal tubules (Figure 1), abutting the site of
occasional focal C3 deposition along the basolateral basement mem-
brane (Figure 5A). In kidneys subjected to 24 minutes of ischemia
this polarity was lost before reperfusion, and Crry could be seen
within the tubule lumen (Figure 5B). As shown in Figure 3, the
tubules showed signs of injury at these early time points but did
not yet show necrosis. The altered Crry expression is therefore a
biochemical change of non-necrotic tubules. After 6 and 24 hours
of reperfusion (Figure 5, C and D), individual tubules demonstrat-
ed little remaining Crry. Those tubules that no longer demonstrat-
ed Crry showed little C3 deposition, whereas those with more severely
altered Crry localization showed extensive C3 deposition. 

Because Crry appeared in the tubular lumen after ischemia, we
examined the urine for the presence of Crry. It is known that PTECs
shed F-actin–containing membrane vesicles into the tubular lumen
after I/R (24). We found that the urinary sediment of mice subjected
to I/R contained both F-actin and Crry (Figure 5F). This indicates
that loss of Crry in the blebs and vesicles shed in the urine after I/R
partially accounts for the absence of Crry in some tubular segments.

Lower-powered views of kidneys stained for Crry and C3 demonstrat-
ed that Crry and C3 did not colocalize along the tubules. At
baseline (Figure 6A), Crry was polarized to the basolateral aspect of
tubules, and sporadic patches of C3 were seen along the tubular
basement membrane. Little colocalization (yellow) was seen.

In order to quantitate colocalization of Crry and C3, we performed
digital microscopy and measured the color intensity at each wave-
length (Figure 6, D and E). Pixels were plotted according to their
Crry and C3 intensity, and the number of pixels within the image
that had a given staining intensity was indicated by color. A histo-
gram of Crry and C3 intensity (Figure 6D) showed that there was
virtually no colocalization of C3 and Crry. Pixels with a high inten-
sity of both markers would map to the upper right quadrant. Very
few pixels were present within this quadrant, indicating that those
pixels with a high intensity for Crry staining did not have abun-
dant C3. After 24 hours of reperfusion (Figure 6, C and E) there was
much greater overall C3 deposition than at baseline, but there was
still little, if any, colocalization of C3 and Crry. Renal I/R injury is
patchy in nature, and some tubules retained their polarity (Figure
6C, arrows) even in areas of extensive injury. Consistent with the
histogram in Figure 6E, those tubules that retained the polarity of
Crry showed little C3 deposition, whereas those with more severely
altered Crry localization showed extensive C3 deposition.

Our results demonstrate that Crry localization was altered
during renal ischemia prior to complement deposition and that
tubules that retained Crry expression showed little C3 deposition.
These data suggest that altered Crry localization is permissive of
complement activation and that altered complement inhibition is
a necessary event in the activation of complement after renal I/R.
Although overall expression of Crry in the kidney did not signifi-
cantly decrease (Figure 4), altered expression of Crry specifically on
the surface of the tubules appeared to permit complement activa-
tion at that specific location.
Altered Crry localization occurs primarily in the proximal tubule. Dual staining of kidney sections for Crry and a distal tubule marker revealed that altered Crry localization occurred primarily in proximal tubules (Figure 7A). It has long been known that Na/K-ATPase, another protein polarized to the basolateral aspect of PTECs, is associated with the actin cytoskeleton. Many studies have shown that I/R causes Na/K-ATPase to lose its polarity, and that after I/R the protein is seen diffusely throughout the cell cytoplasm (25, 26). Dual staining for Crry and Na/K-ATPase demonstrated that these proteins colocalized to the basolateral membrane of proximal tubules at baseline (Figure 7B). After I/R both shifted into the cytoplasmic domain. Colocalization of the proteins was weaker after I/R, however, suggesting distinct subcellular localizations.

Crry shifts from a triton-insoluble to a triton-soluble form after I/R. The polarity of many transmembrane proteins is maintained by attachment to the actin cytoskeleton (27), an association that can render a protein triton insoluble. A change in a renal tubular protein’s solubility in triton after I/R may therefore reflect the disruption of this cytoskeletal attachment (25). To explore the possibility that tubular Crry is attached to the cytoskeleton, we examined the triton solubility of Crry before and after I/R. At baseline, approximately half of the 53-kDa Crry isoform was present in the triton-extractable fraction of whole-kidney sections (Figure 7C). After I/R, however, nearly all of the protein was present in the triton-soluble fraction.

Figure 5
The polarity of Crry is lost after I/R. Immunofluorescence microscopy was performed on tissue sections at baseline and after ischemia and variable times of reperfusion. (A–E) Dual staining for Crry (green) and C3 (red) was performed. (A) At baseline Crry was polarized to the basolateral aspect of the tubules in the kidney and scant C3 deposition was seen. (B) After 24 minutes of ischemia, the polarity of Crry expressed by proximal tubules was lost. Crry was seen more diffusely throughout the PTEC and within the tubule lumen (arrow). After 6 (C) and 24 hours (D) of reperfusion, an increasing percentage of tubules in the outer medulla ceased to express Crry. Those tubules that no longer expressed Crry showed extensive C3 deposition along their tubular basement membranes. Dashed lines indicate the tubule lumen, demonstrating that the tubule was still lined by epithelial cells although Crry was not detectable. (E) By 48 hours of reperfusion, Crry expression and the polarity of Crry was restored to baseline. (F) Staining of urinary sediment for F-actin (red) and Crry (green) revealed that after 2 hours of reperfusion, the sediment contained cellular blebs (as evidenced by the F-actin) and Crry. Original magnification, ×400.

Figure 6
Complement deposition after I/R does not colocalize with Crry. (A–C) Lower-powered views (original magnification, ×100) of the kidney demonstrate that the deposition of C3 during reperfusion only occurred along tubules that lost basolateral expression of Crry. Glomeruli (arrowheads) indicate cortical tissue, and the outer medulla is shown toward the bottom of each frame. (A) At baseline, Crry was heavily expressed in the glomerulus and along the basolateral aspect of the tubules. (B) After 2 hours of reperfusion, the polarity of Crry in many tubules was disrupted. (C) After 24 hours of reperfusion, most of the tubules had diminished levels of Crry. C3 was extensively deposited along the proximal tubules. Some tubules retained normal Crry expression and polarization (arrows) and showed little C3 deposition. In areas of heavy C3 deposition, little colocalization of C3 and Crry (yellow) is seen. (D and E) Histograms of Crry and C3 staining at baseline (D) and after 24 hours of reperfusion (E) further demonstrate that little colocalization occurred. Figures are mapped according to Crry and C3 intensity. (D) The heat map shows the relative population densities depicted by color. Pixels with a high intensity of both C3 and Crry are plotted in the upper right quadrant. (E) Even after 24 hours of reperfusion, when the population of pixels with a high C3 intensity was much increased, virtually none of these high C3 pixels demonstrated concurrent high-intensity staining for Crry.
other isoforms, in contrast, were primarily triton insoluble in both nonischemic and postischemic samples. This supports the hypothesis that the polarization of 53-kDa Crry in tubular epithelial cells is dependent upon attachment to the cytoskeleton, and the disruption of this attachment explains the loss of polarity seen after I/R.

**Deficiency of Crry renders mice susceptible to renal I/R injury.** To confirm that the loss of inhibition by Crry is an important event in the pathogenesis of ischemic ARF and complement activation, we subjected Crry-deficient and wild-type mice to a milder renal insult (Figure 8). Crry<sup>−/−</sup> mice do not directly address the issue of redistribution per se. They do, however, address the issue of the “concentration” of the complement inhibitor at the site of activation. We hypothesized that loss of functional Crry at the basolateral tubule would render the tubules more susceptible to injury, whether due to redistribution, targeted gene deletion, or the exogenous administration of an inhibitor of Crry (28). We therefore predicted that mice expressing less Crry at this location in the first place are more susceptible to complement activation. When subjected to 20 minutes of ischemia and 24 hours of reperfusion, Crry<sup>−/−</sup> mice had significantly greater SUNs (111 ± 18 versus 52 ± 8 mg/dl; *P* < 0.01) and significantly greater morphological injury (average ATN score, 3.9 ± 0.3 versus 2.56 ± 0.6; *P* < 0.05) than did wild-type controls. Thus, mice that express decreased levels of Crry develop more severe injury in response to a mild ischemic insult.

**Crry is critical for protecting PTECs in culture from alternative pathway–mediated injury.** Complement activation may injure the tubular epithelium directly or through indirect effects such as alterations to the vasculature or chemotaxis of inflammatory cells. In order to confirm that PTECs themselves are the direct target of local alternative pathway–mediated injury when Crry function is impaired, we developed an in vitro model of PTEC injury. We incubated PTECs with fresh mouse serum in a buffer solution containing 10 mM EGTA in order to inhibit classical pathway activation. Treatment of these cells with an inhibitory rat IgG1 mAb to Crry in order to suppress the function of this protein caused injury to the cells as assessed by lactate dehydrogenase (LDH) release (Figure 9), whereas the addition of control rat IgG1 had no effect. Similar injury was also seen when Fab fragments of the inhibitory antibody were used, indicating that the injury is not a nonspecific effect of antibody binding or Fc-mediated complement activation. We then added the same inhibitor to a reaction mix in which the alternative complement pathway was selectively blocked with a mAb to factor B. Without an intact alternative pathway, blocking the function of Crry did not result in injury to the cells. Thus, functional Crry on the surface of PTECs is essential to the prevention of alternative pathway–mediat-
ed cellular injury. As described above, Crry was the only membrane-bound inhibitor of complement expressed by PTECs. The loss of complement inhibition by Crry on the surface of PTECs, whether by the application of an inhibitor of this protein in vitro or by the altered expression of Crry after ischemia, is therefore sufficient to permit alternative pathway-mediated injury of the cells.

Local synthesis of alternative pathway components after I/R. In other models, local synthesis of complement by the kidney has been found to be an important cause of complement activation and tissue injury (14). The kidney has been shown capable of synthesizing C3, factor B, and factor H (29, 30), and renal I/R in rats was demonstrated to increase renal C3 expression (31). To determine whether the complement activation that occurs after I/R is enhanced by local synthesis of alternative pathway substrates, mice were subjected to sham surgery or to I/R. After variable times of reperfusion, the outer medulla (where C3 deposition is heaviest) was dissected out, and real-time PCR was performed to quantitate expression of C3, factor B, and factor H (Figure 10). C3 mRNA did not significantly increase at any of the times of reperfusion, although at 24 hours there was a trend toward increased C3 mRNA. Because of the tendency toward increased C3 synthesis after I/R, we more specifically evaluated the tubules of the outer medulla for changes in C3 synthesis using in situ hybridization (Figure 10). Local synthesis of C3 was evident in the outer medullas of sham-treated animals, and synthesis in this region increased after 24 hours of reperfusion. Factor H mRNA in the outer medulla was significantly decreased from baseline after 2 and 24 hours of reperfusion, as determined by real-time PCR. Thus increased local synthesis of C3 and decreased synthesis of factor H may further facilitate alternative pathway activation during reperfusion. Given the basal synthesis of C3 in this region of the kidney, it is also possible that a pool of locally produced C3 contributes to the activation that occurs after I/R. Furthermore, even small increases in C3 synthesis may serve to initiate a much larger degree of complement activation, subsequently utilizing circulating components of the alternative pathway.

Discussion

Complement activation by the alternative complement pathway is an important pathogenic event after I/R (1, 2). C3 deposition primarily occurs along the tubular basement membrane of proximal tubules in the outer stripe of the outer medulla, and Crry is the only membrane-bound inhibitor of complement expressed by the tubular epithelial cells in this segment of the nephron. At baseline, Crry is polarized to the basolateral aspect of the tubules, adjacent to where complement deposition occurs after I/R. Mice that are
increased Crry mRNA in the outer medulla suggests that replacement of Na/K-ATPase, this biochemical shift of Crry probably reflects the occurrence only after the loss of Crry at this location. Despite abundant C3 deposition after I/R there was little colocalization of C3 and Crry, suggesting that the loss of Crry at the basolateral surface is necessary for activation of the alternative pathway. Conversely, those tubules that retained polarized expression of Crry along their surface demonstrated little C3 deposition.

Crry is analogous to other polarized membrane-associated proteins such as Na/K-ATPase, which has long been known to lose its polarity during ischemia (25). Like Na/K-ATPase, Crry within the kidney also shifted from a triton-insoluble fraction to an almost entirely triton-soluble state during ischemia. As is the case with Na/K-ATPase, this biochemical shift of Crry probably reflects the loss of a stabilizing association, such as with the actin cytoskeleton. The rapid change in the triton solubility of Crry and movement of it away from the basolateral membrane preceded detectable activation of complement on the tubular cells.

Treatment with an inhibitor of Crry leads to increased complement activation and spontaneous tubular injury in rats (28). Furthermore, we have found that mice expressing lower levels of Crry (Crry<sup>−/−</sup> mice) were more susceptible to renal I/R. This supports the hypothesis that a decreased concentration of Crry at the basolateral aspect of the PTEC (whether due to a targeted gene mutation, the administration of an exogenous inhibitor, or altered expression in response to I/R) leaves the tubules more susceptible to complement-mediated injury. When the function of Crry expressed by PTECs in culture was blocked, the cells were injured in the presence of alternative pathway–sufficient serum. Blocking the function of Crry did not injure PTECs when the alternative complement pathway in the serum to which they were exposed was also inhibited.

Previous studies have shown that PTECs in culture activate the alternative pathway when they are heat killed (34), and that such activation results in an inflammatory response by the cells (35). It is possible that altered expression of Crry is a general response to injury of these cells, and that complement activation by tubular epithelial cells also occurs in other forms of tubular injury. In support of this, we have observed that glycerol-induced rhabdomyolysis (a model of toxin-induced ATN) is also characterized by decreased tubular Crry and peritubular C3 deposition (our unpublished observations).

We have examined the effects of I/R on the intrinsic inhibitors of complement activity and on local synthesis of the complement components. Several previous studies have highlighted the contribution of locally synthesized C3 to the development of renal injury (14, 31). Real-time PCR of samples made from the outer medulla demonstrated that there was basal synthesis of C3 in this region, and synthesis of C3 was increased after I/R when evaluated.

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research article
Membrane cofactor protein (MCP, also known as CD46), a human functional analog of Crry, is expressed by human PTECs in a polarized fashion (38). Studies have demonstrated that the polarity of PTECs is disrupted in human ATN (39). Loss of polarity of MCP after I/R may also therefore occur, permitting activation of the alternative pathway.

Our results may also have broader implications regarding the regulation of complement activation. The importance of the complement system in the pathogenesis of immune complex diseases as well as the mechanisms of complement activation in these diseases are well established. The complement system has recently been implicated in a large number of Ig-independent diseases (40–42), however, and the cause of activation in these diseases has yet to be fully elucidated. Although genetic complement deficiencies are known to cause autoimmune disease (10, 13), nongenetically determined loss of complement inhibition of a specific organ or cell type may significantly contribute to inflammation and organ injury. Studies in rats have shown that CD59 expression in the myocardium decreases after ischemia, perhaps allowing greater assembly of the membrane attack complex (43), although in that setting complement activation precedes loss of inhibition. Anoxic injury of human endothelial cells in culture also results in diminished expression of cell surface complement inhibitors and activation of the complement system on the cell surface (44). Given the need for constant inhibition of the alternative pathway to prevent spontaneous injury, it is possible that other injuries in various tissues may impede this tonic inhibition and result in alternative pathway activation. Locations such as the renal tubules that rely on only 1 complement inhibitor may be particularly vulnerable to such injuries.

We have demonstrated that altered localization and expression of renal tubular epithelial Crry is an early consequence of renal ischemia. The loss of inhibition by Crry is permissive of complement activation on the surface of epithelial cells and of the resultant complement-mediated injury. These results explain the unique nature of complement activation in the kidney after I/R. Furthermore, these findings demonstrate that complement activation may be a response to any injury that alters host expression of complement inhibitors. Inhibition of the alternative pathway may be an effective means of ameliorating ischemic ARF. The loss of complement inhibition and the activation of the alternative pathway are early events after I/R, however, and exogenous inhibition may need to be performed soon after the ischemic event. The exact localization of cell surface complement inhibitors is clearly very important, and exogenous inhibitors of complement may not effectively prevent activation of complement on the epithelial cell surface if they do not achieve adequate levels specifically at this site.

**Methods**

*Experimental animals.* C57BL/6J mice (The Jackson Laboratory) were used as wild-type mice for experiments examining the relationship of Crry expression and complement activation. Crry-deficient mice were generated as previously described (45). Mice that were heterozygous for Crry (Crry<sup>+/–</sup>) and littermate controls, and C57BL/6J mice were used to test the

### Table 1

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for, forward; rev, reverse.
effects of Crry deficiency since homozygous deficiency is lethal in utero (45). Only male mice were used in this study because they demonstrate a higher level of complement activity and are more vulnerable to complement-mediated renal injury (32). Sprague-Dawley rats were used for isolation of renal basolateral and brush border membranes and for immunogold labeling. The animals were housed and maintained in the University of Colorado Center for Laboratory Animal Care in accordance with the NIH Guidelines for the Care and Use of Laboratory Animals. The study protocol was approved by the University of Colorado Health Sciences Center Animal Care and Use Committee.

Antibodies and primers. The following antibodies were used for immunofluorescence: FITC-conjugated goat anti-mouse C3 (Cappel; MP Biomedicals) diluted 1:50 with PBS, rat anti-mouse Crry (BD Biosciences—Pharmingen) diluted 1:150, rabbit anti-human C3d (Dako) diluted 1:150, mouse anti-Na/K-ATPase α-1 (Upstate USA Inc.) at a concentration of 0.5 μg/ml, rabbit anti-mouse collagen type IV (Chemicon International) diluted 1:400. FITC-conjugated Arachis hypogaea (Sigma-Aldrich) was used as a distal tubule marker (46). A polyclonal rabbit anti-DAF antibody and a mouse anti-CD59 mAb (a generous gift of S. Tomlinson, Medical University of South Carolina, Charleston, South Carolina, USA) were diluted 1:300. To stain F-actin, Alexa Fluor 350 phallolidin (Invitrogen Corp.) was diluted 1:40. For secondary antibodies, FITC-conjugated anti-rat IgG, rhodamine-conjugated anti-rabbit IgG, and FITC-conjugated anti-mouse IgG (Cappel; MP Biomedicals) were used at dilutions of 1:100, and Alexa Fluor 488-conjugated goat anti-rabbit IgG was used at a dilution of 1:400 (Invitrogen Corp.). Two mAbs were used as complement protein inhibitors. The mAb S5D (47) was used to inhibit the function of cell-surface Crry, and mAb 1379 (48) was used to block mouse alternative pathway activity. Fab fragments of S5D were also made using papain-agarose (MP Biomedicals) according to the manufacturer’s instructions.

For Western blot analysis, a peroxidase-conjugated goat anti-mouse C3 (Cappel; MP Biomedicals) was used at a dilution of 0.3 μg/ml. A polyclonal antibody to Crry (23) and a polyclonal anti-type Ila Na/Pi cotransporter antibody (49) were used at dilutions of 1 μg/ml, followed by incubation with peroxidase-conjugated goat anti-rabbit IgG (Sigma-Aldrich).

Induction of ischemic ARF. Mice weighing 20–25 g were anesthetized with 300 μl 2,2,2-Tribromoethanol (Sigma-Aldrich) injected intraperitoneally. After the mice were anesthetized they were placed on a heating pad to maintain their body temperature during surgery. Laparotomies were then performed, and the renal pedicles were located and isolated by blunt dissection. The pedicles were clamped with surgical clips (Miltex Inc.), and occlusion of blood flow was confirmed by visual inspection of the kidneys. The clamps were left in place for 24 minutes and then released. This time of ischemia was chosen to obtain a reversible model of ischemic ARF with a minimum of vascular thrombosis and to avoid animal mortality. For experiments comparing Crry+/− mice with Crry−/− controls, an ischemic time of 20 minutes was used. The shorter ischemic time was chosen because it causes milder injury, allowing for detection of mouse strains with greater susceptibility to I/R injury. The kidneys were observed for approximately 1 minute to ensure blood reflow, then fascia and skin were sutured with 4-0 silk (U.S. Surgical). Sham surgery was performed in an identical fashion, except that the renal pedicles were not clamped. The mice were volume resuscitated with 0.5 ml of normal saline and kept in an incubator at 29°C to maintain body temperature. After 0, 2, 6, 24, or 48 hours of reperfusion the mice were anesthetized, and blood was obtained by cardiac puncture. Laparotomy was performed and the kidneys were harvested.

SUN measurements. SUN was determined for each mouse using a Beckman Autoanalyzer (Beckman Coulter).

Renal morphology. After the kidneys were removed from the mice, sagittal sections were fixed in 4% paraformaldehyde. After being embedded in paraffin, 4-μm sections were cut and stained with periodic acid–Schiff. The sections were evaluated by a renal pathologist (D. Ljubanović) in a blinded fashion. The cortex and outer stripe of the outer medulla were assessed for epithelial necrosis, loss of brush border, tubular dilatation, and cast formation. At least 10 fields (magnification, ×400) were reviewed for each slide, and the percentage of tubules displaying these findings was determined. Kidney sections were scored as follows based on the percentage of affected tubules: 0, none; 1, less than 10%; 2, 10–25%; 3, 26–45%; 4, 46–75%; 5, greater than 75%.

Immunofluorescence and in situ hybridization. For immunofluorescence, sagittal sections of the kidneys were snap frozen in OCT compound (Sakura Finetek). Sections (4 μm) were cut with a cryostat and stored at −70°C. The slides were later fixed with acetone and incubated with the primary antibodies. After hybridization with antibodies was complete the slides were counterstained with hematoxylin (Vector Laboratories). Tissue sections were incubated with secondary antibodies alone to ensure the specificity of antibody binding.

To detect the presence of Crry in the urine, immunofluorescence was performed on urine sediments as previously described (24). Upon sacrifice after 6 hours of renal reperfusion, urine was expressed from the bladder and collected at the tip of the urethra. Urine was centrifuged at 4°C for 10 minutes at 15,996 g. The pellets were then fixed for 1 hour at room temperature in 3% paraformaldehyde in PBS (pH 7.4). The sediment was then washed 3 times in PBS for 5 minutes and recentrifuged for 5 minutes at 15,996 g. The pellets were incubated for 10 minutes in 0.1% Triton X-100 (Sigma-Aldrich) in PBS. Nonspecific binding was blocked with 2% BSA in PBS for 1 hour. The pellets were then incubated with rat anti-mouse Crry and Alexa Fluor 350 phallolidin. To ensure that antibody binding was specific, anti–secondary antibody alone was used as a control.

Imaging was performed on a Nikon T2000 inverted fluorescent microscope and slidebook software (version 3.0; Intelligent Imaging Innovations Inc.). Intensity histograms were obtained using a Zeiss LSM 510 Meta confocal microscope running Zeiss software version 3.2 Service Pack 2.

To examine kidneys for the local production of C3, 16-μm kidney sections were fixed for 15 minutes in cold 4% paraformaldehyde. The sections were then hybridized with a digoxin-labeled antisense probe to mouse C3 (Table 1) according to the manufacturer’s instructions (GeneDetect.com). The bound probe was detected using alkaline phosphatase–labeled antidigoxigenin Fab fragments and NBT/BCIP tablets (Roche Diagnostics Corp.). Digoxin-labeled sense probe to mouse C3 was used to confirm the specificity of the binding.

Immunogold labeling of Crry. Rat kidneys were fixed by perfusion with 2% formaldehyde and 0.1% glutaraldehyde in 0.1 M sodium cacodylate, pH 7.35, as previously described (50). The samples were dehydrated and embedded in LR Gold resin (Polysciences Inc.). Ultrathin sections were then prepared using nickel grids and stained for Crry. The grids were floated on droplets of PBS (10 minutes), 10% whole goat serum (10 minutes), and either droplets of rabbit anti-Crry at a dilution of 20 μg/ml or nonimmune rabbit serum diluted 1:150 (45 minutes). The grids were washed using droplets of PBS for a total of 20 minutes, after which each was floated on 10 μl of goat anti-rabbit IgG Fab2 conjugated to 10-nm gold particles (Ted Pella Inc.). After washing the grids with PBS, they were stained for 6 minutes with 2% uranyl acetate, washed with water, and air dried. The sections were then viewed using a Tecnai G^2 microscope (FEI Co.). After identifying PTECs, images were taken at a magnification of ×18,500 in the apical and basal regions of the cells, and the number of gold particles seen in each view were counted.

Western blot analysis. Renal tissue was homogenized in RIPA lysis buffer containing 1% Triton X-100, 0.5% deoxycholic acid, 150 mM NaCl, 20 mM β-glycerophosphate, 20 mM Tris-HCl (pH 8.0), 5 mM EGTA, 3 mM MgCl₂,
0.1% SDS, 1 mM DTT, 50 mM Na2VO4, and 1 tablet of complete, EDTA-free protease inhibitor cocktail (Roche Diagnostics Corp.). Homogenates were centrifuged at 15,996 g for 15 minutes at 4°C, and the supernatant was collected. Brush border and basolateral membranes from rat renal cortical tissue were isolated as previously described (51). Briefly, kidneys were placed in cold isolation buffer containing 300 mM mannitol, 5 mM EGTA, 0.1 mM PMSF, and 18 mM Tris (pH 7.4). Thin cortical sections were obtained and homogenized. Brush border and basolateral membranes were then isolated by differential centrifugation, magnesium precipitation, and use of a discontinuous sucrose gradient (52). To isolate triton-soluble and -insoluble protein fractions, kidneys were homogenized in a solution containing 0.1% Triton X-100, 20 mM HEPES, 120 mM KCl, 12 mM NaCl, 1.62 mM MgSO4, 1 mM EDTA, 0.35 mM CaCl2, and 1 tablet of complete, EDTA-free protease inhibitor cocktail (Roche Diagnostics Corp.). Samples were centrifuged at 35,000 g for 14 minutes at 4°C to separate the soluble and insoluble fractions. Because approximately two-thirds of cellular proteins are triton soluble (53), the insoluble pellet was resuspended at half the volume of buffer used for the initial homogenization. Protein concentrations for all of the samples were determined using the Bio-Rad protein assay (Bio-Rad). Multiple-group comparisons were performed using ANOVA with post-test according to Newman-Keuls. Comparison between the control and Crry−/− groups was performed with the Mann-Whitney test. A P value of less than 0.05 was considered statistically significant. Results are reported as mean ± SEM.

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