



Published in final edited form as:

Compr Physiol. 2012 April ; 2(2): 1303–1353. doi:10.1002/cphy.c110041.

Pathophysiology of Acute Kidney Injury

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Abstract

Acute kidney injury (AKI) is the leading cause of nephrology consultation and is associated with high mortality rates. The primary causes of AKI include ischemia, hypoxia or nephrotoxicity. An underlying feature is a rapid decline in GFR usually associated with decreases in renal blood flow. Inflammation represents an important additional component of AKI leading to the extension phase of injury, which may be associated with insensitivity to vasodilator therapy. It is suggested that targeting the extension phase represents an area potential of treatment with the greatest possible impact. The underlying basis of renal injury appears to be impaired energetics of the highly metabolically active nephron segments (i.e., proximal tubules and thick ascending limb) in the renal outer medulla, which can trigger conversion from transient hypoxia to intrinsic renal failure. Injury to kidney cells can be lethal or sublethal. Sublethal injury represents an important component in AKI, as it may profoundly influence GFR and renal blood flow. The nature of the recovery response is mediated by the degree to which sublethal cells can restore normal function and promote regeneration. The successful recovery from AKI depends on the degree to which these repair processes ensue and these may be compromised in elderly or CKD patients. Recent data suggest that AKI represents a potential link to CKD in surviving patients. Finally, earlier diagnosis of AKI represents an important area in treating patients with AKI that has spawned increased awareness of the potential that biomarkers of AKI may play in the future.

Introduction

In the century that followed Richard Bright's description of kidney disease in 1827 (72) many case studies of acute Bright's disease associated with a variety of etiologies including infections, toxins, and transfusion reactions were published. However, it was the landmark report by Bywaters and Beal in 1941 (84) linking crush injury to the acute impairment of renal function that stands out as the starting point for modern medicine's discussion of acute kidney injury (AKI). This ongoing discussion in the clinic and laboratory has made significant strides in our understanding of AKI, formerly known as acute renal failure (ARF). Nonetheless, AKI remains a significant and serious problem accounting for the most common reason for nephrology consultation in the hospitalized patient. In fact, the progress of modern medicine and the aging population has primarily rendered AKI a nosocomial disease in the developed world. It is the intent of this chapter to provide our current understanding of cell, organ, and systemic alterations involved in the pathophysiology of AKI that will hopefully serve as a framework on which to build further understanding of this vexing syndrome and ultimately provide meaningful therapeutic interventions in the clinic.

I. AKI: History, Etiology, Terminology and Pathology

I.1. Etiology

One of the most important functions of the kidney is the filtration and excretion of nitrogenous waste products from the blood. The measurements of elevated blood urea nitrogen (BUN) and creatinine serve as indicators of decreased renal function indicative of the decreased clearance of these waste products. AKI is currently defined as a rapid decline in the glomerular filtration rate (GFR) resulting in retention of nitrogenous wastes, primarily creatinine and blood urea nitrogen (542). Consequently, the diagnosis currently is dependent on the serial measurement over time of these substances in the blood of patients. The rapidity of this decline may occur over a time course of hours to months, but typically occurs over the course of hours to days. The ability of these measurements to serve as a surrogate marker of GFR is relatively imprecise (138, 363) and improved methods for evaluating GFR and a direct assessment of renal injury are sorely needed in the practice of medicine (96).

Clinically, AKI can be conveniently grouped into three primary etiologies: prerenal, renal, and postrenal (542). All three etiologies will be discussed briefly here; however, renal etiologies, especially those from ischemic and nephrotoxic injury, will receive the bulk of the discussion in this chapter given the frequency with which they occur and the fact that they are the etiologies associated with frank renal tissue injury.

Maintaining a normal GFR is dependent on adequate renal perfusion. Prerenal azotemia is characterized by a decrease in GFR due to a decrease in renal perfusion pressure without damage to the renal parenchyma (22). The kidneys receive up to 25% of the cardiac output and thus any failure of the general circulation or isolated failure of the intrarenal circulation can have a profound impact on renal perfusion. Causes of prerenal azotemia include: hypovolemia resulting from conditions such as hemorrhage, vomiting, diarrhea “third spacing,” poor oral intake, burns, excessive sweating, renal losses (e.g. diuresis); impaired cardiac output resulting from congestive heart failure or decreased cardiac output states (e.g. pericardial tamponade, severe pulmonary hypertension); decreased vascular resistance (peripheral vasodilation) resulting from conditions such as sepsis, vasodilator medications, autonomic neuropathy, or anaphylaxis; and renal vasoconstriction from vasoconstrictive medications or conditions such as hypercalcemia.

The normal response of the kidney to prerenal conditions is to concentrate the urine maximally and avidly reabsorb sodium in an effort to maintain/increase intravascular volume and normalize renal perfusion (372). In general, therapy aimed at rapidly restoring renal perfusion will promptly improve renal function. However, it is important to appreciate that prolonged or profound prerenal azotemia can result in ischemic damage to the kidney and thus there is spectrum from prerenal azotemia to ischemic AKI.

Postrenal causes of AKI are characterized by acute obstruction to urinary flow. Urinary tract obstruction increases intratubular pressure and thus decreases GFR. In addition, acute urinary tract obstruction can lead to impaired renal blood flow and inflammatory processes that also contribute to diminished GFR (200, 222, 234, 500, 581). Obstruction of the urinary tract at any level may produce AKI. In general, obstruction must involve both kidneys or a solitary kidney to produce significant renal failure. However, a patient with pre-existing renal insufficiency may develop AKI with obstruction of only one kidney.

Urinary obstruction may present as anuria or intermittent urine flow (such as polyuria alternating with oliguria) but may also present as nocturia or nonoliguric AKI. Causes of postrenal AKI include benign prostatic hyperplasia and prostate cancer in men, gynecologic

cancers especially cervical cancer in women, retroperitoneal fibrosis, ureteral stones, papillary necrosis, neurogenic bladder, and intratubular obstruction due to precipitation of various substances such as acyclovir or indinavir (95).

Renal etiologies of AKI can be a challenging form of AKI to evaluate because of the wide variety of injuries that can occur to the kidney. In general, it can be helpful to think of damage to the four major structures of the kidney when considering etiologies of intrinsic renal failure. These four structures are 1) the tubules, 2) the glomeruli, 3) the interstitium, and 4) the intrarenal blood vessels.

Tubular Damage—Acute tubular necrosis (ATN) is the term used to designate AKI resulting from damage to the tubules. The two major causes of ATN are:

1. Ischemic – resulting from severe or protracted decrease in renal perfusion.
2. Nephrotoxic – resulting from a variety of exogenous compounds (e.g. aminoglycosides, amphotericin B, cis-platinum, radiocontrast media) and endogenous compounds (e.g. hemoglobin in hemolysis, myoglobin in rhabdomyolysis) that are toxic or potentially toxic to the kidney.

Historically, classic ATN goes through an oliguric (urine output < 400 mL/24 hours) phase of 1–2 weeks followed by a nonoliguric (urine output > 400 mL/day) phase of 10–14 days with eventual recovery of renal function. However, this description is not the rule as both prolonged oliguric phases and initial nonoliguric phases are common (381).

Glomerular Damage—AKI from glomerular damage occurs in severe cases of acute glomerulonephritis (GN). Acute GN can be due to a primary renal disease such as an idiopathic rapidly progressive GN or as part of a systemic disease such as systemic lupus erythematosus, bacterial endocarditis, or Wegener’s granulomatosis (381, 542).

Interstitial Damage—AKI from interstitial damage can result from acute interstitial nephritis due to an allergic reaction to a variety of medications (commonly antibiotics such as penicillins, cephalosporins, sulfonamides) or an infection (bacterial illnesses such as leptospirosis, legionella, rarely pyelonephritis and viral illnesses such as Hanta virus), (381, 542).

Vascular Damage—AKI from vascular damage occurs because injury to intrarenal vessels decreases renal perfusion and diminishes GFR. Causes of vascular injury include malignant hypertension, atheroembolic disease, preeclampsia/eclampsia, and hemolytic-uremic syndrome (HUS)/thrombotic thrombocytopenia purpura (TTP) (381, 542).

I.2. Epidemiology/Outcomes

As mentioned previously, AKI has primarily become a nosocomial disease in the developed world. AKI has an incidence of 5–7% in hospitalized patients (93, 390, 562). This incidence appears to be rising over time (214, 582), which may reflect greater clinical vigilance and/or differences in how AKI is precisely defined. The etiology of AKI differs somewhat depending on if the onset of AKI occurs in the hospital (hospital-acquired) or prior to hospitalization (community-acquired). Depending on the study, prerenal etiologies account for 25–60% (73, 213, 317, 318) and renal etiologies account for 35–70% of AKI cases (73, 93, 213, 317, 318, 363, 389, 562). Ischemic injury or nephrotoxins contribute to 80–90% of the renal etiologies (213, 317, 318, 363). Postrenal etiologies generally account for <5% of AKI in hospitalized patients but this number increases to almost 20% in community-acquired cases (318). The disease burden of AKI results in an estimated \$10 billion in additional costs to the health care system in the United States (93) and is associated with a

mortality of 45–70% (25, 303, 316, 363, 389, 562, 584). The diagnosis of AKI increases the risk of mortality 5.5 to 6.5-fold as compared to a similarly ill patient without AKI (93, 303). Unfortunately, despite advances in knowledge of the pathophysiology of AKI, this figure has not significantly changed since the advent of dialytic therapy which is required in 20–75% of patients (93, 94, 303, 317, 318, 384, 389, 562, 584). There is burgeoning evidence for the direct negative impact that AKI has on distal vital organs (177). These distant organ effects may significantly contribute to the overall mortality observed in patients with AKI despite the initiation of dialytic therapy. For those patients surviving an episode of AKI requiring dialytic therapy, 10–50% continue dialytic therapy after discharge (93, 94, 227, 303, 317, 318, 336, 384, 389, 477, 562, 584). Furthermore, patients that survive an episode of AKI requiring dialytic therapy and recover function are at increased risk of progressing to ESRD (9, 227, 336, 584). In fact, it has been estimated that these patients make up 3% of the overall yearly incidence of ESRD in North America (583, 584). Clearly, AKI has serious acute and chronic sequelae.

I.3. Clinical Pathological Findings

The term ATN is a misnomer. Indeed, the contradiction between the degree of histological abnormalities on human biopsy samples and the extent of GFR depression has served as a one of the main paradoxes in AKI. While the S3 segment of proximal tubule has been generally viewed as the most important site of injury in human ATN (56, 137, 410, 514) the relative contribution of injury to the distal tubule that sits adjacent to the S3 segment in the outer medulla has been a point of controversy (61, 322). Settling this controversy is hampered by the relative lack of medullary tissue routinely obtained on clinical biopsies of AKI. Outside of rare frankly necrotic tubular cells that occur, the prominent findings in human biopsies include detachment of renal tubular epithelial cells from the basement membrane, sloughing of cells into the tubular lumen, effacement and loss of brush border in proximal tubular segments, and the formation of tubular casts derived from sloughed cells, tubular debris, and protein. While evidence of tubular injury is apparent, evidence of tubular cell regeneration can also be seen alongside the injury (514). Tubular cells with basophilic cytoplasm and hyperchromatic nuclei consistent with regenerating epithelial cells and even tubular cells undergoing active mitosis have been observed in many human biopsies of patients with ATN (514).

Alterations of the interstitium, vasculature, and glomeruli have also been documented in ATN. Interstitial edema is often observed and may develop from leakage of fluid from increased microvascular permeability or backleak of tubular filtrate into the interstitium (511, 514). Peritubular accumulation of leukocytes in the interstitium has also been observed in experimental models of ATN (168, 627) although this does not appear to be a predominant feature in human ATN (514). Congestion of the peritubular capillaries in the outer medullary region is a salient feature of ATN that may play an important role in regional alterations of blood flow and exacerbating tubular injury during the extension phase. Additionally the accumulation of leukocytes in the vasa recta (118, 513, 514, 626) may also be an important contributor extending tubular injury. Changes in peritubular capillaries of the cortical region are less prominent but loss of some peritubular capillaries and dilatation of others has been described (276). While changes in arteries and arterioles are relatively subtle, biopsies of ATN have demonstrated vacuolization of smooth muscle cells as well as alteration of smooth muscle cell actin (285). As may be expected, the most noted glomerular alteration in ATN is collapse of the glomerular tuft due to hypoperfusion. Alterations in glomerular foot processes have also been described (516) but this has not been a consistent finding.

II. Pathophysiology, hemodynamics and inflammation

II.1. Phases of acute kidney injury due to Acute Tubular Necrosis

As mentioned above, the term ATN correctly identifies the site of injury, though relatively few renal epithelial cells undergo frank necrosis. More commonly, sublethal changes in renal tubular epithelial cells are present and thus the term acute tubular injury may be more appropriate. Importantly, even sublethal changes can have a major impact on the decrement in GFR that is the hallmark of AKI.

Clinically, ATN and the associated decrease in GFR can be divided into initiation, extension, maintenance, and recovery phases. These clinical phases directly relate to cellular events that occur during the injury and recovery process (Figure 1). Although a clear mechanistic explanation between tubular injury and a fall in GFR has remained elusive, afferent arteriole vasoconstriction in response to tubuloglomerular feedback, backleak of glomerular filtrate, and tubular obstruction have all been postulated as mechanisms for decreased GFR in ATN (see below). All three of these mechanisms can be directly related to changes occurring in renal tubular epithelial cells.

The initiation phase of ATN occurs when renal blood flow (RBF) decreases to a level resulting in severe cellular ATP depletion that in turn leads to acute cell injury and dysfunction. Renal tubular epithelial cell injury is a key feature of the *Initiation Phase* (532). Renal ischemia *in vivo* rapidly induces a number of structural and functional alterations in renal proximal tubular epithelial cells that are directly related spatially and temporally with disruption of the normal framework of filamentous actin (F-actin) in the cell (14, 502, 542) (Described in section III). The extent of these alterations depends upon the severity and duration of ischemic injury. Although these alterations usually fall short of being lethal to the cell, they do disrupt the ability of renal tubular epithelial cells and renal vascular endothelial cells to maintain normal renal function. Additionally, ischemic injury to vascular smooth muscles cells and endothelial cells during the initiation phase may also contribute to the structural abnormalities observed in the renal vasculature during ischemic AKI (285, 358, 541).

Recent evidence now indicates that “activation” of epithelial and possibly endothelial cells during the early initiation phase results in the up regulation of a variety of chemokines and cytokines (446, 522) that are instrumental in initiating the inflammatory cascade (See below). The cellular localization of these phenomena within the kidney is just beginning to be elucidated (131). As these and other events resulting in cellular alterations are not well identified morphologically, they have been difficult to evaluate using standard methodologies. Therefore, what role these early cellular responses play in further worsening renal perfusion remains to be determined.

The *extension phase* is ushered in by two major events: continued hypoxia following the initial ischemic event and an inflammatory response (Figure 1). Both events are more pronounced in the corticomedullary junction (CMJ), or outer medullary region, of the kidney. Documentation of severely reduced blood flow, stasis and accumulation of RBC and WBC has been historically noted (See Figure 2), however, the epithelial ramifications of these events have only recently been uncovered (329, 446). It is during this phase that renal vascular endothelial cell damage likely plays a key role in the continued ischemia of the renal tubular epithelium, as well as, the inflammatory response observed with ischemic ARF. During this phase, cells continue to undergo injury and death with both necrosis and apoptosis being present predominantly in the outer medulla (257). In contrast, the proximal tubule cells in the outer cortex, where blood flow has returned to near normal levels, actually undergo cellular repair and improve morphologically during this phase. As cellular injury

continues in the CMJ region during the extension phase, the GFR continues to fall. There is continued production and release of chemokines and cytokines that further enhance the inflammatory cascade (132). Interrupting the amplification of this inflammatory cascade may have therapeutic implications. Although the extension phase is probably the most promising phase for successful therapeutic intervention in ischemic ATN, there is a short window of opportunity. Based on animal models of renal ischemia, inflammatory cell infiltration in the outer medullary region of the kidney is significant by 24 hours following ischemia (260, 445, 628) and leukocytes may begin to appear in as early as 2 hours after ischemia (606).

During the clinical phase known as *maintenance*, cells undergo repair, migration, apoptosis and proliferation in an attempt to reestablish and maintain cellular and tubule integrity (Figure 1). The GFR is stable albeit at a level determined by the severity of the initial event. This cellular repair and reorganization phase results in slowly improving cellular function and sets the stage for improvement in organ function. Blood flow returns toward normal and epithelial cells establish intracellular and intercellular homeostasis. During the *recovery phase* cellular differentiation continues, epithelial polarity is reestablished and normal cellular and organ function returns (191, 402, 542). Thus, renal function can be directly related to the cycle of cell injury and recovery and this cell response to injury will be covered to greater extent later in this chapter.

II.2. Hemodynamic changes

Despite the role of tubular injury, the hallmark feature of AKI is a reduction in GFR, which by its essence implies an underlying impairment in hemodynamic regulation (114, 203, 210, 251, 354) and indeed was originally characterized as “vasomotor” nephropathy (210). Typically, the response is a sustained increase in renal vascular resistance (RVR) that may be attributable to a number of factors (62, 424, 495, 532). Renal blood flow is determined by a balance between factors that promote either vasoconstriction or vasodilation. The large number of potential contributors make it difficult to pinpoint a single cause for the increase in RVR. Rather, increased RVR is likely related to a number of variables: 1) alteration of intrinsic mechanisms of constriction; 2) enhanced production of vasoconstrictors or decreased production of vasodilators; 3) the altered response to vasoconstrictors or vasodilators.

II.2.1. Tubuloglomerular feedback and adenosine—Sublethal tubular damage results in impaired proximal sodium reabsorption (See Section III). This impaired tubular transport of sodium is expected to activate tubular glomerular feedback (TGF) mechanisms leading to increased afferent vascular resistance and a concomitant decrease in GFR. Studies by Blantz’s group in the 1980s using micropuncture techniques demonstrated that uranyl nitrate nephrotoxicity reduced SNGFR and that inhibition of the TGF-mechanisms by furosemide, which inhibits solute reabsorption in the TAL, improved SNGFR (434). This observation is not uniform among different models of injury as furosemide was not able to protect functional changes in SNGFR in ischemic kidney injury (53) and in general furosemide has not been demonstrated to be of clinical utility in altering the course of human AKI (206). However, these studies are complicated by the fact that furosemide has a number of potential kidney effects in AKI. Although additional studies in animals (7) and humans (6) have provided evidence for the importance of TGF in AKI, the extent that TGF contributes to diminished GFR in clinical AKI is not fully elucidated.

Adenosine, a potent renal vasoconstrictor resulting from the catabolism of adenine nucleotides, mediates the TGF response via the activity of A₁ receptors (528). In the setting of renal injury, adenosine activity may also lead to increased renal vasoconstriction. Some

studies support the idea that adenosine inhibition may prevent vasoconstriction and preserve injury in the setting of I/R. In rats, administration of the non-selective adenosine receptor antagonist theophylline preserved renal blood flow when administered before ischemic injury and increased renal blood flow after 5 days of recovery (326). In addition, theophylline was shown to preserve GFR in patients following exposure to a contrast agent relative to a placebo group (142).

Although encouraging, other studies from animal models raise questions about the nature of adenosine activity in the setting of injury. For example, transgenic mice lacking A1 receptors were unexpectedly more sensitive to ischemia reperfusion injury (528). In addition, increasing adenosine by inhibiting adenosine deaminase lessens the severity of ischemic injury (64). The protective effects may be associated with A2a receptor activity (409) and is likely mediated via effects on inflammatory cells (287). Thus, it appears that adenosine may be involved in multiple aspects of acute kidney injury and therapies geared toward targeting TGF and other hemodynamic responses still warrant additional consideration.

II.2.2. Sympathetic Nervous System, Norepinephrine and Ang II—Renal vascular resistance during AKI may be enhanced by increased sympathetic nervous activity.

Reductions in effective circulating plasma volume associated with hypotension would necessarily lead to an increase in activation of the sympathetic nervous system. Increase sympathetic activity is highly characteristic in rhabdomyolysis (67). Conger demonstrated that direct infusion of norepinephrine into the renal artery for periods of up to 60 minutes in rats could lead to prolonged reductions in RBF resulting in AKI (99). There are elevated levels of norepinephrine in the renal venous blood following ischemic renal injury (156). In addition, denervation of the kidneys improves RBF and attenuates the subsequent manifestation of acute tubular necrosis in ischemia and endotoxin models of AKI (156, 472, 588). Pharmacological blockade with clonidine, propranolol or guanethidine improves RBF in rat ischemia/reperfusion (I/R) injury (512). Recently, treatment with agmatine was shown to reduce SNS activity and ameliorated the resultant injury due to ischemia/reperfusion injury (527).

Increased sympathetic nerve activity is also a potent stimulus for renin secretion. Ang II, a potent renal vasoconstrictor, may also contribute to decreased blood flow in AKI. Patients with AKI have elevated plasma renin activity and hypertrophy of the granular cells of the juxtaglomerular apparatus (521). It is interesting to note that renal vasoconstriction to Ang II and NE are augmented by adenosine A1 activity (4). Despite this, there seems to be little evidence to support a role for Ang II activity in AKI; In fact, blockade of Ang II with either receptor blockers or ACE inhibitors increases the risk of AKI in patients following cardiac surgery (12).

II.2.3. Endothelin—Endothelin is a potent renal vasoconstrictor and produces profound reductions in GFR (149). Endothelin conveys its effects via at least two distinct receptor subtypes, the ET-A and ET-B receptors. Interestingly, activation of each of these subtypes has the potential to reduce GFR. In anesthetized rats, the selective ETA antagonist, A-127722 inhibited afferent arteriolar constriction, while specific activation of the ETB resulted in both afferent constriction as well as efferent vasodilation (226).

In humans, there is compelling evidence that the endothelin system may be activated in the setting of acute injury. Increased endothelin levels have been reported in patients with AKI due to severe sepsis (436). In a rat model of cold ischemic storage and transplant, pre-pro-endothelin mRNA was upregulated and immunoreactive ET-1 localized in the peritubular capillary endothelial cells following reperfusion injury (605). The role of endothelin in AKI

is supported mainly by pharmacological studies carried out in models demonstrating modest protective effects. For example, both non selective and ET-A specific antagonists have been shown to protect rats from the manifestation of renal I/R injury (83, 215, 277). Gellia et al, demonstrated a significant protective effect of ET-A blockade using an extreme model of IR injury in which 100% of control animals died within 3 days, while there was 75% survival over 2 weeks in animals treated with the ETA antagonist (165). However, the effect of ET antagonism in humans is less clear, as it did not manifest any protection prophylactically for contrast induced nephropathy (586).

II.2.4. Prostaglandins—Arachidonic acid released from phospholipids is converted by cyclooxygenase (COX-1 constitutive, COX-2 inducible) in the kidney to PGI₂, PGE₂, PGF₂, PGD₂, and thromboxane A₂. PGE₂ and PGF₂ are produced predominantly but not exclusively in the renal medulla, whereas degradative enzymes are present in both the cortex and medulla (192). During renal hypoperfusion, the intrarenal generation of vasodilatory prostaglandins, including prostacyclin (PGE₂) promotes the maintenance of renal perfusion.

Cox-2 inhibitors represent a major cause of drug-induced AKI and their use can increase risk for AKI following surgery or administration of radiocontrast agents (66, 70). Inhibitors of COX-2 exacerbate reductions in GFR and medullary blood flow in endotoxemia induced by LPS (590). Conversely, iloprost, the PGI analog, preserved RBF and corticomedullary hypoxia in a similar endotoxemic model (238). Iloprost can ameliorate the degree of renal insufficiency and renal blood flow in response to ischemia reperfusion in dogs (548). These effects are likely to be mediated by vasodilation, however, iloprost also decreases ICAM expression on the endothelium (224). In one study, prostacyclin protective effects were reported to be independent of effects on renal blood flow (148). Recently, some reports indicate that PG analogs may be a useful prophylaxis in preventing AKI clinically (518).

II.2.5. Platelet activated factor—Platelet activated factor (PAF: 1-O-alkyl-2-acetyl-sn-glyceryl phosphocholine) is a lipid mediator with a wide spectrum of biological activity. It can be produced by a number of different cell types including kidney cells. Its effects on renal function include reductions in GFR and RBF, making it a possible candidate for involvement in the development of acute kidney injury (337). Its hemodynamic effects occur by directly stimulating afferent arteriolar constriction as well as mesangial cell activity influencing K_f, and perhaps additional activity related to vascular obstruction (inflammation, *vide infra*) and cell toxicity (337). Acute renal injury may activate PAF via Ca⁺⁺-dependent phospholipase A₂; PLA₂ acts on membrane 1-O-alkyl-2-acylglycerophosphocholine to generate the precursor Lyso-PAF and subsequent conversion to PAF is mediated by an acyl transferase. Renal synthesis of PAF occurs following injury to the kidney in diverse models of AKI (e.g., ischemia, glycerol, gentamicin, endotoxemia) and treatment with PAF antagonists ameliorates the severity of AKI (337).

II.2.6. Alterations in vascular function secondary to injury—In addition to the induction of vasoactive factors, renal injury may directly damage the renal vasculature and alter its activity; such damage may influence vascular responsiveness, barrier function, coagulation cascades and/or inflammatory processes (74, 532, 610).

Studies by Conger and colleagues were among the first to highlight alterations in vascular responsiveness following AKI; these investigators demonstrated that post-ischemic rat kidneys do not autoregulate blood flow, and in fact, manifest vasoconstriction in response to decreased renal perfusion pressure. This occurred even one week following recovery from I/R when total renal blood flow had returned to baseline values (100). Morphological evidence of endothelial and smooth muscle injury was present up to 7 days following I/R

induced by renal artery clamping or norepinephrine infusion. The increased constrictor responses could be blocked by Ca⁺⁺ antagonists (101).

NOS3, (endothelial NOS) is produced at high levels particularly in the renal medulla (359) and its activity may represent an important mechanism to preserve medullary blood flow in response to renal vasoconstrictors such as Ang II (648). However, following renal injury, there is impaired endothelial NOS function, demonstrated by a loss of vasodilator responses to acetylcholine and bradykinin (101). A role for NOS3 in preservation of flow is suggested by the greater degree of renal damage that occurs in NOS3 knockout mice in response to endotoxin (589). Several studies using animal models have been utilized with the goal of increasing renal NO activity and/or preserving endothelial function. For example, administration of L-arginine, the NO donor molsidomine, or the eNOS cofactor tetrahydrobiopterin can preserve renal vascular perfusion and attenuate AKI induced by I/R (112, 161, 244, 483, 485, 525).

Recent studies have focused on rapid changes in endothelial function and their potential role in the initiation or maintenance of injury in the setting of I/R. Yamamoto et al., reported that renal perfusion in peritubular vessels was compromised within minutes of unclamping, characterized by sluggish and occasionally retrograde blood flow (610). Infusion of endothelial cells to athymic nude rats, or other non-endothelial cells harboring the NOS3 gene expression construct, protected against this early compromise in blood flow (74). Arrerio et al. propagated endothelial like cells from mesenchymal stem cells; these cells expressed markers typical of endothelial cells such as Tie-2, flt, flk and NOS3. Administration of these cells to control rats generated short-term engraftment into the vasculature and short term protection from I/R injury (13). Taken together, these data suggest that endothelial function, particularly via NOS3 activity, has protective effects on the severity of AKI.

Reactive oxygen species may influence the effects of vasoconstrictors and vasodilators and lead to an increase in renal vascular resistance. Increased renal superoxide production augments renal vasoconstrictor responses particularly in the renal medulla. Superoxide promotes vasoconstriction and enhances the reactivity of Ang II in the renal cortex and medulla (647). Oxidative stress in the kidney also enhances the vasoconstrictor effects of adenosine (91). Superoxide, by virtue of its known activity to convert NO to peroxynitrite, may block the normal homeostatic mechanism maintaining medullary perfusion and these responses may be normalized with the use of antioxidants (647).

II.2.7. Other factors affecting renal blood flow

II.2.7.1. Cell swelling and Edema: As described below in Section III, alterations in cell energetics leads to alterations in cell shape including swelling induced by enhanced sodium uptake. In addition to potentially affecting cell death, increased cell swelling may increase extravascular pressures on the peritubular capillaries and restrict flow. Flores et al demonstrated that when cellular swelling was reduced by infusion of mannitol, there was an improvement in renal blood flow (152). Tubular obstruction by cellular debris may influence hemodynamic response, In models of direct obstruction by ureteral clamping, there is rapid impairment of renal blood flow and this may be due alterations in tubular pressure and/or alterations in hemodynamic factors liberated secondary to injury (200, 222, 234, 500, 581).

I/R injury in rats alters cytoskeletal organization of small arterioles and endothelial cells (285, 532, 534). These alterations contribute to the breakdown of VE-cadherin in and the integrity of the tight junctions of peritubular capillaries as evidenced by leakiness of high molecular weight dextrans (>300,000 Da) into the interstitial space (534). Breakdown of barrier function may be the result of the activation of MMP-2 or 9 (533). The potential

ramifications of endothelial leakiness on the course of AKI are unclear. Increased edema resulting from leakiness may compromise renal perfusion by compressing peritubular capillaries and/or increasing hemoconcentration and exacerbating RBC trapping (201–203, 532).

II.2.7.2. Vascular injury and thrombosis: Although not well explored, it worth mentioning that endothelial damage in I/R injury may also promote thrombosis and fibrin deposition. The formation of microthrombi have been described in renal IR models as well as in renal transplant biopsies (141, 532). Evidence that endothelial cell damage promotes thrombus formation in renal I/R is minimal and its potential role is suggested mostly from work in other organ systems. Potential microthrombus formation may be associated with loss of surface inhibitory factors, notably thrombomodulin, which allows fibrin formation by disinhibiting thrombin (532). Recently, a soluble thrombomodulin molecule was shown to attenuate the degree of renal injury following ischemia in rats; this was associated with an rapid effect on preservation of perfusion within renal capillaries and decreased leukocyte adhesion (492).

In addition, tissue factor, a membranous glycoprotein that functions as a receptor for the coagulation system and is normally not present on quiescent endothelial cells, was shown to be elevated in peritubular capillary endothelial cells following renal I/R injury. Inhibition by antisense oligonucleotides manifested a significant protection indicated by preservation of overall morphology and mortality in rats (357).

Endothelial cells may also modulate the injury process via binding of activated serum protein C, to the endothelial protein C receptor. Activated protein C exerts both antithrombotic and cytoprotective properties, the latter via modulation of protease-activated receptor-1 (PAR-1) signaling (240). In a recent study, a mutant APC molecule that retained PAR agonist function preserved renal function in AKI, but a mutant APC that retained only anti thrombogenic capacity did not protect against LPS injury (492). In clinical trial of human sepsis patients, Drotrecogin (a recombinant human activated protein C) significantly decreased 28 d mortality over the placebo group(48).

II.2.8. Lack of vasodilator response and role of infiltrating cells and renal injury—The data on animal models suggest that vasoconstriction contributes to the early loss of GFR and triggers injury. While reductions in RBF may represent an important feature in the initiation phase of AKI, increasing RBF after established injury may not resolve GFR. Figure 3 illustrates proposed pathways in which factors independent of direct vasoconstriction may ultimately impact the loss of renal blood flow and GFR in AKI

Numerous examples in the literature demonstrate a lack of effect of vasoactive factors at correcting established AKI. Perhaps the most well known example of this is represented by the use of “renal-dose” dopamine. At low doses (i.e., 3 µg/kg/min) dopamine is vasodilatory and promotes diuresis primarily through the D-1 receptor and has beneficial effects in multiple models of AKI (121). Despite this, multiple studies have consistently demonstrated that renal-dose dopamine or the dopamine agonist fenoldapam, do not influence outcome of AKI in patients despite producing diuresis (45, 254, 255, 288, 560). Similarly, atrial natriuretic peptide can produce diuresis in AKI patients but does not provide any improvement in renal function or recovery (305).

Characteristic of renal injury is the prominent infiltration of leukocytes, primarily neutrophils occurring fairly rapidly following ischemia/reperfusion injury in rodents (160, 375, 628). Leukocyte adhesion to activated endothelial cells following I/R, and subsequent red blood cell congestion results in a further compromise renal function by influencing renal

hemodynamics. Leukocyte adhesion is proposed to impair renal function in AKI in the following ways: 1) Leukocytes may physically impair capillary flow, particularly in the renal medulla and exacerbate hypoxia in this region; 2) Leukocytes may generate molecules exacerbating vasoconstriction, such as ROS and/or vasoactive lipids; 3) Leukocytes may contribute to parenchymal cell injury, the result of which may be to increase tubular lumen pressure and reduce GFR. Leukocyte induced interstitial edema can increase interstitial pressures and retard peritubular capillary flow. These activities may represent a switch to the extension phases of AKI and be the cause of resistance to vasodilator therapy after the initiation of AKI. To our knowledge, there have been no studies correlating the adhesion of leukocytes with reductions in medullary blood flow and subsequent resistance to vasodilator therapy.

II.3. Inflammatory process in AKI

In addition to the myriad of altered vascular functions that may influence the development of AKI, inflammation is mediated in part by the adhesion of leukocytes to affected endothelial cells. Inflammatory cell infiltration is a prominent early feature in the outer medullary vasa recta capillaries following injury. There is an increase surface expression of leukocyte adhesion molecules ICAM-1 and P and E-selectin on endothelial cells in response to injury (158, 261, 383). *In vivo* imaging studies have shown leukocyte adhesiveness to the peritubular capillary wall occurs within hours of reperfusion (259, 492). Treatments geared toward reducing endothelial/leukocyte interactions by targeting these endothelial adhesion molecules preserve blood flow and protects against renal damage in IR induced AKI (158, 261, 383). The endothelium may also be the source of chemoattractant factors, such as fractalkine (CX₃CL1), which is expressed following renal injury and may promote infiltration of macrophages (see below).

On balance, most studies in animal models confirm an important role of inflammation in the establishment of AKI. Studies geared to prevent neutrophil infiltration by interfering with P-Selectin, ICAM/CD18 pathways, mitigate the degree of renal damage in ischemia reperfusion injury (160, 262, 395, 443–445). Both anti-oxidant therapy and minocycline treatment reduced the extent of leukocyte infiltration into post-ischemic kidneys, an effect that was partially attributable to inhibition of ICAM-1 expression (259, 330). Neutrophil extravasation has been shown to be dependent on the activity of the adenosine A_{2A} receptor which minimizes the degree of parenchymal injury (57). In human AKI, ICAM antibody pretreatment protected against delayed graft function in transplant recipients relative to a group of recipients receiving the contralateral kidneys from the same donors. Interestingly, immunoneutralization following transplant did not affect the development of DGF (155, 195, 473). Thus, this treatment may prevent injury, but may be ineffective after initiation of injury.

The degree to which neutrophils actually contribute to renal injury remains somewhat unclear. Neutrophils are not prominent in biopsies of patients with acute kidney injury (155). There are conflicting reports regarding the effect of neutrophil depletion on I/R induced AKI with some studies demonstrating a protective effect while others demonstrating no effect (143, 262, 444). DeBroe's group suggested that the degree of neutrophil accumulation following injury may be overestimated, since traditional neutrophil markers (MPO and chloroacetate esterase) cross react with macrophages (628).

II.3.1. T cells—Recent evidence suggests that monocytes, primarily T-cells, may play an important role in AKI. T-cells are apparent in the medullary vasa recta in animal models of AKI and are present in biopsies of patients with AKI (Figure 4) (155, 627). It is noteworthy that blockade of CD18a/ICAM pathway, thought to be affecting neutrophils, also affects the

deposition of T-cells (78). Evidence supporting an important role for T-cells derives from studies in which T-cells are depleted using either antibody neutralizing approaches (624) or genetic approaches (117) reducing the degree of renal injury to I/R. De Greef and colleagues further demonstrated the importance of the early expression endothelial B7-1 in peritubular capillary endothelial cells on the adherence of T cells and the manifestation of injury following I/R (Figure 4) (118, 537). It remains unclear how T cell deposition contributes to the injury process. However, evidence suggests that there is an important role for T-cell derived IFN- γ since reconstitution studies demonstrate that CD4⁺ T cells derived IFN- γ null mice are not sufficient to recapitulate the injury observed in T-cell replete animals (117).

Conversely, regulatory T cells (Tregs, CD4, CD25, Foxp3) are lymphocytes with immunosuppressive properties that may traffic to areas of inflammation and mitigate immune reactions. Depletion of Tregs with anti CD25 mAb potentiates kidney damage induced by IRI, and is associated with increased neutrophils, macrophage and cytokine responses. When adoptive transfer of T cells from Foxp3 null mice was carried out, renal injury is worsened, while adoptive transfer of Treg alone into T cell deficient mice results in significant protection (274).

A potential problem in understanding the pathological role of T cells in the mediation of AKI is that CD4⁺ cells are classically thought to play a role in antigen specific adaptive immunity that requires 2–4 days for processing. Such activity is inconsistent with a role for these cells in AKI. However, natural killer T (NKT) cells, which express the TCR (T cell receptor) and the marker NK1.1 represent a unique and small subset of CD4⁺ cells. These cells can produce large amounts of cytokines such as IFN γ and TNF (Th1 type), or IL-4 and IL13 (Th2 type). NKT producing IFN γ are present in the post-IR kidney within 3 hours of injury (311). Blockade of NKT cell activation with the anti-CD1d mAb, NKT cell depletion with an anti-NK1.1 mAb in wild-type mice, or use of iNKT cell deficient mice (J α 18^{-/-}) inhibit the accumulation of IFN- γ -producing neutrophils after IRI and prevented AKI (311), suggesting that NKT cells are the major early acting CD4⁺ population acting affecting kidney injury following I/R.

II.3.2. Macrophages—Macrophages are derived from monocytes and have phagocytic capabilities. Macrophages infiltrate the injured kidney within 1 hour of ischemia reperfusion; this activity is mediated by fractalkine (CX₃CL1), which is expressed by injured endothelial cells and is a potent chemoattractant and adhesion molecule for (CX₃CL1 receptors) CX₃CR1 and CCR2 expressed on monocytes and macrophages. Both ischemia and cisplatin AKI induced fractalkine expression on peritubular capillary endothelial cells. Blockade of this pathway with CX3C receptor-1 antibody effectively reduces the severity AKI in mice while adoptive transfer of activated RAW 264.7 macrophages restores sensitivity to injury (312, 407). Macrophages lacking CCR do not infiltrate injured kidneys and the resultant injury is less severe (312). Infiltrating macrophages may exacerbate injury by producing a high levels of the cytokines IL-1a, IL-6, IL 12 and TNF- α and promoting further inflammation (272).

The last several years has seen an increased awareness of the diversity of macrophage phenotypes. Within the setting of kidney injury, macrophages are most abundant during the repair response suggesting a beneficial role for macrophages in recovery. A distinct population of macrophages, referred to as M2, secrete anti inflammatory compounds such as IL-10 and TGF- β and may be involved in tissue repair by secreting potential trophic growth and angiogenic factors (136, 591). Proof of principal for this activity was demonstrated by Want et al., in which ex vivo programming of macrophages toward an M2 phenotype ameliorates chronic renal inflammation (593). Jang *et al.* found impaired recovery from kidney IRI when macrophages were depleted following kidney IRI using liposomal

clodronate (231). Administration of macrophages during the repair phase of ischemic injury increased renal cell proliferation and enhanced recovery (577). In one study, macrophages were ablated using a transgenic approach in which the diphtheria toxin (DT) receptor was expressed on CD11b-positive cells. DT treatment of these mice reduced macrophages and resulted in prolonged and less successful recovery from AKI (327). The reparative potential of macrophages may be mediated by Wnt ligands expressed on the surface of regenerative macrophages, and interact with epithelial cells expressing Wnt-receptors (327).

II.3.3. Dendritic cells—CD11c⁺ MHC class II⁺ DCs are the most abundant leukocyte subset in the normal mouse kidney suggesting an important role in renal immunity and inflammation. When activated due to injury, these cells may release pro-inflammatory factors and interact with NKT cells. However, the role of dendritic cells in AKI is not yet clear. While dendritic cell activation has been suggested to lead TNF- α production in the setting of ischemic AKI (129), there have been few functional studies on dendritic cell activity in the injury process. In one recent study, ablation of CD11c DC using a DT mediated approach, increased in sensitivity to cisplatin induced injury, suggesting that DC have a protective role via an unknown mechanism (536).

II.3.4. Inflammatory signals in AKI—In combination with endothelial adhesiveness, inflammatory mediators are synthesized and released by both tubular epithelial cells, and activated leukocytes. Tubular epithelial cells produce TNF- α , IL-1, IL-6, IL-8, TGF- β , MCP-1, ENA78, RANTES an fractalkines, while leukocytes may produce IL-1, IL8, MCP-1, reactive oxygen species and eicosanoids. These factors act in concert to promote inflammation in a positive feedback loop promoting further kidney injury (63). The control of inflammation and the various factors that regulate it is substantial and the subject of numerous reviews. A limited consideration of some of the major pathways is described below.

For example, IL1 β is a chemoattractant that can recruit leukocytes to areas of injury; pro IL1 β is processed by caspase 1 (ICE) to form mature IL1 β . IL1 β levels are increased in mice following ischemia reperfusion injury (262). Based on studies using receptor antagonists, or IL-1 β null mice, the blockade of IL-1 β reduces the extent of neutrophil infiltration following ischemic injury, but has no effect on the resultant loss of renal function (190). Caspase-1 is also involved in the activation of IL18. IL-18 is enhanced in AKI and may represent an early biomarker of acute injury (77). Conversion to mature IL-18 is not observed in *caspase-1*^{-/-} mice, relative to wild type controls, and caspase 1^{-/-} are protected from ischemic injury. Similarly, the caspase inhibitor Quinoline-Val-Asp(Ome)-CH₂-OPH (OPH-001) also prevented IL-18 processing while IL-18 immunoneutralizing antibodies preserved renal function following I/R (365, 366).

IL-6 may also promote inflammation in AKI; injured proximal tubules produce molecules that activate the expression of IL-6 expression on infiltrating macrophages and IL6 is found at elevated levels in the serum of mice following renal injury (266). IL-6 knockout mice are resistant to AKI, but sensitivity to injury is restored by adoptive transfer of wild-type macrophages (266). However, there are mixed data on the importance of IL6 in AKI, as IL-6 knockout mice are more sensitive to injury induced by cisplatin, possibly via an increase in pro-apoptotic regulatory signals (374).

Toll like receptors (TLR) are a family of pattern recognition receptors that detect motifs on pathogens and host material important in the activation of innate immunity. Located primarily on immune cells, TLR signaling transduces microbe host interactions into an inflammatory state which is required to eliminate invading organisms; when uncontrolled TLR signaling can lead to tissue damage (5). TLRs mediate their pro-inflammatory

responses by a common downstream signaling molecule, MyD88, which leads to the activation of NF κ -B inflammatory signaling.

TLR 4 is the classic endotoxin receptor. C3H/HeJ mice lack the TLR4 receptor, while C3H/JeOuJ mice with intact TLR4 receptors manifest an increased response to endotoxin with greater production of the pro-inflammatory cytokine TNF- α , more pronounced renal damage and neutrophil infiltration when compared to C3H/HeJ (108). The influence of specific TLR may be dependent on the model of sepsis. Using a cecal ligation puncture (CLP) model of sepsis, TLR4 and TLR2 deficient mice were not protected against renal damage and inflammation, while MyD88 knockout mice are resistant (120). TLR 9 may mediate CLP induced renal inflammation because severity of AKI is reduced in both TLR9 knockout mice and following the administration of chloroquine, an antagonist of TLR 3, 7,8, and 9(619).

TLRs also interact with non microbial endogenous substances, which has expanded this field beyond sepsis into conditions of autoimmunity and inflammation (457). For example, HMBG-1 proteins, potentially released from damaged cells, can activate inflammatory signaling in human renal proximal tubules and endothelial cells from wild type mice, but not from TLR 4 knockout mice (90, 280). There is a blunted induction of cytokines and chemokines in TLR4 and TLR2 null mice, as well as reduced neutrophil and macrophage infiltration in response to ischemia and other models of injury, such as cisplatin (113, 295, 609, 638). TLR4 expression is found on vasa recta capillaries within 4 hours of ischemia reperfusion injury; the expression of adhesion molecules CD54 and CD62E in peritubular capillaries was not observed in TLR4 null mice compared to wild type. Secondary increases in TLR4 on proximal tubule cells occur within 24 hours of injury and may amplify the pro-inflammatory state of I/R injury (90). Recent evidence suggests that the TLR 4 pathway may be important in human AKI in the setting of transplant. Donor kidneys with a TLR4 loss-of-function allele contained less TNF α , less MCP-1, more heme oxygenase 1 (HO-1), and exhibited a higher rate of immediate graft function (280).

Another pro-inflammatory regulatory system that can influence acute kidney injury is the complement (C) system. The C system contains three activation pathways that participate in innate immunity (441). Recent studies have suggested that the alternate complement pathway may be active in the setting of acute kidney injury by ischemia reperfusion (643). The complement system is proposed to represent a primary regulator of neutrophil infiltration. Mice deficient in complement factor B (fB $^{-/-}$), an essential component of the alternative pathway, develop substantially less functional and morphologic injury, less tubulointerstitial complement C3 deposition, and less neutrophil infiltration in the outer medulla after I/R, relative to wild type controls (546). Complement processing in AKI may be controlled by the complement inhibitor Crry, which redistributes from the basolateral membrane in response to I/R (547). Evidence in support of this activity derives from studies in which Crry deficient mice demonstrate an increased sensitivity to IR injury (547). Surprisingly, despite a significant increase in C3 generation following endotoxemic (LPS-induced) kidney injury, blockade of the complement cascade does not protect against kidney injury in this model (107). Both C5a and C5b-9 contribute to renal injury independent of neutrophil attraction, perhaps by induction of apoptosis and impairing recovery responses following IR (441, 488). The activation of complement system is required for the production of macrophage inflammatory factor -2 and keratinocyte derived chemokine (KC) in renal tubule cells following I/R in an NF- κ B dependent fashion (545).

In summary, the general observations described above suggest a complex interplay with various pro-inflammatory signals. The sources of these various signals and how they are regulated at the molecular level are only beginning to be understood. For example, the regulation of pro-inflammatory genes may be secondary to the activation of upstream

transcriptional activators of inflammatory genes, such as interferon regulatory factor-1 (IRF-1). In proximal tubule cells, IRF-1 expression is controlled by ROS activity *in vitro*, and it is produced rapidly by outer medullary tubular epithelial cells following renal injury. Null mutations in the IRF-1 gene reduce the degree of inflammation and pro-inflammatory gene expression following ischemic AKI (592). As mentioned, initial increases in endothelial TLR4 expression may be required for subsequent adjacent tubular TLR4 activity serving to amplify pro-inflammatory mechanisms. Future studies will likely be geared toward deciphering the complex interacting network of pro-inflammatory interactions that may aid in developing potential therapeutic targets.

II.4 What is the link between tubular injury and GFR?

A loss of GFR necessarily indicates that Starling forces are impaired during the injury process that results in renal dysfunction. Although the above sections describe a number of parameters that have been proposed (vasoconstriction, tubular congestion, edema, inflammation), no single factor has been demonstrated to be responsible for the entirety of the reduced GFR in response to tubular injury. Rather, the more likely interpretation from the multitude of these studies is the following; 1) Early vasoconstriction observed in many models impairs blood flow and GFR and is likely mediated by a number of redundant pathways. As such, blockade of any one pathway is not likely to produce significant benefits. 2) Features secondary to inflammation or structural changes in the renal parenchyma likely sustain reductions in GFR. At this point, hemodynamic therapies may serve a more limited role and the therapies should be directed toward blocking inflammatory pathways and/or restoring structure. 3) A given therapy may be effective in some settings, but depends on the timing. Therefore, a greater understanding of the phases of injury and methods to evaluate such phases will play an important role in designing effective treatments.

II.5. Remote effects of AKI and possible causes for mortality

In addition to altering kidney function, renal tubular damage may set off a series of events that alter physiological responses in distant organs and that may play a significant role in mortality. AKI increases the risk of death in patients with other co-morbid conditions and death *per se* may not be due to loss of renal function (237). Experimental models of renal ischemia reperfusion lead to increased pulmonary vascular permeability and macrophage infiltration (279). Rabb's group has also shown that renal I/R increases the permeability of the blood brain barrier (335), induces histological evidence of damage in neurons and microglial cells, and increases levels of pro-inflammatory cytokines such as G-CSF in the cerebral cortex and corpus collosum. Post AKI mice performed poorly in an open field locomotor tests to evaluate brain function (335). With regard to cardiovascular function, Kelly et al demonstrated decreased cardiac contractility and cardiac myocyte apoptosis in mice following I/R injury (256).

Based on these observations a developing view is that kidney injury sets in motion a complex systemic inflammatory response that is the basis for organ cross talk and it is likely that liberated pro-inflammatory cytokines (177). In kidney ischemia models in mice, there is evidence for the distinct activation of various circulating cytokines including KC, G-CSF, IL-6, IL-1 β and IL-12 (177). An important potential area of future investigation will be to determine if treating these cytokine responses influence renal as well as the function of distant organs and influence mortality.

The idea that loss of GFR *per se* is not the single critical feature of mortality risk in AKI was addressed by Humes and colleagues. These investigators pioneered the bioartificial kidney which consists of a hemofilter in a continuous venovenous hemofiltration circuit (CVVH)

with a cartridge containing renal proximal tubule cells, also called the Renal Tubule Assist Device (RAD). In nephrectomized dogs treated with endotoxin to simulate sepsis, the RAD helped to reduce overall inflammatory cytokine levels and improved cardiovascular parameters relative to dogs that were similarly dialyzed but without the RAD (150). Consequently, therapeutic strategies that target solute clearance, as well as, remote mediators of inflammation may hold future promise for addressing the full clinical impact of AKI.

III. Cellular response to injury

III.1. Energetics and structure

As described above, AKI is characterized not only by cell death, but also by a large spectrum of cellular injury (573), which may lead to the initiation of hemodynamic and inflammatory pathways reducing kidney function (Figure 2, 3 and 4). Therefore, an understanding of the cellular responses leading to both lethal and sublethal injury represents an important element in understanding the underlying pathophysiology of AKI. The varying sensitivity of renal cells to injury results from physiological parameters including the ratio of the energetic substrate to metabolic demand, the physical location of the cells within the kidney, the degree of regional blood perfusion in the aftermath of the insult, oxygenation status and membrane permeability.

III.1.1. Energetics and sensitivity to injury—In most animal models of AKI, the proximal tubule is considered the most sensitive cellular entity to ischemic, hypoxic or nephrotoxic damage (62, 68). However, the thick ascending limb of Henle is also highly metabolically active and is dramatically affected in certain models of AKI (61). In a rat model of ischemia reperfusion, early morphological alterations (such as tubular dilatation and loss of renal microvilli) are observed in all proximal tubule segments within hours of the initiation of reperfusion (489, 573). Proximal tubule cells within the S1 and S2 segment manifest a largely reversible injury; if cell death occurs, it localized primarily in the S3 segment (573).

Proximal tubule sensitivity relates largely to high metabolic rate and a strong dependence on oxidative phosphorylation. The intact mammalian kidney reabsorbs nearly 80 meq Na/g kidney/day across the renal tubules and accounts for nearly 70% of oxygen utilization by the kidney (306). To meet this demand, tubule cells generate a significant amount of ATP. The preferential energy substrate is non-esterified free fatty acids, primarily palmitate, and to a lesser degree, lactate, citrate and pyruvate (306). Proximal tubules do not utilize glucose, but rather are gluconeogenic (306).

Because of its limited glycolytic capacity, renal ATP concentrations fall to ~20% of control values within 10 min of renal artery occlusion (265, 578). It is interesting that cortical values drop much more dramatically than medullary levels reflecting the higher proportion of proximal tubule cells in the cortex. In addition, a proportion of medullary cells are capable of anaerobic glycolysis. Medullary collecting duct cells appear largely resistant to ischemia and these cells can maintain ATP levels at approximately 60% of control values when oxidative phosphorylation is blocked and can even increase their glycolytic capacity under stress conditions (24). In contrast, both the medullary thick ascending limb and S3 proximal tubule cells have a limited glycolytic capacity and are unable to sustain normal ATP levels in the face of oxidative stress (42, 561).

Hypoxia, ischemia/reperfusion and nephrotoxic agents can have rapid and sustained effects on cellular ATP levels in vivo or in vitro. (241, 281, 355, 401, 524). Using both standard biochemical approaches and ³¹P NMR, Stromski et al., demonstrated a rapid reduction of

renal content of ATP within minutes of the onset of ischemia (524). The restoration of ATP is biphasic upon reperfusion with a rapid initial phase and a more prolonged phase up to 2 hours. If ischemia is short (i.e., ~ 15 min or less), ATP levels quickly return to control values. However if the ischemic time is extended (e.g., > 30 min), tissue ATP recovers only partially within 2 hours. There is a significant negative correlation of the 2 hr ATP level and the resultant degree of renal function at 24 hours (524). Similarly, in a model of hypotension induced AKI, the early recovery of ATP was impaired in animals that subsequently developed kidney damage, while animals that manifested rapid restoration of ATP were spared from significant injury (454). These observations suggest a tight link between tubular energetics and a transition from pre-renal azotemia to intrinsic renal failure.

Both ATP depletion and associated cell damage are readily reversible provided substrate can be replenished and the machinery for ATP generation has not been dramatically compromised. In a series of studies from Siegel's group, ATP-MgCl₂ administration to post-ischemic rats accelerated the recovery of tissue ATP that was associated with the preservation of GFR, renal blood flow, and several indices of cellular injury (162–164, 501, 503). Supplementation with inosine, an ATP substrate precursor, and guanine analogs to maintain GTP levels in both in vivo and in vitro models of injury have been shown to have a modest degree of renal protection (146, 258).

III.1.2. Loss of structure with impaired energy status—As described above, one of the earliest morphological features in acute injury is the rapid loss of the renal brush border, which occurs within 15 min of reperfusion and is due to the breakdown of cytoskeletal elements (573). The cytoskeletal response is attributed to the sensitivity of actin polymerization on cellular ATP levels. The reduction in cellular ATP results in a loss of (filamentous) f-actin from apical microvilli, terminal web junctional complexes and stress fibers (253, 377, 378). This activity also disrupts cytoskeletal association of protein complexes mediating cell-cell interaction as well as cell substrate interactions, which contribute to the loss of cell polarity, the mixing apical and basolateral transporters and loss of tight junctions (377).

The dynamic processes of actin polymerization/depolymerization become dysregulated when ATP levels are reduced. There is a counter intuitive increase total cellular (filamentous) f-actin formation and a decrease in monomeric g-actin content that occurs within 5 minutes of ATP depletion resulting in the formation of punctuate f-actin structures in the perinuclear region (382). Under physiological conditions, ATP-G-actin monomers, via their association with a class of cytosolic proteins known as thymosins, are prevented from polymerizing with established actin filaments or other G-actin monomers. In the setting of ATP depletion, thymosin dissociates from G-actin monomers, predisposing unregulated f-actin formation (18).

Ashworth and colleagues suggested that the disruption of the actin network is mediated by cofilin/actin depolymerizing factor (ADF). ADF is dephosphorylated and localizes from a diffuse cytosolic location to the apical microvilli and submembrane spaces within minutes of renal ischemia (490) (Figure 5). ADF has been localized in brush border membrane vesicles sloughed into the tubular lumen (15) and can be found in microvesicles in the urine samples from patients with AKI (376). The dependence of ADF activity on the stability of the actin network was demonstrated by studies in which a constitutively active form of ADF disrupted cortical f-actin structures in kidney cells, while cells containing only an inactive mutant of ADF/cofilin did not manifest breakdown of the cortical f-actin network in response to ATP depletion (15).

ADF/cofilin activity is regulated by the Rho GTPase, RhoA. RhoA is a member member of the Ras superfamily of p21 G-proteins that are regulated by guanine nucleotide binding. When GTP is bound to RhoA it is active, but intrinsic GTPase activity results in an inactive GDP-bound form (52). GTP levels fall in parallel to ATP in the setting of ischemic injury or in renal cells in response antimycin A and this decreases abundance of the GTP-bound form of RhoA (186). Constitutively active RhoA mutant proteins (RhoV14) preserves cortical f-actin networks and cytoskeletal ezrin activity. Inactivation of Rho with the bacterial toxin C3 mimics ischemia, results in stress fiber reduction and an impairs recovery of the cortical actin network with the re-establishment of ATP levels (186).

Alterations in cytoskeletal structure lead to a loss of epithelial polarity and barrier function. Cell polarity is established in part by the function of tight junctions and adherens junctions, both of which have a similar architectural network consisting of transmembrane bridging proteins and cytoplasmic proteins linking to the actin cortical network (290). Not surprisingly, ATP depletion alters the cellular distribution of TJ proteins occludin ZO-1, ZO-2 and cingulin (21, 80, 176, 558). In cultured cells and in response to ischemia, there is a degradation of E-cadherin and a disruption of its interaction with catenin both in cultured cells and in ischemic kidney of rat (348). The loss of these junctional structures allows transmembrane proteins with restricted domains to move freely to other membrane spaces. The Na-K ATPase moves from a Triton X-100 insoluble fraction to a Triton X soluble fraction following ischemia in vivo or ATP depletion in vitro. This event is indicative of this protein's release from a restricted basolateral localization, which contributes to the loss of vectorial transport of Na in the proximal tubule (379, 380). The re-localization of Na-K ATPase can occur with repletion of ATP by a recycling of Na-K-ATPase subunits (569).

Alterations in cell-cell junctions result in decreased in transepithelial electrical resistance suggesting an increased leakiness in the paracellular pathway (348). This reduced transepithelial resistance is associated with increased backleak of filtrate and was suggested to occur in vivo by studies in which ^3H - inulin microinjected into the lumen of post-ischemic proximal tubules was subsequently recovered in the urine formed by the contralateral kidney (133). Similar results have been described in human transplant patients. In a study by Kwon et al., the fractional excretion of graded molecular weight dextrans was increased in patients with sustained AKI, indicative of backleak of inulin during the clearance procedure (284).

Proximal tubule cells require attachment to the basement membrane for proper integrity and function. Cell-matrix adhesions are mediated by the integrin family of cell adhesion receptors, multi-subunit proteins that span plasma membrane and mediate substrate attachment at focal adhesions (FAs)(187, 649). β 1-Integrin is the most prominent integrin mediating PTC-substrate adhesion and is lost from the basolateral membrane region during the ischemic period (173, 649). Goligorsky et al., have proposed that the loss of the basolateral expression of integrin receptors is responsible for the exfoliation of viable proximal epithelial cells and that the redistribution of integrin receptors from the basolateral to the apical surface of epithelial cells facilitates self-adhesion and clumping of detached cells in situ (173). These two processes culminate in tubular obstruction and may severely increase tubular pressures and impair fluid flow. This hypothesis was supported by a series of studies in which cyclized RGD peptides inhibiting β 1 integrin binding mitigated the formation of obstructive luminal cells, decreased intratubular pressures, and improvement in serum creatinine following renal ischemia reperfusion (400, 459).

III.1.3. Impaired renal energetics and the role of reactive oxygen species—

Replenishment of energetic substrates can resolve the early alterations in cell morphology, but irreversible cell damage may ensue if not adequately restored. Impaired energetics

represents a potential link toward intrinsic tissue damage and results from various inter-related features of metabolism.

1. **Substrate availability:** Prolonged ischemia/hypoxia may impair the cellular substrate pool for the re-synthesis of ATP, as originally proposed by Fernando et al. (146). When ATP is degraded, ADP and AMP are formed. The activity of 5' nucleotidase with AMP forms adenosine, inosine and hypoxanthine. Although the nucleotides (ATP ADP AMP and IMP) have low permeability to the cell membrane, the nucleosides adenosine and inosine, and hypoxanthine are permeable and can readily diffuse out of the cells (599). Furthermore, hypoxanthine can be converted by xanthine oxidase to uric acid. Thus, as a result of either diffusion out of the cells or metabolism, there is a decrease in the substrate pool for ATP synthesis (599).
2. **Renal hemodynamics:** The renal outer medulla sits on the “brink of hypoxia” with pO₂ values between 10–20 mm Hg (compared to 50–60 mm Hg in the cortex). With regard to energetics, impaired renal medullary blood flow can further exacerbate hypoxia in the early phases of ischemia reperfusion in rodents (70). In humans, hypoxia is worsened in delayed vs. immediate function grafts following transplant as demonstrated using blood oxygen level dependent (BOLD) MRI (126). The sustained hypoxia may inhibit the resynthesis of ATP accounting for a greater degree of injury in this region.
3. **Impaired mitochondrial activity** plays a significant role in the injury process and has implications regarding energetics, activation of cell death pathways and the genesis of reactive oxygen species. The effect of mitochondrial function in the setting of injury has been a central focus of not only acute kidney injury, but also injury in myocardium, brain, liver and other organs susceptible ischemic and toxic insult (44, 299, 415).

Increased mitochondrial cell swelling is a common morphological feature of early response to hypoxia, ischemia/reperfusion and nephrotoxins, *in vitro* or *in vivo*. For example, mitochondrial swelling is evident by electron microscopy within 1–2 hours of ischemia reperfusion injury in both proximal convoluted and proximal straight tubules, and the increased severity of ischemia was associated with the severity of mitochondrial swelling (169–171). Mitochondrial swelling is also observed in response to gentamicin treatment (506) and mercuric chloride. *In vitro*, mitochondria of isolated proximal tubules swell in response to hypoxia/reoxygenation (491). Human kidneys subjected to anoxia during cold storage manifest mitochondrial swelling, release cytochrome C and demonstrate bax translocation consistent with the activation of apoptotic pathways (see below; (471)). Thus, early alterations in mitochondrial structure represent an important early event in AKI.

III.1.3.1. Respiration and Mitochondrial Permeability Transition: Impaired mitochondrial respiratory capacity appears tightly linked to altered mitochondrial structure. Studies using isolated mitochondria from kidney or cultured kidney cells report reduced ATP synthesis, and respiratory capacity secondary to ischemia, cisplatin (69), lead administration (154) or gentamicin treatment (308, 506). Using both *in vivo* models or preparations of isolated tubules, state II, state III and 2,4 dinitrophenol uncoupled respiration are reduced within 1–2 hours of hypoxia or administration of HgCl₂ (126, 159, 164). Mitochondrial functional deficits also develop due to the loss of mitochondrial membrane potential (144) and the impaired ability to utilize complex I substrates (602).

The connection between swelling and impaired energetic function may result from an increase in the permeability of the inner mitochondrial membrane (178). Mitochondrial permeability transition (MPT) refers to the process by which the mitochondria lose the

ability to maintain the H^+ ion gradient necessary for the production of ATP via the F_1F_0 -ATPase. Substantial evidence suggests that the permeability is due to the opening of a non-specific pore with a high conductance of molecules less than $M_w \sim 1200$ (105, 267). The result is the loss of the mitochondrial transmembrane potential $\Delta\psi$ accounting for the de-energization of mitochondria and impaired ATP synthesis. The increased permeability can result in the influx of ions such as K^+ down its concentration gradient, which is thought to contribute to the observed swelling (105, 267).

MPT is commonly studied by evaluating swelling using changes in light absorbance and this approach has been used to identify factors that activate MPT. Most notably Ca^{++} can activate MPT in isolated mitochondria from liver hepatocytes (300) or in permeabilized renal tubule cells at concentrations as low as 35–45 μM (145). MPT can also be stimulated by reactive oxygen species by addition of hydrogen peroxide (538), tert-butyl hydroperoxide (268) or xanthine/xanthine oxidase (344). Other potentially important signals associated with renal injury may participate in altering mitochondrial potential. Phospholipase A2 (PLA2) participates in MPT activation suggesting a role for fatty acid (273). Interestingly, Feldkamp et al. demonstrated that delipidated BSA, which can bind free fatty acids in permeabilized tubules, can protect against MPT in response to hypoxia (145).

Permeability transition is thought to be mediated by the opening of the MPT pore, proposed to comprise the adenine nucleotide transporter (ANT) and the voltage dependent anion channel (VDAC) normally associated in the outer membrane. An in depth summary of the MPT pore is outside the scope this discussion and the reader is encouraged to see recent review articles on its biochemical properties (105, 267). Evidence for a specific MPT pore as a physiologically regulated entity derives in part from the discovery that cyclosporine A inhibits MPT at low and saturable concentrations (299). Cyclosporine reduces ischemic myocardial and hepatic injury (11, 299), and it also reduces MPT in rabbit renal tubules exposed to hypoxia (144). However, cyclosporine is not useful in AKI due to its vasoconstrictor effects which likely contribute to worsened kidney injury following ischemia (49). Because cyclosporine binds cyclophilin D (CypD), it has been hypothesized that cyclophilin D, may be a part of and/or regulate the MPT pore complex (397). Neurons from cyclophilin D knockout mice, have a reduced level of MPT in response to oxidant stress when compared to neurons from wild type animals (478). The importance of cyclophilin D in MPT in kidney injury was recently highlighted by Devalaraja-Narashimha et al., who reported that cyclophilin D null mice were functionally and morphologically protected from renal ischemia reperfusion injury (122). In addition to cyclosporine A, MPT activity can also be inhibited by Mg^{++} , low pH and opening of mitochondrial K channels by valinomycin (297).

Despite tremendous interest in MPT, other mechanisms for mitochondrial swelling may play a role in tubular cell injury. For example, cadmium may cause swelling independent of the classic MPT; in renal tubule cells cadmium induced mitochondrial swelling that was not blocked by cyclosporine A, but rather was blocked by $AgNO_3$ and the mitochondrial calcium uniporter (MCU) inhibitor Ruthenium Red (294).

The potential to study MPT within intact cells and tissues has been enhanced by advances in fluorescence imaging and the development functional mitochondrial probes. For example, in cultured rat hepatocytes, the low molecular weight fluorescent compound calcein distributes from cytosol to mitochondria in response to hypoxia, suggesting increased mitochondrial permeability to low-molecular weight substances. Simultaneously, tetramethylrhodamine methylester (TMRM), a red-fluorescing cationic dye that accumulates electrophoretically into mitochondria is depressed in hepatocytes in response to hypoxia (440). Similar results have been shown in kidney tissue slices in vitro. Hall et al., using 2 photon microscopy,

demonstrated that proximal tubule mitochondrial transmembrane potential $\Delta\psi$ was severely disrupted in slices subjected to chemical anoxia (Figure 6)(185, 601). Theoretically, these approaches could be applied to intact blood perfused whole organs (474). Other potentially important markers of mitochondrial membrane potential including JC-1 and safronin-O have been utilized in models of culture cell injury and isolated tubules (145, 602, 618), but have not yet been optimized in intact organs with live imaging techniques.

Brooks et al have reported mitochondrial fragmentation may contribute to outer membrane permeabilization. Mitochondrial size and structural integrity is mediating by processes of fission and fusion. Mitochondrial fission is more pronounced following ATP depletion in cultured proximal tubule cells and this change is thought to contribute to outer membrane permeabilization and consequent apoptosis (76). Mitochondrial fission is regulated in part by Dynamin related protein-1 (Drp1), which translocates to the mitochondria during early tubular injury. Inhibition of Drp-1 via gene knockdown or dominant negative approaches protected against mitochondrial fission and cell death. A pharmacological inhibitor of Drp-1, known as Mdivi-1, was able to block cell death in vitro as well as in response to ischemic or nephrotoxic damage in vivo (76).

III.1.3.2. Reactive oxygen species: Reactive oxygen species formation in the kidney likely plays a significant role in the genesis acute injury. Evidence has implicated ROS activity as being important in AKI in models of ischemia/reperfusion (106, 127, 392), cisplatin (23), mercury (417), gentamicin (338, 351), endotoxin (620) and glycerol (631). ROS may also influence hemodynamics and inflammation (See Section II). In this section, we will focus on the generation of ROS as a cellular response to injury, and the influence of cellular defense mechanisms in response to injury, occurring secondary to alterations in tubular energetics and mitochondrial function.

The generation of ROS secondary to disturbances in intermediary metabolites in AKI has been the subject of several reviews (392, 521, 599). Reactive oxygen species of significant interest to renal injury include superoxide anion, hydrogen peroxide, hydroxyl radical as well as nitrositive reactive products nitric oxide and peroxynitrite.

Superoxide is generated as oxygen accepts a single electron. This is a highly reactive short-lived radical form that is dismutated to hydrogen peroxide in the presence of superoxide dismutase (SOD) (392, 521, 599). Studies in the rat post-ischemic kidney suggested that type D xanthine oxidase (xanthine dehydrogenase) is converted to type O xanthine oxidase, which uses oxygen, rather than NAD as an electron acceptor (392). During ischemia, hypoxanthine and xanthine are produced as biproducts of ATP metabolism. A role for xanthine oxidase activity in AKI has been suggested by studies in which allopurinol reduced oxidant activity and mitigated injury secondary to I/R (456). However, several other studies have found inhibition of xanthine oxidase to have minimal effects in models of AKI (128, 634). Moreover, the activity of this enzyme is relatively low in human kidney vs rodent kidney, suggesting that its impact in human AKI is unlikely(392).

The altered fidelity of the mitochondrial electron transport chain represents another potential source of ROS contributing to AKI. Significant levels of reactive oxygen species have been measured in isolated mitochondria from injured rat kidney, despite an overall decrease in oxygen consumption (175). This may be due to impaired electron flow at succinate-cytochrome c stage and ROS production at the NADH dehydrogenase and ubisemiquinone sites (175). Blockade of the mitochondrial electron transport chain increases superoxide detectable by dihydroethium fluorescence in kidney slices (185). Infiltrating leukocytes with high levels of NADPH oxidase activity represent a further source of superoxide in AKI (392).

Sirt1, a NAD-dependent protein deacetylase, is reported to regulate intracellular metabolism and attenuate reactive oxidative species generation from mitochondrial activity. When stimulated pharmacologically, Sirt 1 can protect renal tubules against oxidant induced injury (159). When overexpressed in proximal tubules of transgenic mice, Sirt1 transgenic mice are protected from cisplatin induced injury and show reduced ROS stress and an increase in anti-oxidant enzyme expressions such as catalase (193).

Sirt-1 is also an activator of the peroxisome proliferator activated receptor γ coactivator1 α (PGC-1 α) a master regulator of mitochondrial biogenesis and adaptative response to hypoxia (499). The up-regulation of this pathway via adenoviral constructs promotes energetic recovery of tubules following oxidant or toxic injury (453). Recently, isoflavones have been shown to upregulate the activity of this pathway to promote mitochondrial biogenesis via effects on Sirt-1(452).

Liberated iron may contribute to oxidant stress via the catalysis of the Fenton reaction (392, 521, 599). Iron, typically found in storage sites in plasma proteins and heme groups, may be liberated in response to injury and become available for interaction with existing hydrogen peroxide to generate the more insidious hydroxyl radical. Studies using iron chelators such as deferoxamin reduce oxidant associated lipid peroxidation and improved function following ischemia reperfusion or cisplatin (27, 216, 419, 420).

In order to neutralize the effects of reactive oxygen species, renal cells contain a substantial antioxidant defense mechanism. The kidney contains 3 forms of superoxide dismutase, (SOD1/ CuZn SOD, SOD2 Mn SOD, and extracellular SOD/ecSOD/SOD3), catalase and glutathione S transferase. These defense systems are impaired during injury and their activity may modulate the severity of AKI. Cu/Zn SOD, catalase and glutathione levels and glutathione S transferase expressions are reduced in most models of renal injury, although there may be a modest increase in Mn SOD following ischemia reperfusion injury (127, 221, 289, 485, 508, 625). The further impairment of these intrinsic defense mechanism results in worsened outcome in AKI; for example, SOD1 null mice develop more severe AKI to ischemia reperfusion than corresponding wild type mice (611), and chronic exposure of rats to a hypoxic environment reduces renal SOD expression and results in increased sensitivity to AKI (517). It is interesting to note that a recent report demonstrated a significant association between serum catalase activity and the severity of AKI in patients with catalase allele C to T at position -262 (431), suggesting that the intrinsic antioxidant defense pathways of individuals may vary and influence the course of AKI.

Enhancement of intrinsic anti-oxidant mechanisms has protective benefits in models of AKI. Administration of superoxide dismutase decreases severity of AKI in rats (43, 421, 482, 556) and dogs (278), while an increase CuZn SOD gene expression by adenoviral gene transfer protects renal cell death in vitro and in vivo in response to ischemia reperfusion (314, 621). Pharmacological agents such as tempol, which possess SOD mimetic activity, attenuate the degree of renal injury in endotoxemia and I/R models (87, 289). Similarly, MnTMPyP, possessing both SOD and catalase activity and protects against renal ischemia reperfusion (386). Supplementation with glutathione (1, 184, 418, 600) or the glutathione precursor N-acetylcysteine (NAC) prevents the drop in renal glutathione content in response to injury, attenuates oxidant stress and the severity of renal injury in a variety of AKI models (98, 125, 398). However, these approaches tend to have small modulatory roles on the extent of injury and may have limited therapeutic potential. In clinical trials, NAC did not influence AKI incidence or length of stay following cardiac surgery (2, 220).

III.2. Cell death in the setting of AKI

An examination of cellular injury in AKI inevitably results in a discussion regarding the mode of cell death that occurs in the kidney and the potential distinction between necrotic death and apoptotic (also referred to as programmed cell death, PCD) forms of cell death. The reader is referred to several excellent reviews dedicated to this subject (415, 475). The terms apoptosis and necrosis arise from classical pathological terms based on well-defined morphological distinctions. Both apoptosis and necrosis utilize distinct and common biochemical processes; as this represents an area in which potential therapeutic targets may emerge, the study of these processes is an area of intense investigation.

III.2.1. Apoptotic and necrotic cell death and their presence in AKI—The term apoptosis (from greek: apo-from, ptosis falling) was originally described by Kerr in 1972, emphasizing its role in normal cell turnover and distinguishing it from necrotic cell death following traumatic injury (263). In its classical form, apoptosis is a highly ordered process mediated by pre-existing enzyme cascades and requiring energy in the form of ATP.

The earliest morphological features of apoptosis are defined by cytoplasmic and nuclear shrinking, chromatin fragmentation and the breakdown of the cell into multiple spherical bodies (i.e., apoptotic bodies) that retain membrane integrity (415). A hallmark feature of apoptosis is the retention of membrane integrity, which prevents spilling of cellular/lysosomal contents into the interstitium, preventing an inflammatory reaction and limiting the extent of surrounding tissue damage (415). Although membrane components remain intact, phosphatidyl serine residues that are localized in the inner membrane are exposed on the outside of plasma membrane prior to cellular fragmentation where they are involved in targeting cells of the resultant apoptotic bodies for phagocytic engulfment by macrophages or neighboring parenchymal cells. Therefore, when apoptosis occurs on a limited basis during normal cell turnover and in response to modest injury, there is little effect on the surrounding tissue (415).

In contrast, the morphology of necrosis is distinctly different and is characterized by swelling and degeneration of the entire cytoplasm. The chromatin is not packed into discrete membrane bound particles, but forms irregularly shaped clumps. There is dilation and fragmentation of the cisterns of the rough endoplasmic reticulum and polyribosomes are dissociated and dispersed throughout the cytoplasm. The plasma membrane does not remain intact and there is blebbing of the plasma membrane with shedding of the cytoplasmic contents, likely exacerbating inflammatory reactions and expanding the degree of local injury (Figure 7) (415).

Within the setting of acute kidney injury, both apoptosis and necrosis may occur simultaneously. Using protocol biopsies following transplant or post mortem histological analysis of patients with AKI, demonstrates the presence of both necrotic and apoptotic cells (554). In rat ischemia reperfusion injury, Schumer et al, demonstrated that a dose dependency of apoptosis and necrosis in response to ischemia reperfusion injury in rats such that low levels of ischemia (~15 min) manifested only apoptosis and no necrosis. Increasing levels of ischemia were associated increasing appearance of apoptosis as well as necrosis (489).

Although apoptotic and necrotic pathways will be discussed below, we suggest these distinctions should not be viewed sharply. Rather, within the context of renal injury, we suggest that apoptosis and necrosis are not functionally opposed forms of cell death, but rather represent manifestations of death within a spectrum of injury, as put forth by Lieberthal et al., (319, 321)(Figure 8). Because apoptosis requires ATP, and ATP prevents the onset of necrotic cell death, the form of death is dependent on the degree to which ATP is depleted.

Using in vitro models of proximal tubule cells subjected to chemical anoxia, when ATP levels fall below 15% of control values, necrosis ensues. In contrast, when ATP levels are reduced to between 25–50% proximal tubule cells undergo predominantly apoptotic cell death (321). Dose dependency has been observed in response to cisplatin (286), cyclosporine (199), oxidant injury by H₂O₂ and other ROS generating models of injury (564).

The blurred distinctions between apoptosis and necrosis has led Lemasters to propose the term necrapoptosis, to describe the observations demonstrating that both forms of cell death can be observed in tissues subjected to ischemic or toxic insults (298). The term necrapoptosis suggests that both apoptosis and necrosis can be triggered by a common mechanism, but culminates in either cell lysis (necrosis) or programmed self destruction (apoptosis) depending on the nature of other modifying factors (298). The hypothesis is predicated on the idea that the biochemical pathways leading to classical necrosis and apoptosis overlap and interact. Primary among these pathways is the central role that mitochondrial permeability transition plays in triggering the activation of classic apoptotic stimuli and regulating cellular ATP levels (298).

Viewed in this way, the hypothesis put forth by Leiberthal et al., is similar to the idea of necrapoptosis proposed by Lemasters. During the course of apoptotic cell death, if injury is severe and ATP depletion occurs, a secondary necrotic death will intervene. If ATP levels are maintained, cell death occurs primarily by apoptosis. The dose dependency of ATP on the type of cell death may account for the in vivo observation that necrosis is more prominent in the outer medulla where pO₂ reductions are exacerbated by vasoconstriction or congestion, thus prolonging or exacerbating the degree of ATP depletion in this region as compared to a relatively better-perfused cortical zone.

In AKI, the preponderance of apoptotic vs. necrotic cell death may differ substantially depending on the experimental parameters such as the agent, dose and species being investigated. It is widely recognized that differences in sensitivity to injury are prevalent across species. Rats and mice both show overt necrosis in response to I/R injury. Mice typically manifest significant injury with shorter ischemic times in the range of 25 minutes (609) with some reports of significant necrosis resulting from an injury in as little as 17 minutes (266). Studies in rats utilize longer ischemic times (60–60) minutes of ischemia (554) although some studies have demonstrated necrosis in rats with as little as 25 minutes of ischemia (573). As mentioned earlier, although evidence of tubular damage is seen in human kidneys, necrosis is patchy following ischemia suggesting an overemphasis of cell death in rodent models (323). This may relate to the size of the organism and the degree to which juxtamedullary blood flow is affected in larger kidneys. In dogs, juxtglomerular flow is not dramatically reduced following release following clamps over 120 minutes. There is little evidence of overt necrosis despite the persistent evidence of sublethal injury (323, 458). Similarly, pig models show modest necrosis with up to 3 hours of ischemia (26, 462).

Although mercuric chloride toxicity has long been reported to be associated with the rapid development of renal epithelial necrosis (566), models using toxins to induce AKI are usually executed to develop more slowly and show a more limited necrotic response. This may be related to the doses and timing used for each study. Cisplatin injury shows a peak onset of injury as late as 3 days following the administration of the agent and is characterized by a mix of apoptotic and necrotic death, although apoptotic cell death is readily identifiable in this model is early as 2 days post-injection (362, 450). Typically, many nephrotoxic agents (e.g., cisplatin, gentamicin, mercury, chromium), will manifest evidence of necrosis by 3 days, while manifesting evidence of sublethal damage at earlier time point (211, 442, 644). All of these agents are reported to induce apoptotic pathways in

vivo and in vitro (211, 281, 388, 442, 450, 466, 493) suggesting that much of the necrosis observed is secondary.

III.2.2. Biochemical pathways mediating necrosis—Necrotic cell death is historically viewed as a passive process secondary to the breakdown of cellular energetics. Reductions in cellular ATP levels result in reduce Na-K ATPase activity and net Na influx associated with swelling of the cytosolic compartment (62, 353). Although the site of passive sodium entry in ATP depletion remains elusive (415), the associated cell swelling is thought to be antecedent to necrotic death. Zager et al., demonstrated that mannitol protected against cell swelling in ischemia reperfusion, but did protect against oxidant stress, mitochondrial swelling or energetics (630). Similar protection from cellular swelling was also observed in rhabdomyolysis model induced by glycerol, however mannitol actually impaired energetic recovery in this model (633).

Histological evidence of necrosis is typically, but not exclusively, observed in proximal tubules, while sparse in the distal tubule and TAL. Among the possible explanations for this difference is the relatively high hydraulic conductivity of proximal tubule membranes that contribute to swelling secondary to Na influx (61). Aquaporin-1 is the primary water channel in the proximal tubule that contributes to water reabsorption and therefore, it may contribute to cell swelling. In contrast, the plasma membranes of the TAL and DT are largely impermeable to water. The idea that aquaporins contribute to swelling associated with necrosis is supported by studies showing that aquaporin 4 null mice are protected from edema in various brain injury models (e.g., glutamate, focal ischemia) (349). Although aquaporins have been investigated with regard to diuretic activity in acute kidney injury, no studies have examined their potential contribution toward osmotic cell swelling in AKI.

The potential role of Ca^{++} ion in mediating renal damage in acute injury has been a major area of research for over 30 years. A role for Ca in the contribution to renal injury is suggested due to the protective effects of Ca^{++} channel blockers (486, 487). Verapamil, a slow Ca^{++} channel antagonist, conveys protection in a model of norepinephrine induced AKI (486, 487). Both verapamil and diltiazem have been used to block AKI associated with transplant and both provided significant protective effects to renal allografts (116). However, verapamil does not directly protect renal tubule cells in suspension, suggesting that Ca^{++} channel antagonism is directed at renal vascular impairment rather than cell injury pathways (60).

Nevertheless, injured or dying cells accumulate cytosolic Ca^{++} thought to derive from both intracellular and extracellular sources (60). There is a ~10,000 fold gradient of Ca^{++} concentration (1 mM extracellular vs 100 nM intracellular) that is normally maintained by active energy requiring extrusion mechanisms. The sarcolemmal Ca-ATPase actively pumps Ca^{++} into the ER compartment, representing a high capacity reserve depot for excess Ca^{++} . This system is compromised when ATP levels are reduced (60). Proximal tubular free ionized cytosolic Ca^{++} rises from values of 170 to 390 nM during a 5 min hypoxic injury (486, 487). The increased Ca^{++} is readily reversible with brief hypoxic insults but with prolonged hypoxia or ischemia, Ca^{++} remains elevated and may participate in injurious processes (vide infra) Reductions in extracellular Ca or intracellular sequestration of intracellular Ca^{++} conveys cytoprotection to renal cells in suspension or in culture (607).

The Na/Ca exchanger typically couples Na-entry with Ca^{++} release; under conditions of elevated intracellular Na, this exchanger may work in reverse mode and promote the build up of intracellular Ca^{++} (60, 415). Mice heterozygous for the NCX1 gene express approximately 50% of Na/Ca exchanger protein in proximal tubules. Cultured proximal tubules from these mice demonstrate reduced Ca^{++} influx to hypoxia, while kidney function

and cellular damage in vivo are minimized following ischemia reperfusion injury (612). Moreover, drugs which prevent the Na/Ca exchanger from operating in reverse mode attenuate kidney injury in response to ischemia reperfusion injury (406, 612).

Ca⁺⁺ regulates a number of processes that influence cell injury. As described above, increased cytosolic Ca⁺⁺ may influence the process of MPT. Ca⁺⁺ may also participate in activation of Ca⁺⁺ dependent proteases, phospholipases and endonucleases; processes described further in the following sections.

III.2.3. Biochemical pathways involved in apoptosis and their involvement in AKI—Apoptosis is regulated by several interrelated biochemical pathways targeted toward the programmed self-destruction of the cell in an orderly fashion (Summarized in Figure 9). A hallmark feature of apoptosis is degradation of DNA by specific endonucleases. Endonuclease activity at accessible areas between nucleosomal regions results in the characteristic “DNA ladder”, while necrotic cells demonstrate a less orderly smearing pattern. Nuclease activation represents a point of no return in the injury process. Basnakian et al, identified abundant expression of the gene encoding DNase I in rat kidney and demonstrated that its activity was increased in response to ischemia reperfusion injury and cisplatin (39, 41). Specific inhibition of DNase I with antisense oligonucleotides protected tubular epithelial cells against hypoxia reoxygenation in culture (41).

There is also evidence for other endonucleases activated in response to injury. For example, Endonuclease G (Endo G) is a small 27 kDa protein residing in the inner membrane space of mitochondria and can traverse to the nucleus under apoptotic conditions (313). In rat kidney tubular epithelial cells, Endo G was transported to the nucleus in response to hypoxia reoxygenation. The increased mitochondrial permeability was shown to be dependent the production of ceramide, since the process was inhibited by fumosin B1, an inhibitor of spingomylinase (40). Both DNase I and Endo G may play complimentary and synergistic roles in endonuclease activity in models of AKI; using DNase I null mice, Endo G was found in nuclei of proximal tubule cells following cisplatin injury while inhibition of Endo G in vitro protected against DNA strand breaks in response to cisplatin (622). In addition, another endonuclease, termed apoptosis inducing factor (AIF), may be activated in response to cisplatin injury (333).

III.2.3.1. Initiation and downstream signaling of apoptosis in kidney injury: Caspases, or cysteine-aspartic proteases, are a family of enzymes which are thought to play an essential role as primary initiators of cell death. Caspases are constitutively expressed in a latent form and activated in response to a variety of potentially apoptotic stimuli. They are typically described as comprising a canonical pathway in which initiator caspases (caspases-2, -8, -9, 10) are sensitive to an initiating signal, self activate, and then act upon subsequent “effector” caspases (caspase-3, -6,-7). Effector caspases are in turn responsible for activating pathways associated with the orderly destruction of the cell (e.g, DNase and protease degradation). Historically, apoptotic pathways are considered to be activated from signals associated with intrinsic cell stress, or from factors extrinsic to the cell (Summarized and reviewed in (413, 415, 563)). Therefore, cell death may occur via common mechanisms regardless of the stimuli, as initiator caspases associated with extrinsic and intrinsic pathways both converge on the common effector caspases. Both caspase 3 and caspase 6 are expressed in kidney and activated in response to injurious stimuli (143, 252, 366, 493) and caspase 6, localizes to the nucleus following ischemia reperfusion injury (507). The importance of both intrinsic and extrinsic pathways in multiple models of AKI is generally well-accepted based numerous studies, a portion of which are described below.

III.2.3.2. Extrinsic pathway: The extrinsic pathway is one typically associated with activation of a subset of TNF receptor family cell surface “death” receptors which includes Fas/CD95, TNFR1, and death receptor-3 (415). Engagement of death receptors by their ligand invokes receptor trimerization and complex formation within the cytoplasmic bounds of the receptor with molecules such as FADD/MORT, TRADD and RAIDD, which contain death effector domains (DED). Caspase-8 represents the key initiator caspase of the extrinsic pathway and it is presumed that the association of procaspase-8 with these DED permits its autoproteolytic activation to caspase-8 and subsequently caspase-3.

Tubular epithelial cells undergo apoptosis in response to members of the TNF ligand superfamily, including TNF- α , FasL (CD95) and TRAIL (475). Administration in vivo of CD95 (Fas) results in endothelial apoptosis, which can be blocked by the inhibitor to caspase-8 or with the z-IETD peptide that selectively inhibits caspase-8 (232). These pathways are particularly relevant in models of sepsis. Cunningham demonstrated that TNF-receptor null mice were resistant to the injurious effects of endotoxin on renal function and apoptotic death (107). Other studies have shown that peptide based inhibitors of caspase 8 as well as caspase 3 attenuated endotoxemic injury (179). In a cecal ligation model of sepsis, the administration of siRNA to block either caspase-8 or Fas improved outcome and organ apoptosis, although the kidney injury was not a target in that study (603).

The activation of the extrinsic pathway is not exclusive to sepsis models of AKI; Fas mRNA was increased, and immunoreactive Fas protein localized to proximal tubule cells, within hours of renal ischemia reperfusion injury (399). Dendritic cells may be the source of TNF α following renal ischemia reperfusion that could activate this pathway (129). Inhibition of caspase 3 and 8 with siRNA significantly attenuated renal ischemia reperfusion injury in mice, however, since both caspase 3 and 8 were inhibited in this study, it is difficult to distinguish the effects from the intrinsic pathway, which is also active in these settings (640).

The extrinsic pathway may be activated secondary to other inflammatory or modulatory factors. In cultured renal tubular cells, pro-inflammatory cytokines IFN- γ and TNF α increase NOS-2 activity and lead to the formation of active caspase-8 and apoptotic cell death. This effect was blocked not only by L-NMMA, but also several strategies to inhibit caspase 8 such as the caspase-8 inhibitor z-IETD-fmk, the overexpression of the endogenous caspase-8 inhibitor cFLIP, and silencing of caspase-8 (134). The pro-inflammatory cytokine IL-2 also causes apoptosis in proximal tubule cells, and this effect is reduced in cells from Fas $^{-/-}$ mice and by c-FLIP (135). Natural Killer T cells, (NKT) may also modulate the extrinsic pathway in ischemia/reperfusion injury. The NKT receptor Rae-1 is expressed on the surface of tubular epithelial cells following IR injury; NKT cell interaction can cause cell death in a FasL dependent fashion (641).

III.2.3.3. The intrinsic pathway: The intrinsic pathway is responsive to cell stress and is activated by oxidative stress/reactive oxygen species, radiation, ceramide, and growth factor deprivation. The loss of normal morphology or contact with substratum may also induce apoptosis. It is interesting that the stabilization of the actin cytoskeleton reduces apoptotic death in early IR injury (166).

Features typically associated with the intrinsic pathway are the release of mitochondrial apoptogenic molecules, particularly cytochrome c, the activation of procaspase-9, and the interaction of bcl-2 family members. The release of cytochrome c into the cytosol from the inner membrane space of the mitochondria is sufficient to induce apoptosis in some cell types (413, 415, 563). Cytochrome c binds and oligomerizes with the adaptor protein apaf-1,

which in turn recruits and pro-caspase -9 leading to the formation of the apoptosome, and the autocatalytic activation caspase-9 (413, 415, 563).

Activated caspase 9 can proteolytically activate caspase-3. A central role of cytochrome c/ caspase 9 in the intrinsic pathway is demonstrated clearly in classic studies using cell lines derived from cytochrome C $-/-$ mice, which do not activate caspase-3 and are resistant to the proapoptotic effects of UV irradiation, serum withdrawal, or staurosporine (vs. wild type derived cells). In contrast, these cells remain sensitive to extrinsic pathway stimulation (310). Similarly, thymocytes derived from Caspase 9 $-/-$ mice, show reduced caspase-3 activation and cell death induced by etoposide, dexamethasone, and γ -radiation when compared to wild type cells, and cell death induced by Fas antibody was similar in thymocytes from wild type and caspase 9 $-/-$ mice (282).

Given the importance of mitochondrial cytochrome c release (and other mitochondrial derived pro-apoptotic factors), the understanding of this step is significantly important to understanding the injury process. The control of mitochondrial permeability is often attributed to members of the bcl-2 family of proteins classified as either pro-apoptotic (bax, bad, bak) or anti apoptotic molecules (Bcl-2, bclXL, Bcl-w, Mcl-1, and A1). Pro-apoptotic Bcl-2 family members (e.g., bax) insert into the outer membrane and increase permeability of the membrane to proteins such as cytochrome c, either by forming channels, interacting with membrane proteins to facilitate pore activity (i.e., VDAC), or by contributing to membrane depolarization, in which cytochrome c would be released secondary to swelling (595).

Translocation of bax from cytosolic compartment to the mitochondrial compartment precedes the release of cytochrome c following injury. Saikumar et al., demonstrated that hypoxia leading to ATP depletion resulted in a translocation of bax protein to the mitochondria and release of cytochrome c (469). Caspase inhibitors did not prevent the translocation of bax and the release of cytochrome c; interestingly, caspase inhibition prevented apoptotic cell death but not necrotic cell death, as mitochondria did not recover (469). Others have demonstrated translocation of bax, as well as another pro-apoptotic member, bak, in culture models of ATP depletion (368), cisplatin (235, 388), and in vivo within 6 hours of I/R injury (438). Bax mRNA and protein are up-regulated rapidly following I/R injury in the kidney (36, 172). Bax null mice were resistant to cytochrome c release, apoptosis and had preserved renal function and morphology following administration of cisplatin (597).

Two other pro-apoptotic members of the Bcl-2 family have also been identified in kidney injury models; both bad and bid translocate to mitochondria in hypoxic tubule cells and in response to ischemia reperfusion injury similar to bax (596). Data from biopsies suggest that these pathways may operate in human kidneys following transplant (85). Bid activation is mediated by a cleavage event inhibited by caspase 8 or caspase 9 inhibitors, suggesting a potential link between extrinsic and intrinsic pathways. In kidney cells, bid sensitivity to caspase -9 inhibitors suggests that it may amplify the mitochondrial pathway (596). Bid deficiency in transgenic mice was shown to alleviate the extent of renal damage using a model of ischemia reperfusion (598).

The regulation of bax activity involves, in part, interactions with other proteins resulting in both complexity as well as potential avenues for therapeutic intervention. A well-studied area of investigation involves the interaction that occurs with other Bcl-2 family members. Heterodimerization with Bcl-2 can nullify the efficacy of Bax to release cytochrome c (412). Using kidney tubular epithelial cells, Wang et al., demonstrated that minocycline induced the expression of Bcl-2, accumulating in the mitochondria and interacting with Bax, Bak and

Bid; specific inhibition of Bcl-2 using an siRNA strategy, reduced the protective effects of minocycline in response to hypoxia, azide, cisplatin and staurosporine (587). Transgenic overexpression of Bcl-2 or BclXL attenuates damage associated with in cardiac ischemia reperfusion and acute liver injury (92, 103, 119, 373, 430). Alterations in the activity of Bcl-2 family members may underlie the differential sensitivity of proximal tubular vs. distal tubular injury. Cuttle et al, demonstrated that distal tubule cells in culture express the anti-apoptotic protein of Bcl-X_L, which translocates to the mitochondria in surviving distal nephron cells in response to peroxide treatment, while proximal cells do not express prominent of Bcl-X_L or Bcl-2 and die from apoptosis (111).

The reasons why pro-apoptotic Bcl-2 family proteins migrate to the mitochondria remain elusive. It may involve conformational changes affecting protein-protein interactions. JNK- a pro death signaling molecule in injured tubule cells, affects bax translocation (296, 559). Similarly, inhibition of p53 activity protects in renal ischemia reperfusion injury, in part, by blocking bax translocation (115). A newly identified factor, nucleophosmin is a bax chaperone that may facilitate bax delivery to the mitochondria (264).

III.2.5. Autophagy—Autophagy is a cellular process of bulk degradation of damaged organelles, protein aggregates and macromolecules in the cytoplasm (432). It is a stress response suggested to contribute to cell death. Recently, Periyasamy-Thandavan et al., suggest that this process is renal protective in AKI. In a model of cisplatin induced injury, autophagic vesicles and autophagosomes were identified in tubular cells prior to apoptotic cell death (433). When autophagy was blocked pharmacologically with 3-methyadenine or bafilomycin, tubular cell apoptosis was enhanced. Similarly, knockout of the pro-autophagic gene, beclin, also enhanced the apoptotic response to cisplatin (433). However, the role of autophagy as contributing toward cell death or cytoprotection in the setting of AKI is controversial. This area is likely to garner increased attention in the near future (130).

IV: Repair and Regeneration

IV.1. Adaptive nature of kidney and proximal tubules

As described above the cellular response to injury is heterogeneous, with some cells undergoing necrosis or apoptosis, while others are sublethally injured. The severity and duration of AKI may be related to the proportion of sublethally damaged cells that are able to maintain viability and contribute to a coordinated repair process restoring kidney structure and function. Renal tubular cells are remarkable for their adaptive nature to threatening situations and maintain viability. An examination of gene expression following renal injury reveals groups of genes that are associated with adaptation to injury (123, 529). The activation of such pathways represent potential therapeutic targets to lessen the severity of AKI, while the down-regulation of such pathways may exacerbate injury and may represent clinically relevant risk factors for the development or outcome of AKI.

IV.1.1. The renal stress response—Injury to the kidney activates well-defined cytoprotective pathways, referred to as the renal stress response, which can be classified broadly in at least 3 pathways with multiple targets.

IV.1.1.1. Heme Oxygenase (Inducible HO, HSP32) and anti-oxidant genes: The heme oxygenase enzyme system catalyzes the rate-limiting step in heme degradation, producing equimolar quantities of biliverdin, iron, and carbon monoxide (CO). Biliverdin is subsequently converted to bilirubin by biliverdin reductase. Two isoforms of heme oxygenase have been characterized: an inducible enzyme, HO-1, and a constitutive isoform, HO-2 (505). Activity of HO in response to various signals associated with injury is thought to contribute to cytoprotection. In animal models, HO-1 is rapidly induced following

ischemia/reperfusion (347), glycerol-rhabdomyolysis (391), and nephrotoxic injury (3). HO-1 expression is controlled primarily based on its transcriptional activity and can be influenced by multiple factors including, growth factors, Ang II, inflammatory signals, NO and heavy metals. However, alterations in cellular redox state represents one of the most powerful regulators of HO-1 activity (505). The redox sensitive transcription factor, Nrf2 represents a key regulatory molecule in the activation of antioxidant genes in kidney injury and is a well-described modifier of HO-1 transcription. Mice with a null mutation in the Nrf2 gene demonstrate significantly reduced activation of a number of antioxidant genes such as HO-1 (334). In relation to the mouse promoter, the human HO-1 promoter is much less responsive to factors such as hypoxia, interferon- γ and oxidized lipids. However, the human promoter is very responsive to induction by heme (505). Increases in heme may be attributed to increased filtration in the setting of rhabdomyolysis or may be liberated from destabilized proteins in response to ischemia or nephrotoxin induced damage; such increases precede activation of HO-1 expression (505).

The functional importance HO-1 in AKI derives from multiple studies in which HO activity is impaired. For example, administration of tin mesoporphyrin, which blocks HO-1 activity, worsens renal function in response to ischemia/reperfusion (498). Mice with a null mutation in HO-1 show greater loss of kidney function and increased mortality in response to ischemia reperfusion (437), glycerol (391), and LPS (367). Similarly, Nrf2 null mice with reduced HO-1 induction also manifest greater injury to ischemia and cisplatin (334). Conversely, induction of HO-1 expression by infusion of hemoglobin intravenously prior to injury, or viral delivery of HO-1 expressing gene, protects from injury (392, 393). Other strategies have been used to increase HO-1 activity such as adenoviral gene transfer (55), and cobalt chloride administration (96, 185), which protect against renal injury induced by either warm ischemia reperfusion injury or cold ischemia in transplant models. HO-1 minimizes oxidant stress by virtue of its degradation of heme, a toxic pro-oxidant, as well as its generation of bilirubin, a peroxy radical scavenger that inhibits lipid peroxidation. In addition, CO is byproduct of the HO-1 activity, providing a potential vasodilatory effect on the renal vasculature and a restoration of blood flow (393). HO-1 expression is present in renal biopsies following transplant and its levels correlate with the degree of post transplant function (301).

IV.1.1.2. Heat shock proteins: Sublethal heat stress transiently increases the ability of a cell to withstand an otherwise subsequent lethal heat challenge. Similar protection is routinely observed in response to diverse stressors such as hypoxia, ischemia, heavy metal exposure and oxidative stress. Interest in understanding this protective response led to identification of a fairly large family of proteins known as heat shock proteins; a multigene family of proteins ranging in size from 10–150 kDa and found in all major cellular compartments (46). Heat stress has similar consequences as hypoxia at the cellular level. For example, both heat stress and hypoxia are settings where damage to the cytoskeleton occurs, where there is mitochondrial swelling, and where there is uncoupling of oxidative phosphorylation. Hence, the term “heat stress protein” has been proposed to reflect this more generic association with cellular stresses (46).

One of the earliest reports suggesting a role for heat shock factors in protection from acute kidney injury was from Emami et al (140); transient minimal ischemia (15 min) resulted in an increase in heat shock protein72 (HSP72), which was associated with protection to subsequent ischemia/reperfusion injury. Whether the protection was actually attributable to HSP72 was not addressed in that study as methodologies to specifically inhibit HSP72 specifically were not available at the time.

It is now clear that not only HSP 72, but also other HSP family members including HSP10, HSP 25/27, HSP47, HSP70 and HSP90 and $\alpha\beta$ -crystallin are substantially induced in diverse models of renal injury including ischemia reperfusion, cisplatin and unilateral ureteral obstruction (340, 484, 494, 509, 570, 642). The mechanisms of gene induction of the various heat shock proteins is quite heterogeneous, however, many of the inducible forms of HSP (i.e., HSP 72, HSP 27) contain heat shock elements in their promoters, which are activated upon binding of the regulatory protein, heat shock factor (HSF) (521, 570, 571). Under non-stress conditions HSF is bound to the constitutively-express HSP referred to as heat shock cognates (HSC), often HSP 70. Increased demand for the use of HSC following a stress leads to a dissociation of HSC/HSF complex, thus liberating this regulatory molecule and activating transcription (Figure 10) (521, 570, 571).

The protective effect of HSPs is widely attributed to their activity as "molecular chaperones," which are proposed to assist in the assembly or repair of newly synthesized or damaged proteins (65). *In vitro*, chaperones function to prevent aggregation of other proteins under conditions of stress and to promote restoration of enzymatic activity of denatured protein substrates or enzymes on removal of the stress (46, 65). HSPs are found in multiple protein aggregates following the induction of ischemic kidney injury. Hsp72 binds to aggregated cellular proteins, including Na/K-ATPase, and is released upon addition of ATP *in vitro*, a feature typical of its chaperone function (19).

A prominent role of HSPs is associated with the stabilization and recovery of cytoskeletal structure following injury. During recovery following ischemia, the Na, K, ATPase reassociates with the cytoskeleton, as evidence by its reappearance into a Triton X100 insoluble pellet. HSP-70 is also present in the Triton X100 insoluble pellet and correlates with the reappearance of the Na, K, ATPase. When antibodies against HSP-70 are included with the insoluble fractions, the re-association of the Na, K, ATPase with the cytoskeletal fraction is inhibited, while the addition of HSP-70 protein promoted Na, K, ATPase/cytoskeletal interaction *in vitro* (50). HSP72 co-precipitates with focal adhesion component paxillin following ATP depletion; the overexpression of HSP 72 hastens the re-establishment of focal adhesions following transient ATP depletion (350).

HSP27 has also been identified in cytoskeletal protein aggregates from injured kidney (567). In cultured proximal tubule cells, HSP27 migrates to the basolateral cell boundaries following ATP depletion. Overexpression of either HSP-72 or HSP-27 preserves the alterations in cytoskeletal architecture in response to ATP depletion (135, 176). HSP-27 may preserve tight junction formation via an indirect effect on Src-activation (198).

HSP-chaperone activity may confer cytoprotection by interacting with, and inhibiting the biochemical pathways leading toward cell death. HSP 72 associates with AIF (apoptosis inducing factor) and prevents leakage of AIF from mitochondria, the nuclear accumulation of AIF and subsequent DNA fragmentation in ATP depletion (461). Furthermore, HSP-27 antagonizes conformational changes in Bax and protects mitochondrial injury (197).

HSP family members may also function as cellular antioxidants. In MDCK cells exposed to oxidant stress, HSP-70 prevented the increase in superoxide production by increasing the activity of glutathione peroxidase and glutathione reductase (180). HSP-90 coupled to NOS3 preserves NO synthesis, while when uncoupled from NOS3, leads to the formation of vasoconstrictive superoxide (439). Hypoxia increases HSP-90-NOS3 coupling in coronary endothelial cells. More recently, HSP-90 has also been shown to modulate the vasorelaxation induced by NOS1. Agents such as radicicol inhibit HSP-90/NOS3 interactions and lead to an acute reduction in renal blood flow (451). Therefore, cumulative HSP activity may improve renal blood flow in the regenerating kidney by improving the

nitric oxide/superoxide ratio. HSPs may also provide protection in the setting of injury by attenuating inflammation. Ischemia/reperfusion–induced NF- κ B activation, MCP-1 expression and monocyte infiltration was suppressed by heat preconditioning. Inhibition of HSP-70 by quercetin reversed these effects, while liposomal delivery of HSP 72 into renal proximal tubule cells blocked the activation of NF- κ B in response to in vitro hypoxia-reoxygenation (71, 99).

Until recently, in vivo studies on HSP function in AKI have been lacking due to the paucity of transgenic models. Transgenic mice overexpressing HSP-70 are protected in a model of cerebral infarction (448), but these mice have not been evaluated in AKI. Recently, Kim et al., demonstrated that the selective overexpression of HSP-27 targeted in proximal tubules conferred resistance to ischemic AKI (270). Genetic factors may influence HSP activity; higher basal levels of HSP-72 and HSP-27 were reported in Brown Norway rats which are resistant to AKI relative to commonly utilized Sprague Dawley rats (30). Other approaches such as preconditioning (described in more detail below) are consistent with the idea that up-regulation of HSP provide protection in AKI. In patients, it has been suggested that genetic polymorphisms in HSPs, specifically, the Hsp72 1267-GG allele may increase the risk of acute kidney injury in low birth weight neonates (572).

IV.1.1.3. Stress activated protein kinases: Cellular stress induced by oxidants and DNA-damaging agents activates two related parts of the mitogen-activated protein kinase signaling pathway, or MAPKs—the extracellular-regulated protein kinases, or ERKs, and the stress-activated protein kinase (SAPK), also known as Jun N-terminal kinase (JNK and p38). Both of these systems are activated in response to kidney injury and/or renal tubular cell stress in vitro (464). Jun N terminal kinases (JNK and P38) are activated by inflammatory cytokines such as TNF-alpha and IL-1 and are only minimally activated by growth factors (464). JNK is activated in both proximal and distal tubules following ischemic injury (124). JNK down regulation ameliorates proximal tubule injury in response to peroxide induced injury (10)

ERK1/2 activity is typically activated by growth factor receptors, particularly, the EGF-receptor and other vasoactive molecules. The role of ERK in acute kidney injury is somewhat obscured due to conflicting results using different model systems. One point of view suggests that ERK activity is associated with cellular protection and facilitation of renal repair. Safirstein and colleagues demonstrated that ERK, but not JNK was activated in thick ascending limb cells following renal ischemia (464) and suggested that this was a means by which this nephron segment was protected from injury. Overexpression of ERK by adenoviral expression vectors in mouse proximal tubules resulted in protection from peroxide induced injury in vitro. When kidneys are protected via preconditioning, ERK activation is induced, while JNK activation is suppressed. Activation of the EGF receptor may be important in kidney regeneration (see below); activation of ERK via EGF receptor activity may convey protection to cells undergoing regeneration or to neighboring uninjured cells in a hostile environment. The non-specific MAPK inhibitor genestein blocked ERK induction following glycerol induced injury and this was associated with a less effective proliferative response (228).

In contrast, however, a significant literature exists demonstrating an important role for ERK activation in the induction of apoptotic cell death (reviewed in(646)). Inhibition of ERK activity blocks apoptosis associated with cisplatin nephrotoxicity in opossum proximal tubules in vitro (271). Pretreatment of MEK inhibitor, U0126, decreased ERK1/2 phosphorylation following cisplatin administration with significant functional and histologic protection (236). Inhibition of ERK can effect both intrinsic and extrinsic apoptotic pathways; ERK activation was shown to be upstream of caspase-3 activation (236). In

response to cisplatin induced injury, inhibition of ERK may protect cells by repressing bax expression and/or inhibiting p53 activity (78, 206). The reason for the discrepant results remains unclear; it is likely that these are due to differences in injury models (i.e., cisplatin, hypoxia, reactive oxygen species) and the integration of down stream signaling pathways under different experimental conditions (e.g., JNK activation, growth factor activity, severity of injury).

IV.1.1.4. Other adaptive and reparative pathways: In addition to classic stress response pathways, additional pathways are activated in response to injury that may play an important role in stabilizing the injured kidney and/or promoting cellular repair processes.

The hypoxia-inducible factor-1 or 2, (HIF1 and HIF 2) are oxygen sensitive helix-loop-helix transcription factors that regulate biological processes which are geared toward facilitating oxygen delivery and/or providing cellular adaptation to hypoxia (183). This pathway is regulated by post-translational modification of HIF. Prolyl hydroxylase continuously modifies the HIF protein; this activity promotes the binding of HIF molecules to the pVHL-ubiquitin-ligase complex and directs HIF toward proteosomal degradation. Under hypoxic conditions, prolyl hydroxylase activity is reduced, minimizing the HIF degradation and allowing for HIF to interact with HIF response elements on a multitude of potential target genes (183). HIF target genes, if activated in AKI would clearly be beneficial to adaptation and repair, such as HO-1 and VEGF.

HIF1 α / HIF1 β dimerization and nuclear translocation has been reported in kidneys following ischemic injury (139), and in response to hypoxia by vasoconstriction or anemia and by cobalt chloride. HIF1 has been observed in tubular epithelial cells while HIF2 staining was observed in endothelial cells (137, 138). In cultured proximal tubule cells, HIF1 competent cells are more resistant to hypoxic injury than HIF null cells due to increased glucose uptake capacity in HIF-competent cells (51); however, since proximal tubule cells in vivo do not utilize glucose primarily, the importance of this observation is unclear. Recent studies have demonstrated that both HIF1 α and HIF2 α heterozygote mice or HIF2 null mice manifest more significant injury than corresponding wild type mice in ischemia reperfusion injury (205). Strategies to induce HIF-1 activity have also been evaluated; prolyl hydroxylase enzyme inhibition results in renal HIF-1 activation with significant preservation from injury (205). Similarly, Morg1 (MAPK organizer 1) an enzyme that interacts prolyl hydroxylase, was evaluated in transgenic animals; HIF activation was increased and renal damage attenuated in Morg1 heterozygote mice relative to wild type controls (189). Although this latter report describes a reduced induction of inflammatory factors, the HIF target genes that mediate these protective response have not been well-delineated.

IV.2. Preconditioning and pharmacological maneuvers of cellular protection

Preconditioning is a powerful experimental maneuver that reproducibly protects the heart against subsequent ischemic challenge (41, 107). The concept of preconditioning applies readily to the kidney. Multiple studies carried out by several groups have demonstrated that prior stress or injury to the kidney provides resistance to subsequent injury (59). Interestingly, preconditioning effects by nephrotoxic agents were reported by MacNider as early as the 1920s (343).

Preconditioning has been studied using several experimental approaches. For example, an “early window” preconditioning occurs in response to one or several bouts of short ischemia (varying between 4–15 minutes), a short recovery time (on the order of a few hours) followed by longer bout of ischemia that would be considered injurious (e.g., 30–60 minutes). Several examples of early-window preconditioning in the kidney exist (291, 555).

It is typically interpreted that this preconditioning relies heavily on non-genomic activity to provide protection. Evidence suggests that preconditioning increases renal NO production leading to preservation of renal perfusion (233, 613). Other evidence suggest that the rapid activation of adenosine A1 receptors may mediate the preconditioning effect as A1 agonist treatment mimicked the pre-conditioning response, while an A1 antagonist prevented the preconditioning effect (291). Activation of the adenosine A3 receptor subtypes worsened renal injury.

“Late” window preconditioning refers to a process in which kidney injury is induced and is resistant to a second challenge during or shortly after the normal recovery response. The protective effects can be seen even very early during the repair phase of injury, i.e., within 1 day (636). The effect can also be prolonged; using a model of glycerol-induced injury, Nath et al., demonstrated protection from a second insult up to one month following recovery from the first injurious stimulus (394). In a mouse model of ischemia reperfusion, Park et al observed protection to a second injury between 5 and 14 days following recovery, while the protective response was gradually reduced between 14 and 28 days of recovery (424). The preconditioning effect may depend on injury in general and not the specific nature of the insult since other types of injuries induced by either cisplatin or transient ureteral obstruction can also have prolonged protective effects (122, 197, 203).

The time course of resistance following such injuries suggests production of adaptive molecules may convey protection. A substantial number of genes are thought to be modulated by the preconditioning process (174). These include the afformationed stress response genes such as HSP25, HSP 72, p38 MAPK, HO-1, and HIF-1 (59, 124, 393, 425, 568, 615) which may play a modulatory role in protection. Interestingly, injury itself may not be required; repetitive bouts of systemic hypoxia, in and of itself not injurious, can increase the expression antioxidant genes, such as SOD-1 or invoke anti-apoptotic genes and protect against I/R (88, 615). Alterations in cellular lipid composition are also present in the preconditioned state. A consistent finding secondary to multiple models of acute injury is the accumulation of cholesterol due to new synthesis of HMG CoA reductase (632, 635). Blockade of the HMG CoA reductase gene was shown to abrogate the cytoprotective effects of ischemic preconditioning (637).

Several strategies have been attempted to tap into the powerful protective effects conveyed by preconditioning. One strategy derives from the what is termed “remote” preconditioning, which refers to a process where injury in one tissue conveys protection at a distant organ site (449); this phenomenon suggests that circulating factors derived from ischemia may convey the protective effects. This cross talk appears to apply to injury in multiple organ systems; for example, short term ischemia in the kidney protects subsequent myocardial ischemia and these effects can be inhibited by adenosine A1 antagonists and mitochondrial KATP channel blockers (196). Conversely, liver injury can protect against subsequent injury to the kidney (17). One potential strategy might involve the induction of transient limb ischemia which has remote effects on preserving endothelial function (339).

A second strategy involves the activation of stress response pathways by various chemical and pharmacological agents. Preconditioning effects can be mimicked by multiple agents such as anesthetics like xenon, which activates ERK or HIF pathways (341). Inhaled CO and administration of a CO-releasing compound protect against AKI or transplant associated injury (396, 575). Several other factors that potentially influence stress response pathways and have protective effects include glutamine (157), erythropoietin (616), geranylgeranylacetone (535) and tin protoporphyrin (243).

IV.3. Regeneration

The restoration of kidney structure following injury has been the subject of substantial investigation. Evidence of recovery responses in the post injury period has been observed in patient biopsies following ischemic or nephritic injury or in biopsies from patients with delayed graft function (447, 511, 514, 515). In both rats and mice, severe acute injury following ischemia or nephrotoxic insults results cellular proliferation as shown by ³H-thymidine (109, 110, 361) or BrdU incorporation (219, 325, 369, 370, 520) or by the expression of PCNA (360) or Ki67 antigen (481) usually apparent in less than 1 day following injury. The vast majority of this proliferation is present in the proximal tubule cells.

In addition to cell proliferation, the restoration of proximal tubule structure and function requires a variety of complex and well coordinated cellular activities (See Figure 11) (402). For example, in the early post-ischemic period, gaps of naked basement membrane can be found from which necrotic or desquamated cells have detached. Newly formed proximal tubule cells are thought to migrate rapidly to realign the proximal tubular basement membrane (402). These newly formed proximal tubule cells are dedifferentiated and have a very different gene expression profile vs. fully differentiated proximal tubules. Typical markers of epithelial cells such as cytokeratin and E-cadherin are lost and there is expression of markers of undifferentiated cells, such as vimentin (608). In renal injury models using young healthy rodents, the proliferative response is more robust than is required to replenish the proximal tubule. Shimuzu and Yamanaka demonstrated that the post mitotic, post-ischemic rat kidney have proximal tubules with an increased number of smaller tubular cells per cross sectional area than is observed in an uninjured kidney (497). Occasionally, these surplus cells can be observed to form papillary structures evaginating into the lumen of proximal tubule (497). These hyperplastic epithelial cells begin to take on the appearance of a normal proximal tubule cells by hypertrophy and differentiation and this is manifested as a decrease (back toward control levels) in the number of nuclei per cross sectional area (497), the loss of vimentin staining and the reappearance of differentiation markers (402). Concomitant with this process is a wave of apoptosis that restores normal tubular cell density that may last from between 1 week to several months of recovery (497).

IV.3.1. Proliferation of proximal tubules and potential role in recovery—The peak of proliferative activity coincides in time closely with the severity of injury, suggesting that the induction of proliferation represents an intrinsic defense mechanism. Cultured proximal tubule cells proliferate in response to a wide variety of mitogens such as epidermal growth factor (EGF) (403), hepatocyte growth factor (HGF) (523), insulin-like growth factor-I (IGF-1)(54), and fibroblast growth factors (FGFs) (639).

A significant advance was realized in a landmark paper published 1989 by Humes et al, investigating the effect of exogenous EGF on the course of kidney injury in response to ischemia reperfusion in rats. Administration of EGF to rats at the time of ischemia reperfusion injury reduced the severity of the initial injury measured by serum creatinine, and hastened the recovery back to levels of sham control levels (217). This was also associated with a significant increase in the number of mitotic proximal tubule cells in kidney of injured rats treated with EGF (217).

This finding spawned a number of similar studies investigating a variety of polypeptide growth promoting factors in various models of acute kidney injury, such as HGF (369) and IGF-I (38, 103). IGF-I was effective at ameliorating AKI when provided prophylactically to rats or when administered up to 24 hours following ischemia reperfusion injury (371). Using a dog autotransplant model, supplementation of the graft storage solution and treatment of recipients with IGF-I protected the graft and hastened return to renal function (435). In

addition to growth promoting effects, IGF-I increases renal blood flow in animals models and healthy human control subjects via a NO-dependent mechanism (207). When IGF-I was administered to late stage CKD patients it improved GFR, and was well-tolerated in most subjects (405). However, in a multicenter clinical trial, IGF-I failed to demonstrate any effect on recovery from renal function in patients with established AKI (208).

IV.3.1.1. Evidence for endogenous growth factors in renal repair: While the experiments described above provide proof of principal that growth factors may facilitate renal regeneration, the extent to which endogenous growth factor activity mediates repair response is less clear. For example, IGF-1 expression is enhanced in damaged and regenerating tubule cells (356). However, IGF-I knockout mice are not viable and there are no reports demonstrating a definitive role for this factor in the repair of the kidney (332). Similarly, while the molecular expression of hepatocyte growth factor (HGF) and its receptor c-met increase in kidneys in rat models of AKI (223) and HGF is elevated in the urine of patients with acute kidney injury (539), there is no definitive data supporting a role for endogenous HGF activity in proximal tubule injury models. Although it has been reported that a targeted mutation of the c-met receptor in collecting duct impairs recovery from transient ureteral obstruction and hastens the development of fibrosis (342).

Epidermal growth factor, EGF, is down-regulated in response to multiple different forms of AKI, and in fact the EGF gene is one of the most sensitive genes to ischemia-reperfusion injury (467). Despite its down-regulation, pre-pro-EGF present in the distal nephron is processed to an active form where it may participate in the renal repair response (476). A related growth factor, HB-EGF (heparin binding EGF), is expressed in the proximal tubule and is associated with the outer surface of the cell membrane where it can interact with the EGF-receptor (c-erb-B) in a juxtacrine/autocrine fashion. Injury activates the proteolytic processing HB-EGF to liberate a soluble form of the molecule (470). The importance of a functional EGF receptor in kidney repair was established using *waved-2* mice (*wa-2*), which have mutated EGF-receptors with ineffective tyrosine kinase activity and decreased receptor signaling. These mice manifest a significant impairment in structural and functional recovery from mercuric chloride induced AKI (594).

Fibroblast growth factors- FGFs represent a family of heparin binding growth factors with potent proliferative potential of proximal tubules. Various members of the FGF family are induced following the induction of ischemic and nephrotoxic injury. FGFs share a series of common receptors; recent studies using local administration of antisense oligonucleotides against the FGF receptor showed a moderate effect on the expression of nephrogenic genes following ischemia reperfusion (576). Although not clearly established, Klotho, a senescence related molecule appears to interact with various FGFs; Klotho facilitates the binding of FGF23 to the FGF-R1 in kidney (565). Renal Klotho expression is suppressed following AKI, but adenoviral gene transfer of Klotho is cytoprotective in AKI (526)

BMP-7 has received considerable attention for its role not only in stimulating cell proliferation, also as a potential anti renal fibrotic agent in chronic kidney disease. Exogenous BMP-7 hastens recovery from AKI (579). Evidence for the involvement of endogenous BMP-7 in renal regeneration derives from studies using USAG-1 (uterine sensitization associated gene-1) knockout mice. USAG-1 is a powerful endogenous antagonist of BMP-7 and it is prominently expressed in kidney. USAG-1 $-/-$ mice are resistant to AKI and have enhanced recovery following cisplatin AKI (614) These protective effects were abolished by the administration of BMP-7 neutralizing antibodies (614).

IV.3.1.2. Immediate early genes and cell cycle signaling: In most models of renal injury there is a rapid increase in immediate-early gene expression. Activation of immediate early genes such as JE, KC (463) and Cyr61 have been reported (387), which may participate in early growth and or inflammatory processes. In ischemia, c-fos mRNA peaks as early as 1 hour and declines within 4 hours (463). Other “immediate early genes” associated with growth responses are also induced following injury including c-jun, c-myc and Egr1 (414, 463). These genes encode transcription factors that regulate growth processes suggesting a potential role in regeneration.

Injury and cell stress activate downstream signaling pathways associated with cellular regeneration conveying prosurvival signals. The EGF-receptor (EGF-R) is activated not only in response to EGF-related ligands (as described above) but other factors depending on the nature of the injurious stimulus. For example, following replating of tubule cells, EGF receptor activation and subsequent cell proliferation depends on the release of the HB-EGF ectodomain by the metalloprotease ADAM17 (645). In contrast, in response to oxidative stress induced by H₂O₂, EGF-receptor activation and subsequent cell proliferation is dependent on the activation of Src (645).

Cell proliferation in response to injury begins with entry into the G1 phase of the cell cycle, typically under the stimulation of extracellular signals derived from growth factors or integrin-linked kinase receptors. The coordinated synthesis and degradation of cell cycle inhibitors regulate important checkpoints within the cell cycle to ensure the successful completion of cell division (468). One cell cycle inhibitor of significant interest in the renal regeneration response is p21. p21 is a cyclin dependent kinase inhibitor, which controls the G1 to S phase checkpoint. Expression of p21 is associated with terminal differentiation and senescence of many cell types, but it is rapidly induced in the kidney following various forms of acute injury in such models as cisplatin or ischemia reperfusion (360, 362). p21 ^{-/-} mice demonstrate a more rapid and pronounced level of renal injury when compared to wild-type mice, characterized by elevated BUN, and necrotic death even extending into the cortical S1 and S2 segments, and increased apoptosis (360, 362). Increased BrdU labeled nuclei were found in kidneys of p21 ^{-/-} mice, however many of these cells had a polyploidy DNA content, suggesting that the lack of p21 regulatory control results in aberrant repair due with an uncontrolled cell cycle (360, 362).

IV.3.2. Migration and extracellular matrix—Severe renal injury leaves behind denuded areas of the basement membrane and also activates a variety of proteases with the potential to degrade the tubular basement membrane. Studies by Zuk et al., (649) and by Walker (585), demonstrated a decrease in the collagen content and laminin content in the early post-injury phase, which is gradually restored over the course of several days of recovery. Given the close association between matrix and cell adhesion, cellular and matrix remodeling events are likely well coordinated, but are certainly poorly understood.

It is likely that the re-establishment of the basement membrane is critical to the re-establishment of proximal tubule function for proper presentation of integrin binding. There is increased expression of basement membrane ECM components by tubule epithelial cells such as collagen IV and fibronectin mRNA during the regeneration process following AKI (37, 650). An alternative splice variant of fibronectin (EDA, EIIIA) is expressed in the regenerating kidney in a TGF-beta dependent fashion (37); this form of fibronectin may modify cellular integrin binding and affect adhesion and cell migration (194). The significance of these findings on renal repair in vivo has yet to be established. To date, the primary method to address potential migratory activity utilizes migration assays to scrape injury in vitro. Using these approaches, plasminogen activator and EGF promote migration

and repair of wounded monolayers of proximal tubules, while TGF-beta inhibits this process (402).

IV.3.3. Recapitulation of a renal development—During renal development, the formation of a morphologically polarized epithelium from committed renal mesenchymal cells requires an external signal for mitogenesis and differentiation (188). Many cellular events are common in both renal development as well as renal regeneration such as induction of growth factor activity, high rates of DNA synthesis, apoptotic remodeling, the production of ECM, and the interaction of surface receptors leading to an early organization of actin cytoskeleton. The plasma membrane organizes to define apical and basolateral domains and the formation of cell-cell contacts results in a polarized epithelium. These similarities prompted Bacallao and Fine to hypothesize that kidney regeneration utilizes an intrinsic developmental program as a basis for recovery from kidney injury (20).

Although difficult to prove, the notion that renal regeneration recapitulates development is supported from a variety of molecular/genomic studies demonstrating that the a substantial complement of genes typically restricted to renal embryonic development are re-expressed during kidney regeneration. As mentioned above, vimentin, is expressed in regenerating epithelia as well as developing epithelia during fetal development (28). However studies examining specific gene pathways as well as studies employing mRNA expression profiling have revealed a substantial number of kidney developmental genes expressed during renal regeneration; these genes include those involved in growth regulation, extracellular matrix factors such as FN EIIIA, transcription factors Pax2 and Wnt-4 (225, 540).

Epigenetic mechanisms including histone acetylation play a key role in organ development and cellular homeostasis by regulating gene expression in a tissue- and developmental stage-specific manner (530). Increasing histone acetylation, the levels of which are determined by the balance between the activities of histone acetyltransferase (HAT) and histone deacetylase (HDAC), generally stimulates gene transcription by relaxing the chromatin structure. Recent studies by Murano et al., suggest that modification of HDAC activity may provide an important regenerative feature of damaged kidneys. In cultured proximal tubules subjected to energy depletion, there was a decrease in histone acetylation and HAT activity. During recovery, the HDAC isosyme HDAC5 was downregulated as histone acetylation returned. Reduced HDAC 5 levels were associated with an increase in BMP7 levels. In vivo, HDAC expression was repressed in regenerating tubules on the outer medulla of post-ischemic mice kidneys, where BMP-7 levels were elevated (352)

In addition to effects on potential repair mechanisms, it has recently been shown that the transcription factor ATF3 interacts directly with histone deacetylase 1 (HDAC1) and recruits HDAC1 into the ATF/NF- κ B sites in the IL-6 and IL-12b gene promoters. Based on studies using ATF1 knockout mice, this interaction results in the condensation of chromatin structure, resulting in the interference of NF- κ B binding, and inhibition of inflammatory gene transcription after I/R injury (309).

The epigenetic control of gene expression regulating injury and repair processes in the setting of AKI is likely to be the source of significant new research findings in the coming years. Such advancements may likely provide a new viewpoint by which to understand events underlying remodeling occur. As suggested by Surendren and Kopan, the flexibility of histone remodeling may explain how various cell types can alter their differentiation status and re-enter the cell cycle, restore normal morphology, and provides a basis for regenerative processes to occur without a requisite requirement for stem cell niches (530).

IV.3.4. Tubular cells comprise the majority of the regenerative cells during repair—Interest in stem cells as possibly participating in kidney regeneration derives historically from observations in kidney transplant patients where host cells have been reported to incorporate into the renal tubules of the graft (181). Thus, populations of circulating stem cells, presumably bone marrow derived, could home to the kidney and participate in renal recovery. However, the identification of these cells provides little insight into their participation in overall kidney repair.

Some reports have suggested that BM derived cells may transdifferentiate into tubules or may fuse with endothelial cells following injury (82, 324). However, the proportion of bone marrow derived cells that integrate stably into tubular epithelial cells or endothelial cells is small. In a study by Lin et al., a transgenic cre-lox approach was used to label kidney epithelial cells and track their progeny following I/R injury. They demonstrated that BrdU was localized in dedifferentiated cells that expressed the Cre-induced transgene, suggesting that the source of regenerating cells was resident, non lethally injured epithelial cells (Figure 12). In the same study, this group demonstrated transplanted bone marrow GFP+ cells localized to the kidney interstitium one month after injury, but not to renal tubules (325). Other groups have subsequently reaffirmed that the primary source of new cells during regeneration is sublethally damaged kidney tubule cells (219), while no studies using bone marrow transplants have demonstrated that such cells comprise a significant source of regenerated epithelium. Thus, there is little remaining controversy regarding the source of new cells during the regenerative process post-AKI. However, as described below, it remains possible that circulating stem cells may augment renal repair mechanisms and that stem cell niches within the kidney may participate in regeneration.

IV.3.5. Potential contribution of bone marrow derived stem cells—Despite the lack of convincing evidence that bone marrow derived cells contribute to the formation of new cells during renal repair, recent studies clearly suggest a beneficial role for bone marrow derived cells in aiding the regeneration process and provide a possible therapeutic modality. Two distinct stem cells reside in the stroma of the bone marrow; the hematopoietic stem cell (HSC), undifferentiated cells capable of self renewal and giving rise to the specialized cells of the blood including, erythrocytes, thrombocytes and leukocytes. Mesenchymal stem cells (MSC) are of mesodermal origin and are characterized by the expression of CD90, CD73, CD104, CD44 and CD29 but do not express markers of hematopoietic lineage, such as CD34 and CD45 (82). These multipotent cells give rise to cells of connective tissue such as bone, fat, cartilage and muscle.

IV.3.5.1. Hematopoietic stem cells in renal repair: Generally speaking, HSCs are not thought to contribute prominently toward a repair process in the setting of AKI. Some reports have demonstrated that HSC may incorporate into and differentiate into renal tubular cells following injury, however the number of cells of this nature is quite small (324). Typically, administration of undifferentiated HSC or their mobilization worsens kidney injury by exacerbating the extent of renal inflammation (549, 550). In contrast, other reports have demonstrated a beneficial effect of HSC mobilization repair (230). The reason for the discrepant reports is unclear but may relate to the specific methodologies used to induce mobilization.

In addition, HSC may also give rise to a population of cells referred to as endothelial progenitor cells, or EPC. EPCs represent a population of cells derived from HSC which express proteins common to endothelial cells (e.g., VEGFR-2 and V-cadherin), but also markers of hematopoietic lineage (e.g., CD34 and CD133) (455). There is significant confusion in the literature regarding the nature of these cells as the term EPC is likely a misnomer. These cells do not give rise to mature endothelial cells and do not stably integrate

into blood vessels. However, they likely do help facilitate vascular growth and repair processes and therefore may be properly termed “angiogenic” macrophages (623).

Regardless of the terminology, these cells can home transiently to sites of vascular injury and can produce pro-angiogenic factors thought to aid in the angiogenic process and influence mature endothelial cells (455, 623). Several clinical studies suggest that host EPC may migrate and incorporate into tissue grafts following renal transplant (455) but the importance of this observation on maintenance of renal function and repair is not known. In a mouse model of ischemia reperfusion injury Patschan et al, demonstrated that ischemic preconditioning increased the deposition of CD34+/Flk+ cells which are also positive to c-kit/and Tie2. When these cells were isolated and administered to naïve mice, they preserved renal vascular function in response to ischemia reperfusion (426). Recently, Li et al., demonstrated the human CD34+ HSC accelerated recovery following I/R injury in immunocompromised mice (307). Administration of factors such as erythropoietin or uric acid may serve as a trigger to increase the number of EPC migrating to the kidney and have a beneficial effect on vascular injury (209, 427).

IV.3.5.2. Mesenchymal stem cells in renal repair: In contrast to HSC, MSC have received considerable attention in the setting of acute kidney injury. MSC do not elicit immune responses, may be harvested from non bone marrow depots (such as adipose) and can be expanded in vitro making them potential useful therapeutically (82, 557). Several groups have reported a beneficial effect of MSC to experimental animals following AKI. MSC treatment provides protection when administered at the time of injury and hastens repair after established AKI induced by I/R (385, 553). Despite the differentiative potential of MSC there is no evidence to suggest that these cells integrate stably into the repairing kidney. Togel et al., demonstrated that MSCs traffic only transiently in the post-ischemic kidney declining within 1 day with the majority of cells homing in lung and spleen (552). Therefore, it is suggested that the reparative effects of MSC is via the production of endocrine or paracrine factors. For example, MSC produce high amounts of VEGF and IGF-I, which when repressed via siRNA impair MSC effects on recovery from AKI (551). It is likely that increasing MSC homing will enhance the regenerative properties provided by MSC. Following glycerol induced injury, MSC homing was suggested to be due to CD44 interaction with hyaluronic acid as inhibition of CD44 on transplanted cells resulted in less homing and less repair (204). Homing to the kidney may not required at all; Lee et al., have recently demonstrated that MSCs after homing to lung activate the production of the anti-inflammatory peptide, TSG-6, which facilitates recovery from myocardial ischemia (293). Recently, clinical trials with promising potential have been initiated and may represent the advent of a novel therapeutic avenue in the treatment of AKI (553).

Whether endogenous MSC might participate in renal repair has not been adequately addressed. However, recent studies from Goligorsky’s group have identified a population of MSC that lie within the renal capsule. Cell-tracker labeling in vivo demonstrated that these cells have the ability to proliferate and preferentially home to the ischemic perivascular space in the outer medullary region within 3 days of I/R injury. Decapsulation of the kidney modestly attenuated the recovery from renal injury suggesting a partial role for endogenous MSC in renal repair (423).

IV.3.6. Renal derived stem and progenitor cells—Other investigations have been directed toward the concept that renal repair is mediated via intrinsic renal stem and/or progenitor cells following injury. Stem cells, by definition are clonogenic cells capable of both self renewal and multilineage differentiation. In contrast, progenitor cells are defined as immature and proliferative cells which are limited in their differentiation potential to only one cell type (460). A lack of clear markers on what defines a population of renal stem/

progenitor cells, as well as the possibility that there are likely multiple populations of cells with progenitor activity have contributed to some controversy in this area. To date, several different stem and progenitor cells have been identified and may participate in kidney repair in animal models.

For example, Bussolati et al., have isolated cells from human and animal kidneys expressing CD133, a well accepted stem cell marker, using magnetic sorting techniques (81, 82). These cells lack hematopoietic markers but express the renal developmental transcription factor, Pax-2. These cells also show limited self renewal and could differentiate into either epithelial or endothelial cells in culture (81, 82). Administration of these cells into SCID mice with glycerol induced tubular injury resulted in their homing and their integration into tubular structures (81, 82). However, there is no direct evidence that CD133 expressing cells within the kidney participate substantially in repair in response to an injury.

Gupta et al., isolated a unique population of cells from adult rat kidney cells that displayed features of a multipotent renal progenitors (termed MRPC). These cells exhibit self renewal and expression of vimentin, CD90, Pax2 and Oct 4 without evidence of senescence. Oct4 expression was seen in some tubular cells in the adult kidney, suggesting these cells may be candidate renal stem cells. The cells can differentiate into renal tubules when injected under the capsule of an uninjured kidney or intra-arterially after renal ischemia-reperfusion injury (182). Again, evidence that endogenous MRPC participate in the injury process has remained elusive.

Finally, a population of progenitor cells has been identified in the renal papilla based on their ability to retain BrdU label when administered during the early embryonic or neonatal development (345, 411). These “label retaining cells” (LRC) cells can be isolated and expanded in culture and display activity consistent with stem cells as they give rise to multiple different cell types (345, 411). There are multiple conflicting reports on the fate of LRC that may be based in part on different protocols used to identify them. In one report, Maeshima infused BrdU into rats for 7 days followed by 2 weeks of chase. Subsequent injury by ischemia reperfusion resulted in a transient increase in the number of LRC many of which were dedifferentiated and expressed markers of proliferation (346). In contrast, when LRC were labeled in mice by embryonic BrdU administration and studied at 2 months of age, LRC were found in the outer stripe of the outer medulla but most Ki67 cells did not colocalize with these LRCs (269).

IV.3.7. Impaired recovery in aging and CKD—Elderly patients are more prone to develop AKI and to have a less successful outcome (93, 227, 479, 480). It needs to be emphasized that most of our understanding of the reparative responses of the kidney derive from studies on young healthy animals and such mechanisms are likely not as efficient in older animals (479).

Studies in rats and mice have demonstrated an impaired recovery response in older animals vs. younger animals (89, 481). The increased susceptibility is likely the result of multiple factors including impaired vascular reserve, as well as an impaired regenerative capacity. In older animals in response to injury, less cell proliferation was observed and the intrinsic growth potential (i.e., colony forming potential) of proximal tubule cells was significantly reduced in older animals vs. younger animals (481). A comparison of gene expression demonstrated that tubules from older animals express greater amounts of zinc α 2 glycoprotein-1 (Zag-1), a putative inhibitor of proliferation (481). Chen et al (89) demonstrated that the age related sensitivity to injury could be ameliorated by caloric restriction. Up to 92 different genes were altered in the aging kidney that were attenuated by with caloric restriction including genes associated with tight junction activity (Claudin-1)

and genes associated abnormal matrix regulation, such as MMP-7. Aging likely influences a number of processes regulating epithelial growth potential and/or senescence.

Shortened telomeres are associated with cell senescence and the biological significance of this in recovery from AKI has been studied in a mouse model of telomere shortening. Mice with null mutation in telomerase gene (*Terc* $-/-$) develop significantly shortened telomeres within 4 generations and have increased sensitivity to AKI. Moreover, these mice manifest a less successful recovery response that is associated with interstitial fibrosis. At the molecular level, there is an increased expression of the cell cycle inhibitors p21 and p16^{INK4a} relative to age matched wild type controls (604). Interestingly, the cell cycle regulator p16^{INK4a} is highly expressed in aging human kidney; following renal transplant, its expression is elevated in failing renal grafts while it low transplant with immediate function (364).

Impaired recovery from renal damage may also be a function of pre-existing renal disease. Epidemiological studies suggest a less successful recovery of individuals from AKI with established CKD (227). Although early impairment in regional blood flow is worsened in diabetic db/db mice, there is a paucity of studies investigating renal repair potential in models of CKD (496). However, several studies have investigated recovery of AKI in conjunction with reduced renal mass. Such studies show evidence of impaired recovery characterized by incomplete return of GFR and impaired differentiation of regenerating proximal tubules (617). As described below, impaired recovery may relate in part to the predisposition of chronic kidney disease following AKI.

IV.4. Chronic effects of AKI

Incomplete repair or conversion from physiological to a pathological repair response could result in declining renal function and chronic kidney disease following AKI. Although typically AKI is not thought to predispose the development of long-term kidney complications, it has been known for several decades that many adult patients who suffer from AKI never recover renal function completely, manifest decrements of GFR and have impaired urinary concentrating ability (58, 71, 147, 275, 304). Pediatric patients have a particularly high risk of developing CKD and hypertension following AKI (8, 16). As the incidence of AKI has risen, so too has interest in the long-term sequelae of AKI and several studies now suggest patients with AKI progress to ESRD (9, 97, 227, 543).

From a pathophysiological perspective, animal model data support the connection of AKI with CKD and offer potential mechanistic insights into the process. Nath et al., used a model of cutaneous glycerol injection to induce AKI resulting in a significant but reversible decrease in creatinine clearance in rats. Repetitive administration of glycerol (once monthly over 6 months) resulted in a slow progressive loss of GFR associated with increased TGF- β expression and progressive interstitial fibrosis (394). The development of CKD occurred despite the presence of an adaptation to the subsequent repetitive glycerol injections, (i.e, preconditioning) suggesting a disconnect between the acute and chronic manifestations of this insult. Moreover, the results imply that CKD may result from a series of small repetitive acute injuries that may not manifest clinically as AKI.

Single episodes of AKI have the potential to alter renal function on a permanent basis. Chronic reductions in urinary concentrating ability have been observed in post AKI rats following either cisplatin or ischemia reperfusion induced injury (32, 465). Although early elevation in diuresis may be due to increased Na excretion or impaired aquaporin activity, concentrating ability remains chronically impaired even following the return of %FENa to sham levels and the resolution of the response to dehydration stress. Rather it was suggested that the urinary concentration defect persisted due to impaired ability of the kidney to generate a hypertonic medullary gradient (32, 465).

Following a single bout of ischemia reperfusion, proximal tubule repair is typically complete within 4–6 weeks in rats. The renal expression of TGF- β and its cognate receptors are elevated transiently and return to control levels within approximately 4 weeks. With increasing recovery times (16–40 weeks), features of CKD ultimately become manifest, including proteinuria, interstitial fibrosis associated with a secondary increase TGF- β (32).

Renal regeneration is less complete and progression of CKD more rapid and severe when AKI is carried out in the setting of reduced renal mass (31, 167, 416, 574). In an elegant study by Patgulan et al., morphometric analysis of ultra thin serial sections revealed AKI resulted in nephron dropout, presumably due to loss of contact between the injured tubule and its parent glomerulus (416). Post-AKI kidneys in reduced renal mass models have a high proportion of dilated tubules containing cells which have not undergone redifferentiation. Venkatachalam et al., have recently focused attention on the existence of dilated tubules containing dedifferentiated epithelial cells and suggested that these may play a central role in the conversion AKI to CKD. According to the hypothesis, dedifferentiated cells express soluble profibrotic factors which act initially in the physiological repair response; however sustained expression of such factors (e.g., PDGF) may switch the normal repair process to a pathologic/fibrotic response (574). It is possible this relates to the recent observation by Yang et al., in which cells that are growth arrested in G2/M express high levels of profibrotic cytokines such as TGF-beta and CTGF (617). It is currently unclear why dedifferentiated cells are more prominent when regeneration occurs in the presence of reduced renal mass. One possibility may rest with a persistent expression of the TGF-beta axis, which may inhibit redifferentiation of proximal tubule cells (167). However, contributory factors related to reduced renal mass such as exacerbated hypoxia, increased SNGFR and/or exhausted renal compensatory reserve are additional possibilities that remain to be explored. Epigenetic factors may influence gene expression in the post injury state; in a recent report, Zager and colleagues demonstrated that histone modifications may be responsible for sustained profibrotic gene transcription leading toward CKD following an I/R insult (636).

IV.4.1. Impaired vascular recovery following AKI: a possible contributor

toward CKD—Although total renal blood flow typically returns to baseline values between 1–4 weeks of recovery, our laboratory hypothesized that there is chronic impairment of medullary vascular function following AKI. This hypothesis was based on early pathological observations suggesting vasa recta capillaries from post AKI patients were separated from nearby cells of the loops of Henle, potentially compromising the efficiency of the processes driving the generation of the medullary concentration gradient. Using microfil to generate vascular casts, we demonstrated between a 30–50% reduction in the total number of microvessels, most pronounced in the inner stripe of the outer medulla, after recovery from AKI (Figure 13) (32). It is important to note that such alterations in renal blood vessel density do not necessarily indicate a reduction in total RBF and GFR as these values were typically restored to normal levels when vessel density measurements were made (102, 250, 629). Recently published studies have shown reductions in peritubular capillary density in transplanted renal grafts; in these studies, the degree of capillary rarefaction is more pronounced in grafts experiencing delayed function (283, 510).

These observations indicate that the regenerative capacity of the vasculature is reduced relative to that of the tubular epithelium; this impaired vascular regeneration may predispose CKD via a number of possible mechanisms. Hypoxia is a well-known contributor to interstitial fibrosis and is thought to directly stimulate the expression of genes with profibrotic activity (404). Recovery from AKI was shown to be associated with persistent hypoxia using the 2-pimonidazole technique. AKI in combination with reduced renal mass (unilateral nephrectomy) exacerbated hypoxia and accelerated the development proteinuria

and interstitial fibrosis. Chronic L-arginine supplementation beginning after the establishment of injury, did not protect microvessel dropout, but did increase renal blood flow, reduced hypoxia and attenuated the development of interstitial fibrosis and proteinuria (31).

In addition to its role in hypoxia, reduced medullary blood flow is a characteristic of many models of hypertension (104). Johnson and Schreiner suggested a cause and effect relationship between peritubular capillary loss and salt sensitive hypertension in response to various acute insults (239). When post-AKI rats are provided elevated dietary salt after the completion of the initial recovery response (5 weeks), hypertension develops along with substantial interstitial fibrosis (519). There were no differences between post AKI and sham animals in terms of plasma renin activity or circulating Ang II levels. However, when normotensive post-AKI rats were analyzed acutely after 5 weeks of recovery, there was a rightward shift in the acute pressure natriuresis relationship. In addition, there was impaired medullary flow and interstitial hydrostatic pressure responses to elevations in renal perfusion pressure (429). Thus, impairment of the renal microvasculature represents a primary cause of salt-sensitivity in post AKI rats.

Administration of VEGF-121, a non-heparin binding form of VEGF, can attenuate the loss of renal microvessels, and/or may induce new vascular cell proliferation in various models of progressive renal fibrosis such as aging or reduced renal mass via 5/6 nephrectomy (247, 248). VEGF-121 administration did not affect the initial loss of renal function (i.e., serum creatinine) following bilateral ischemia reperfusion injury, but did mitigate the loss vascular density measured at 5 weeks of recovery. VEGF 121 treatment ameliorated the secondary salt sensitive manifestations of CKD following AKI (302). Importantly, VEGF-121 did not protect against CKD when administered after 3 weeks of recovery, suggesting that there is an effective window of opportunity for intervention in the early injury/repair process. A similar protective effect on capillary preservation and subsequent fibrosis was observed in post-ischemic mice treated with cartilage oligomeric matrix protein/angiopoietin-1 (COMP-Ang1) a variant of the native angiogenic factor, angiopoietin 1 (242). Taken together, these data highlight the potential importance of the renal microvascular stability in the function of the post-AKI kidney.

IV.4.2. The basis of vascular loss and impaired recovery following acute injury

—There is no clear consensus regarding the cellular basis of vascular cell loss following injury, which complicates further our lack of understanding of suboptimal vascular repair responses. There is an early alteration in endothelial structure that promotes vascular leakiness (534), but it is not yet clear if this is related to permanent alteration in vascularity. It is certainly reasonable to suggest that apoptotic cell death underlies vascular (i.e., endothelial) cell loss following the induction of acute injury. Lee et al., demonstrated peritubular capillary endothelial cell apoptosis following severe liver I/R in mice which results in a subsequent AKI (292). However, Hörbelt et al., used Tie-GFP mice to label renal endothelial cells and were not able to identify apoptotic endothelial cells in response renal ischemia reperfusion, but did identify apoptotic endothelial cells in response to CD95 ligand. Therefore, it seems that the mode of cell loss (apoptosis or not) may be due in part to the nature of the underlying injury (212).

We recently provided evidence that endothelial mesenchymal transition, a process by which endothelial cells transform into a mesenchymal phenotype, may contribute to vascular cell loss following AKI. There was a significant increase in cells that co-localize both endothelial markers (CD31 or cablin) and mesenchymal markers (S100A4) following I/R injury (35). Fate tracing of endothelial cells using Cre-Lox labeling indicated that interstitial

cells observed 2 weeks following injury were of endothelial origin and the presence of these cells were reduced by treatment with VEGF-121 (35).

Finally, Lin et al, have posited that vascular loss may be mediated by a loss of the critical interaction of the endothelial cells with their neighboring pericytes in response to injury (328). In this hypothesis, the loss of trophic support from surrounding pericytes predisposes the loss of endothelial cells and results in impaired remodeling.

Regardless of the mode of cell loss, the persistent reduction in vascular density highlights the lack of regenerative capacity of renal vascular cells. The basis for the resulting lack of repair may stem from an impaired cellular responsiveness and/or combination with a lack of appropriate growth and/or trophic support factors and evidence exists for both possibilities. Indeed, in contrast to the enhanced expression of factors promoting the growth of epithelial cells, AKI induces a shift toward factors that may prohibit endothelial growth. TGF- β is a negative regulator of endothelial proliferation and, in certain endothelial cell culture systems, induces endoMT. TGF- β is prominently expressed during the regeneration process in ischemic and nephroxic AKI models (38). Inhibition of TGF β using antibodies preserved vascular density and attenuated the deposition of peritubular fibroblasts (520). In addition, other inhibitors of endothelial cell growth are expressed following AKI including angiotensin, sFLT, ADAMTS-1, endostatin, arrestin and canstatin (33, 34, 212).

Renal tubular expression of VEGF is thought to be the primary source of local vascular trophic support. Reductions in VEGF expression as a function of age correlate with vessel loss (544). Similarly, other models of CKD are characterized by reduced VEGF and loss of peritubular and/or glomerular vessels (239, 404). Because hypoxia is considered a primary trigger for VEGF expression via the HIF1 α system (183), it is surprising that VEGF expression decreases rather than increases in developing CKD. Following AKI, VEGF expression has been reported to be either unchanged or transiently reduced (33, 246). It is possible that a transient reduction in VEGF expression may underlie vascular dropout since administration of VEGF-121 could attenuate vascular loss when administered within the first few days of ischemia (302). Although the balance of research done to date indicate an increase in vascular destabilizing factors and a decrease in pro-angiogenic factors following AKI, angiopoietin 1, a vascular stabilizing factor was reported to be prominently and persistently induced in the mouse kidney following AKI (212).

Impaired vascular growth may also be due to impaired cellular responses to paracrine cues. Indeed, while new cell proliferation of glomerular endothelial cells has been demonstrated in the early stages of diabetic glomerulopathy or other models of glomerular injury (151, 249), there is little evidence that peritubular capillary endothelial cells undergo proliferation in disease states. In a recent study, no BrdU positive endothelial cells were detected within 2 days of ischemic AKI, and less than 1% of BrdU cells labeled endothelia after 7 days of continuous BrdU administration. VEGF-121 preserved vessel integrity but did not influence endothelial proliferation (35); this suggests that VEGF protects via non proliferative mechanisms (e.g., perhaps blocking transition), but also suggests that renal endothelial cells have intrinsically poor growth potential. It is also noteworthy that endothelial cells of the kidney are notorious for being difficult to grow in culture. The difficulty in growing renal endothelial cells may reflect an intrinsic impairment of their growth potential relative to other endothelial cells derived from different vascular beds.

IV.4.3. Fibroblasts and myofibroblasts—AKI is associated with the deposition of interstitial fibroblasts or myofibroblasts, an important factor necessary for the development of interstitial fibrosis (520, 574). The source of these cells may be secondary to proliferation, transdifferentiation and/or migration or infiltration. Studies investigating the origin of renal

fibroblasts can be complicated by the use of different markers for these cell types and controversy regarding their appropriateness. Nevertheless, some rather interesting concepts are worth summarizing.

The Ca⁺⁺ binding protein S100A4 has also been termed fibroblast specific protein-1 (FSP-1) and has been proposed as a marker of interstitial fibroblasts. In a classic series of studies by Neilson's group, overexpression of FSP-1 in cultured proximal tubules led to the transformation of these cells into a fibroblastic phenotype (408). This led to the suggestion that fibroblasts derive from epithelial cells via a process called epithelial mesenchymal transition (EMT), contributing to the development of interstitial fibrosis (245). Moreover, colocalization of FSP-1 with epithelial markers was subsequently considered suggestive, but not definitive evidence for EMT. A more definitive approach uses epithelial cells labeled with Cre-Lox approaches for fate tracing; When tubular epithelial cells are labeled with a reporter gene using Cre-Lox, ureteral obstruction resulted in interstitial expansion of labeled cells which stained positively for FSP-1 (229). However, other investigators using similar approaches have not been able to confirm these data (218), so the role of EMT in interstitial fibrosis represents a controversial area.

In ischemic AKI, a role for EMT is not well-supported by the literature. In the study described above by Lin et al., Cre-lox labeled tubular epithelial cells were shown to be the source of proliferating immature cells during regeneration. However, the study is also important as it demonstrated that labeled cells remained restricted to the tubule after the final recovery response and did not migrate to interstitium, strongly arguing against EMT (325). In contrast to epithelial mesenchymal transition, recent evidence from our laboratory suggested that endothelial mesenchymal transition (EndoMT) may not only contribute to vessel dropout, but also deposition of interstitial cells (35).

It is possible that fibroblasts/myofibroblasts derive from the circulation and target the injured kidney, perhaps to participate in physiological or pathological repair. Bone marrow transplant of transgenic animals expressing a trackable marker (i.e, beta gal, or GFP) demonstrate bone marrow derived cells in the interstitium following injury, some of which express S100A4 (75). These cells may be macrophages, since S100A4 expression has been described in this cell type. In contrast, these cells may also represent a form of a circulating cell type known as the fibrocyte (580).

An additional hypothesis recently put forward for the genesis of myofibroblasts was recently articulated by Humphries et al. in which a cre expression was driven in non SMA positive pericytes using the Col I promoter (218). Ureteral obstruction promotes SMA expression in these pericytes accounting for a large number of interstitial myofibroblasts. It will be of interest to see if other models of AKI are also associated with expansion and activation of pericytes as a prominent feature of fibrosis.

IV.4.4. Contribution of immune and inflammatory factors—It is clear that immune and inflammatory factors participate in the progressive process following AKI. Not surprisingly, a number of studies derived primarily from the transplant literature demonstrate that immune suppression blocks the expansiveness of nephropathy following an acute injury. A comprehensive assessment of the long term immune responses following AKI is beyond the scope of this review, however there are a few important points worth highlighting. An important role of both lymphocytes and macrophages has been demonstrated in progression following acute injury. Forbes et al, demonstrated that an early treatment with an endothelin antagonist blocked the deposition of ED-1 positive macrophages and attenuated the late manifestations of acute kidney injury (153). Secondly, there is clearly an important yet unexplained role for T cells in CKD development following

AKI. In a classic study by Chandrakar et al, the B7 costimulatory pathway was blocked one month following unilateral I/R with reduced renal mass, resulting in a dramatic decrease in the development of proteinuria and interstitial fibrosis with recovery times up to 6 months (86). Burne-Taney et al carried out adoptive transfer of splenocytes from post-I/R or sham-operated mice and followed their renal function over time. Splenocytes from injured mice promoted proteinuria non injured recipients within 12 weeks of injury, suggesting that immune responses were sufficient to mediate some of the abnormal responses (79). Finally, Pechman et al, blocked lymphocyte activity with mycophenolate mofetil beginning 5 weeks following recovery from I/R injury. This treatment prevented the development of hypertension and the dramatic increase in interstitial fibrosis concomitant with elevated sodium intake (428). Taken together, it would seem that the full manifestation of CKD following AKI relies on several important features working together including, improper tubule repair, vascular dropout, and persistent activation of immune function. Activities for progression are fueled by features such as reduced renal mass, elevated sodium intake and perhaps, even dietary proteins sources or underlying genetic factors. A schematic summarizing these events is shown in Figure 14.

Conclusion

Acute kidney injury remains a vexing problem in the clinical setting. The last four decades of research have substantially advanced our understanding of the cellular and pathophysiological processes in AKI and the recovery of organ function following established injury. Such work has provided an ample number of potential therapeutic targets yet effective therapies have remained elusive. What then, are the challenges in AKI research that should be addressed in the coming years? In our opinion, technological advances in genomics, proteomic and engineering will be at the forefront of exciting new work on the field of AKI. However, such technologies will likely be focused on addressing established problems not yet adequately addressed. For example:

1. Studies that will identify renal injury earlier and that will more precisely stratify patients based on the stages of injury will aid clinical and preclinical targeting of important mechanisms of kidney dysfunction in AKI. Factors controlling the initial fall in GFR may not be the same as those leading to sustained reductions in GFR during the extension and maintenance phases of injury and as described previously, the assessment of renal function based on surrogate markers of renal injury and GFR is fairly imprecise. Markers such as creatinine change slowly over time following injury to the kidney (138, 363). Improved methods for evaluating GFR and a more rapid assessment of renal injury are likely to improve the ability of the practicing nephrologist to diagnose AKI earlier and perhaps identify the clinical course more accurately (96). Based on our current understanding of the clinical phases of injury and repair processes, a more reliable charting of the course of individual patients may allow therapies tailored for each phase.

Over the past decade, significant advancements in the field of clinical proteomics have been targeted toward identifying novel plasma and urine proteins that may characterize the course of renal injury. The advent of the field of biomarkers and its potential impact on AKI is vast and a thoughtful consideration of its role in diagnosis and treatment options on AKI is beyond the scope of this chapter. However, it must be pointed out that potential markers such as NGAL, KIM-1, cystatin C, IL-18 and NAG have all been postulated to represent early markers of renal injury and the reader is encouraged to seek reviews dedicated to this topic (138, 422, 504). Of note, NGAL, a protein that is upregulated in the proximal tubule following acute injury, and KIM-1, a membrane glycoprotein whose

ectodomain is shed into the urine following renal injury, have received a significant attention and potential early markers of renal injury.

This field has moved rapidly in the last decade from identification of these molecules in the preclinical models, to the initial validation in limited clinical settings (47, 315, 331). Although these studies suggest a potential predictive power, the next several years should provide important information on whether such tools will enable the clinician to initiate treatments earlier and to determine if stratification of patients based on such markers will provide any substantive impact in terms of outcome.

2. Studies that provide a better understanding of the link between tubular injury and the fall in GFR need to be performed. Crosstalk between tubules, blood vessels/ endothelium, inflammatory cells, and resident supporting cells may be key in defining this area and warrant further investigation.
3. A greater recognition of the simplicity of our preclinical models and a more precise understanding of the factors in AKI that lead to death, including but not limited to issues related to organ cross talk and the development of therapies geared toward sustaining multi organ function.
4. A continuation of work on recovery response of AKI will continue to be of significant interest, and important advances will likely be influenced by advances genomic and epigenetic technology. In our opinion, an understanding of the factors that lead from physiological repair to a pathological response leading to CKD will represent an increasingly important area; therefore, work leading to the identification of patients at high risk for developing AKI may help to improve long term outcomes. All in all there is still much work to be done. However, recent efforts provide optimism that some of the long standing hurdles to progress may soon be overcome.

Acknowledgments

The authors are supported by NIH grants DK063114 (DB), and DK-077124 (TS) and pilot funding from the George M O'Brien grant, P30 DK79312 (Bruce Molitoris, Program Director), and Bridge Funding from the Indiana University Research Foundation.

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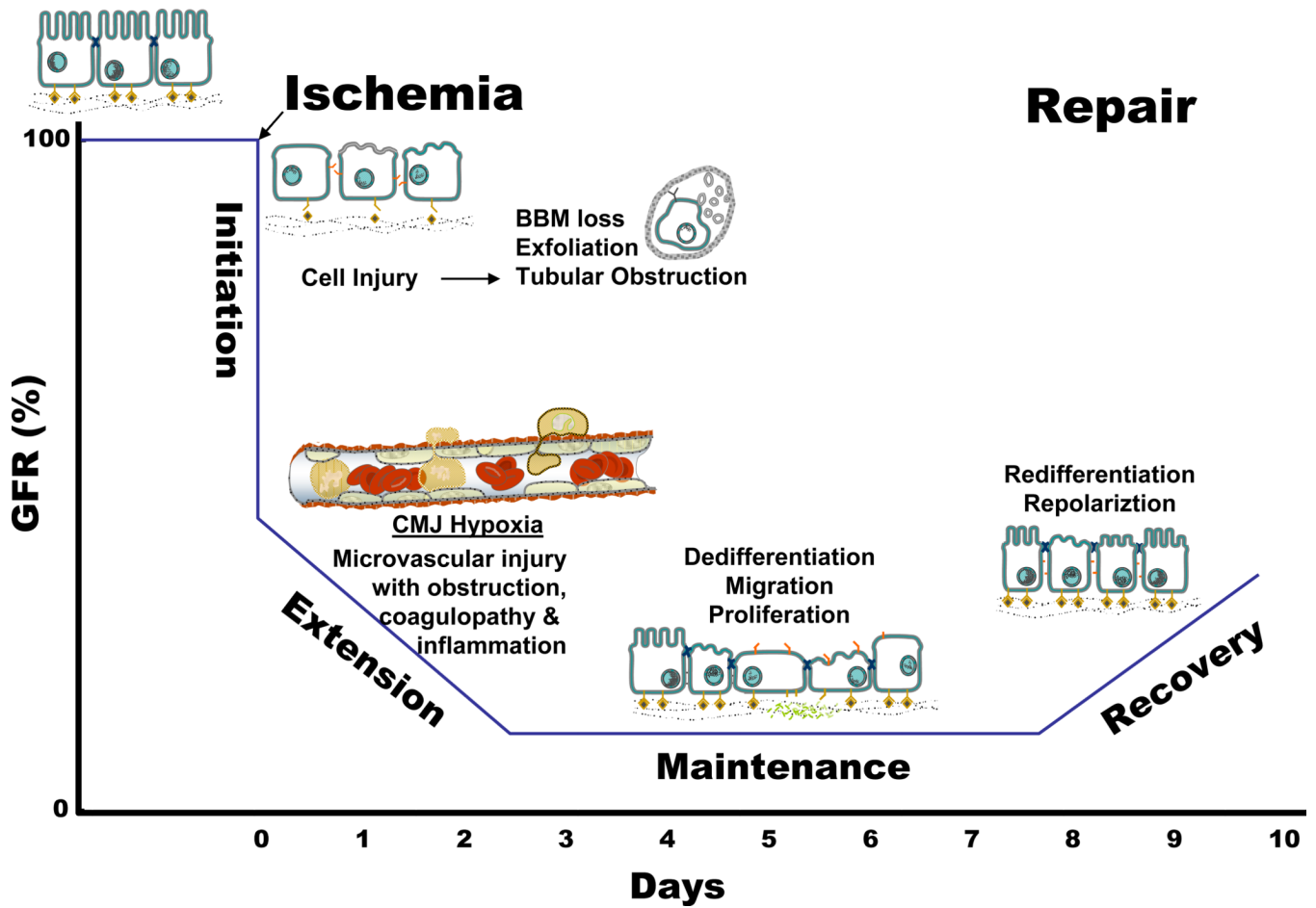


Figure 1. Relationship between the clinical phases and the cellular phases of ischemic acute kidney injury (AKI), and the temporal impact on organ function as represented by glomerular filtration rate (GFR)

Prerenal azotemia exists when a reduction in renal blood flow causes a reduction in GFR. A variety of cellular and vascular adaptations maintain renal epithelial integrity during this phases. The initiation phase occurs when a further reduction in renal blood flow results in cellular injury, particularly the renal tubular epithelial cells, and a continued decline in GFR. Vascular and inflammatory processes that contribute to further cell injury and a further decline in GFR usher in the extension phase. During the maintenance phase, GFR reaches a stable nadir as cellular repair processes are initiated in order to maintain and re-establish organ integrity. The recovery phases is marked by a return of normal cell and organ function than results in an improvement in GFR. Adapted from (532).

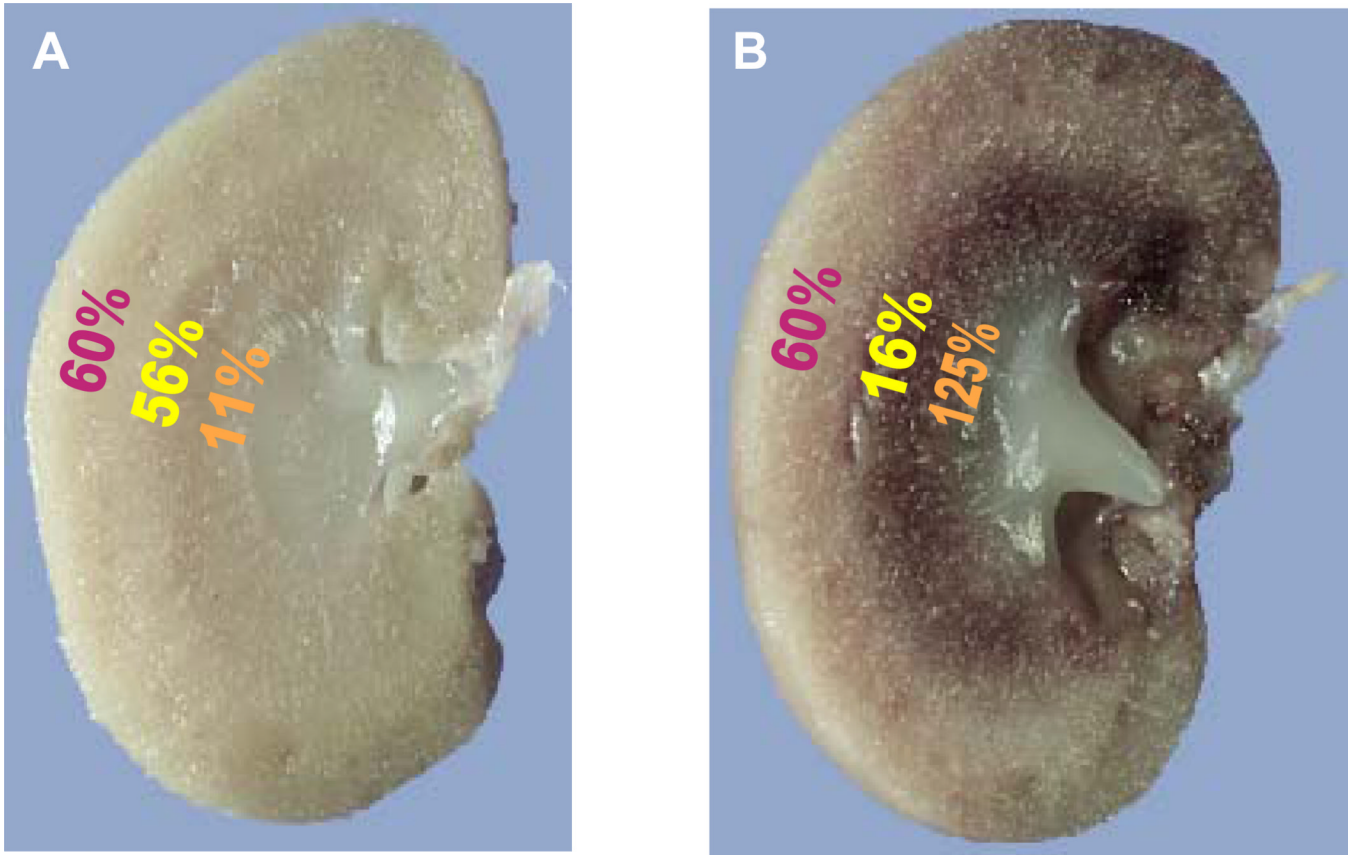


Figure 2. Regional blood flow is altered following injury in ischemic AKI

Immediately following ischemic injury total renal blood flow is reduced but more striking are the regional deficits in blood flow that exist in the cortex, outer stripe of outer medulla and inner stripe of the outer medulla as indicated in panel A (data from (251)). As overall blood flow starts to recover in the ensuing hours after injury, profound regional alterations in blood flow remain with progressive and profound reduction of the blood flow to the outer stripe of the outer medulla as indicated in B (data from (202)).

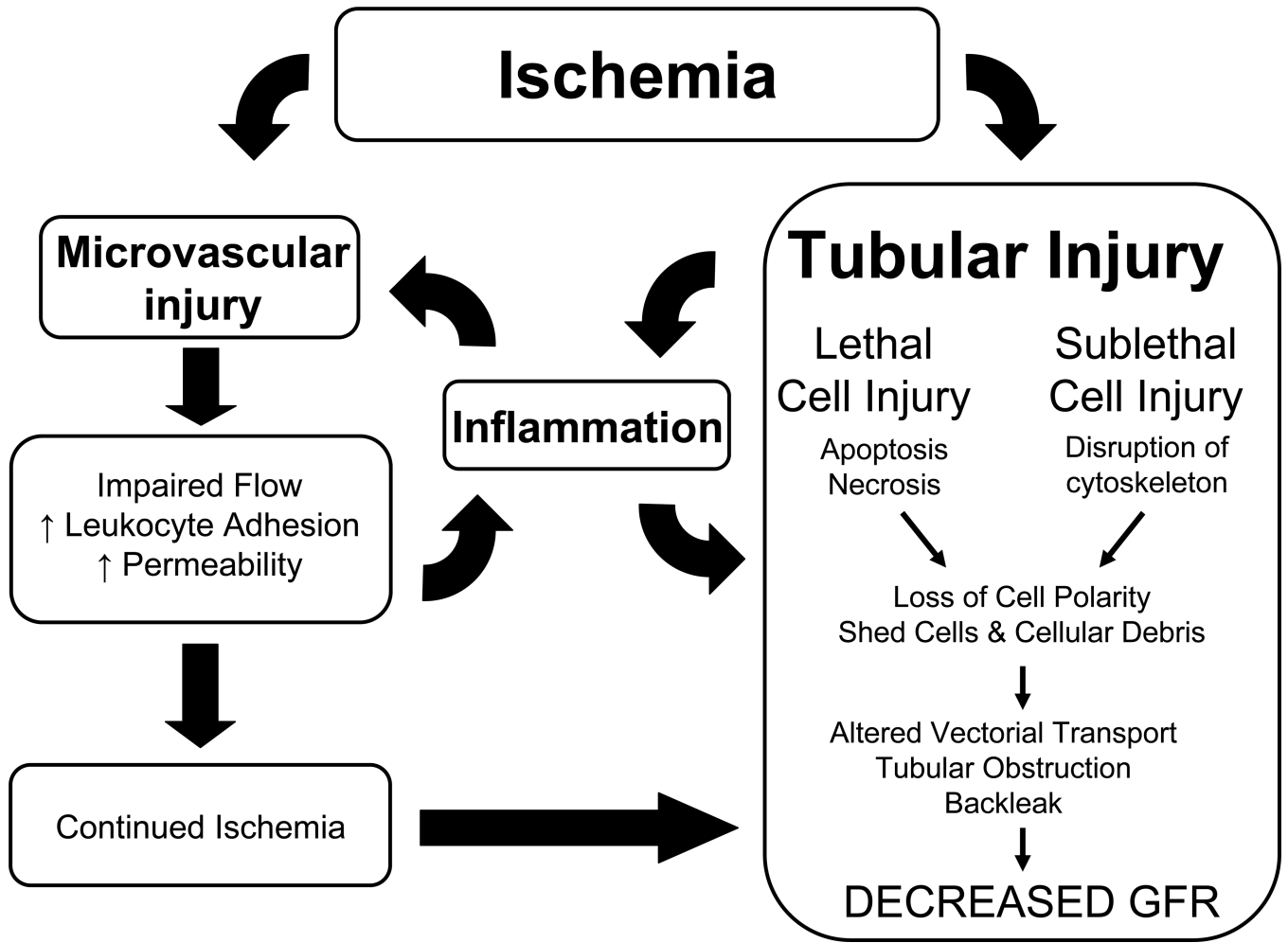


Figure 3. Interplay between tubular and vascular injury leading to sustained reductions of GFR in the extension phase of AKI

Injury induced by ischemia can result in damage to both the tubular as well as the microvascular compartment. Resolution of vasoconstriction appears effective at reducing injury when administered prophylactically, but not following established injury. Resistance may be due to exacerbated inflammation, which may impart reductions in RBF and GFR insensitive to vasodilator therapies. Of central importance in this process is the activation of inflammatory processes which are influenced by factors released by damaged proximal tubules as well as adhesion of damaged microvascular cells. Infiltrating leukocytes may impinge on RBF either by secreting vasoactive factors, or by contributing to the disruption of flow by physical interference. In addition, exacerbated hypoxia leading to tubular obstruction may contribute to reductions in GFR independent of vasodilator therapy. From citation (531)

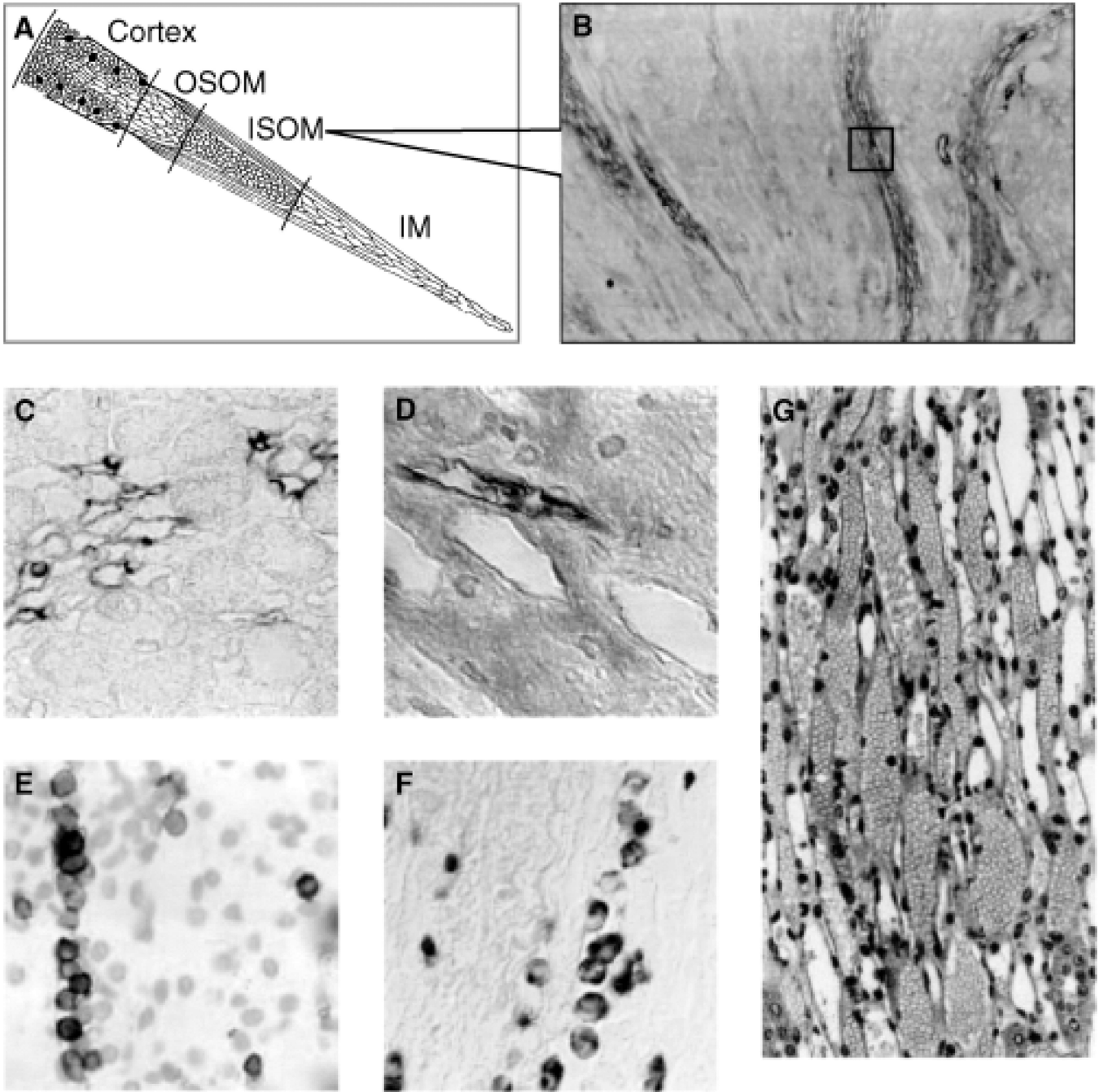


Figure 4. Organization of vascular compartment in the kidney

(A) The medulla is arterial supplied from the efferent arterioles of the juxtamedullary glomeruli, giving supply to the descending arterial vasa recta, and further to the ascending venous vasa recta, draining into the arcuate veins. Abbreviations are: OSOM, outer stripe of outer medulla; ISOM, inner stripe of outer medulla; IM, inner medulla. (B) Very strong expression of b130-1, 2 hours after ischemia/reperfusion (I/R) injury of the kidney, at the level of the ascending vasa recta. (C) Detailed expression of b130-1, 2 hours after I/R injury of the *rat* kidney, at the level of the ascending venous vasa recta. (D) Detailed expression of b130-1, 2 hours after I/R injury of the *human* kidney, at the level of the ascending venous vasa recta. (E) Trapping of CD28-expressing T cells in the ascending vasa recta (HIS-17

staining). (F) Trapping of monocytes/macrophages in the ascending vasa recta (ED-1 staining). (G) This trapping of leukocytes in the ascending vasa recta results in an upstream congestion at the ascending arterial vasa recta. This congestion, or no-reflow, represents a well known phenomenon in acute ischemic injury, exacerbating during reperfusion the ischemic damage. From citation (626).

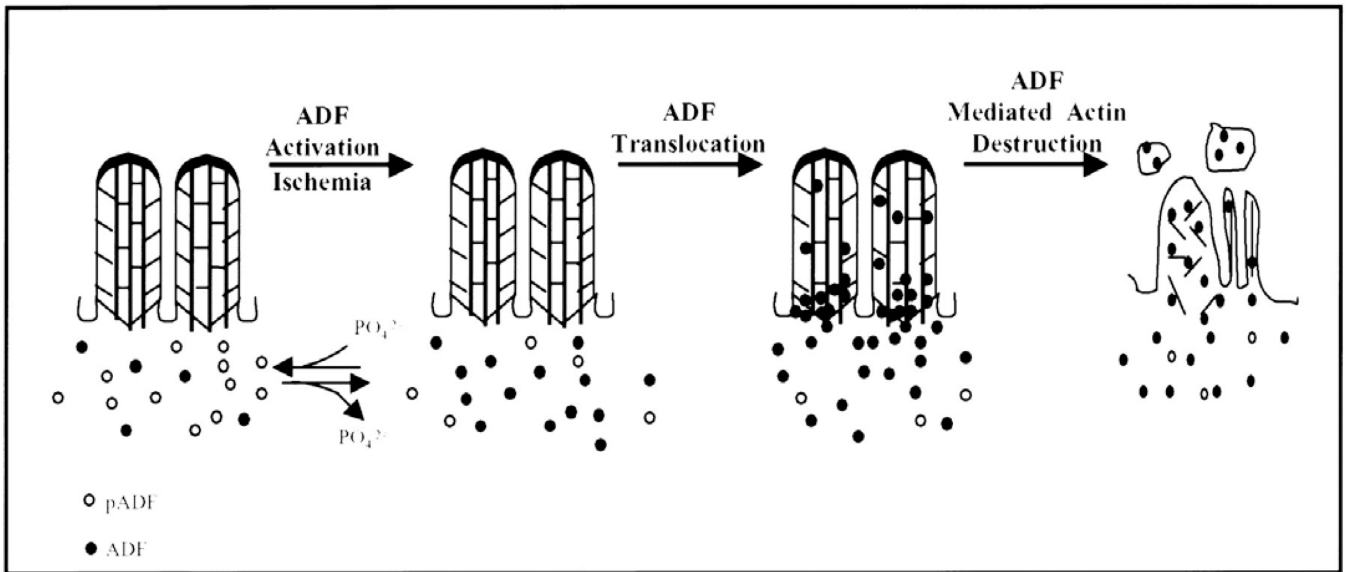


Figure 5. A proposed model for the dephosphorylation, activation, and translocation of ADF to the apical microvilli during ischemia

Under physiological conditions, the distribution of ADF and phosphorylated ADF (pADF) in proximal tubule cells is diffuse throughout the cytoplasm with little or no localization to the apical microvillar region. With ischemia, pADF is dephosphorylated and, therefore, activated. In addition, the diffuse cytoplasmic localization of ADF changes with ADF now concentrating at the apical membrane region of the cell. It is hypothesized that ADF relocates to the apical microvillar region and binds the microfilament core, resulting in markedly enhanced filament severing and depolymerization. Breakdown of the microfilament core is accompanied by dramatic changes in the overlying microvillar membrane. The microvillar membrane is internalized or extruded as membrane vesicle or blebs. These vesicles contain both ADF and monomeric actin. Figure and legend from citation (15)

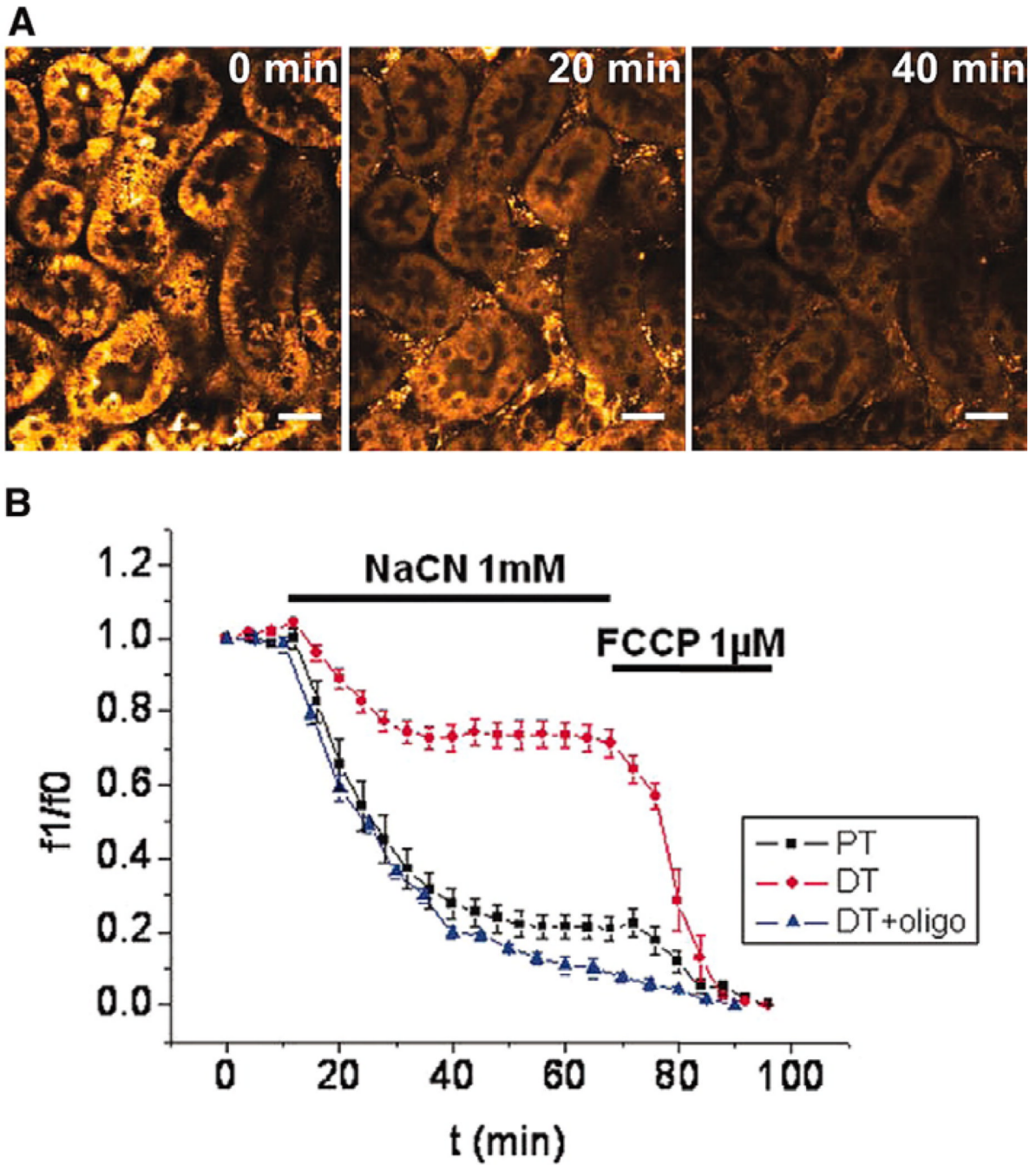


Figure 6. Evidence of mitochondrial depolarization in kidney tissue slices by chemical anoxia using multiphoton imaging
 PTs loaded with TMRM showed rapid depolarization of $\Delta\psi_m$ after chemical anoxia. Bar = 20 μm . (B) In the DT, the decrease was slower and $\Delta\psi_m$ was not completely depolarized after 60 min of anoxia; however, in the presence of oligomycin (5 $\mu\text{g}/\text{ml}$), $\Delta\psi_m$ depolarized rapidly in distal tubular cells when exposed to anoxia. Data are means \pm SE signal per tubule from a total of 15 PTs, 15 DTs without oligomycin, and 29 DTs with oligomycin from three separate slices for each experiment. The data were normalized from 1 (value at $t = 0$, taken as resting $\Delta\psi_m$) to 0 (minimum value after FCCCP, taken as 0 mV). Data from citation (185).

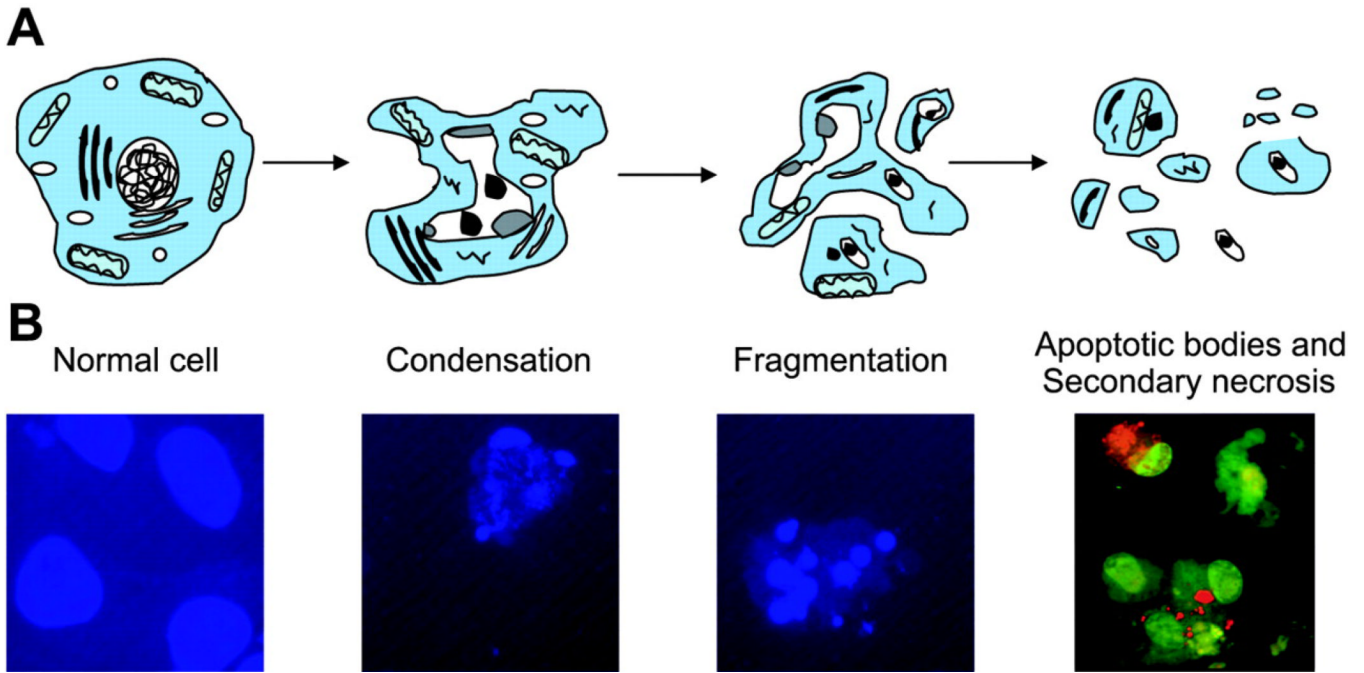


Figure 7. Illustration of the various stages of apoptotic cell death

A: depiction of the stereotypical changes including condensation, changes in nuclear structure, and fragmentation of the cell into small apoptotic bodies. In vivo, the apoptotic bodies are phagocytosed by neighboring cells, whereas in vitro they undergo swelling and eventual lysis (secondary necrosis). B: photographs of LLC-PK1 cells undergoing apoptosis at the corresponding stages as shown in A. Apoptosis was induced by overnight exposure of the cells to 50 μ M cisplatin. The cells in the first 3 photographs were stained with Hoechst dye, and the cells in the last photograph were stained with acrydine orange and ethidium bromide. In the last photograph, viable cells appear green, whereas the apoptotic cells with intact plasma membrane appear green with yellowish dots representing condensed chromatin; apoptotic cells and bodies that are undergoing secondary necrosis appear bright orange or red due to the plasma membrane damage and entry of ethidium bromide. Illustration from citation (415).

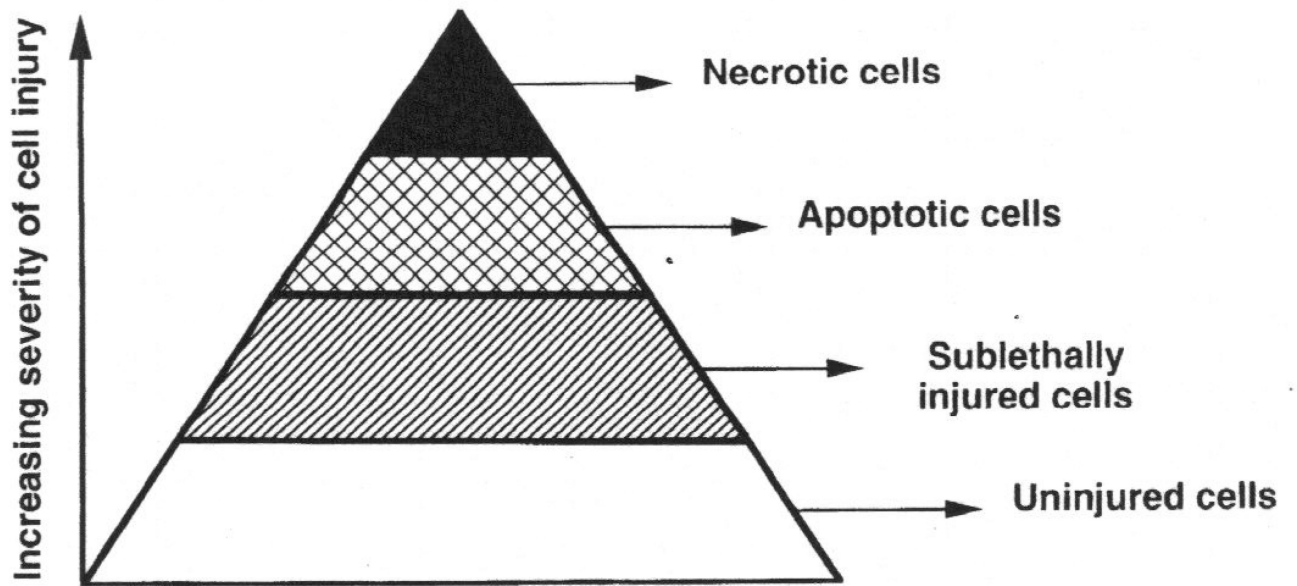


Figure 8. The continuum of renal cell damage

Individual renal tubular cells are likely to respond in different ways to injury depending upon the severity of the noxious stimulus. The majority of cells presumably remain viable, either because they escape injury altogether, or because they are only sublethally injured and able to recover. More severe injury likely results in apoptosis, whereas necrosis only occurs when cells are subjected to extremely severe injury that leads to critical energy depletion and subsequent metabolic collapse. Legend and figure from citation (320).

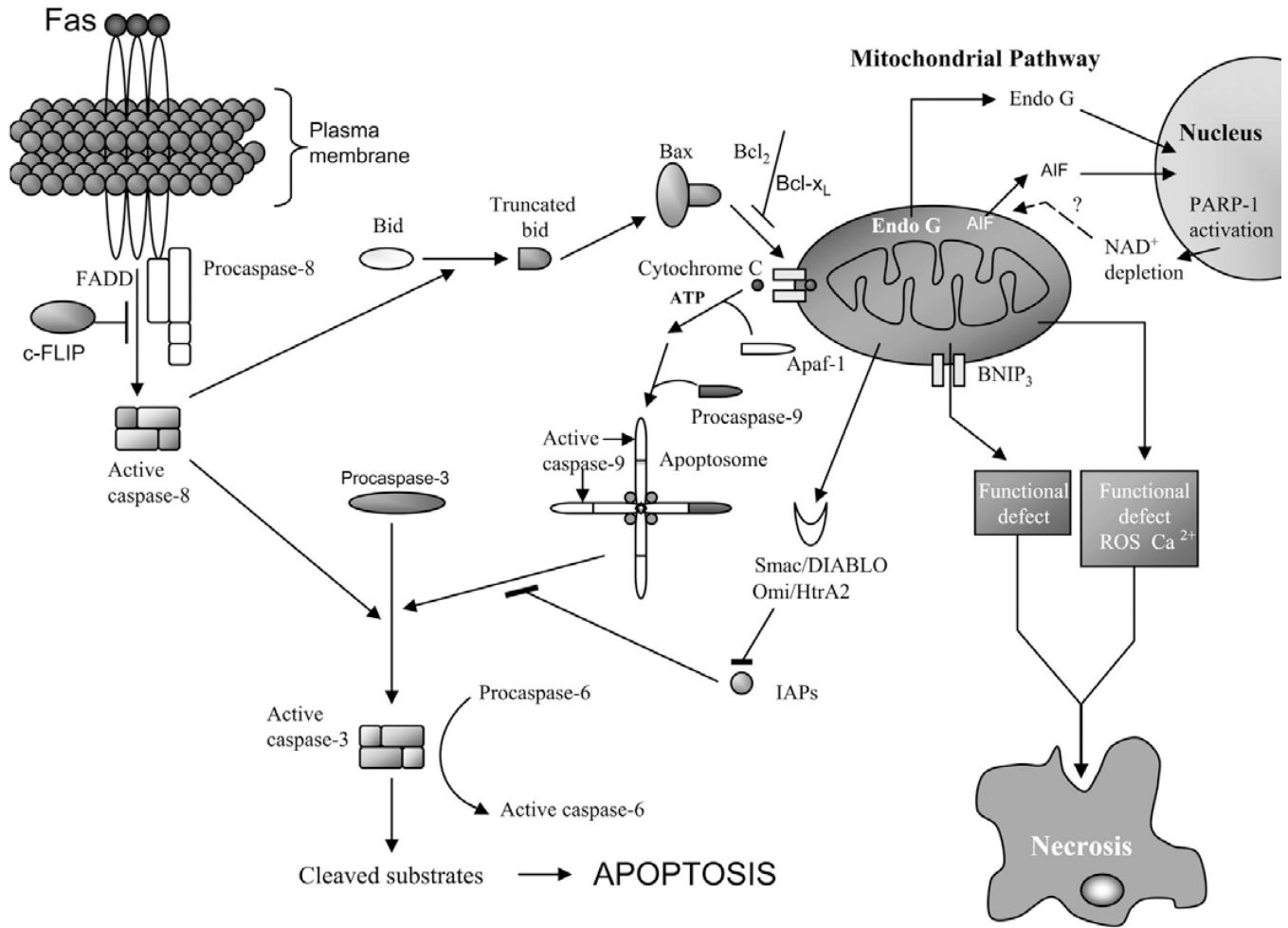


Figure 9. Overview of death-signalling pathways in mammalian cells

The death receptor pathway (left) is initiated by the binding of a ligand (Eg: FasL) to its receptor Fas, which results in the sequential recruitment of FADD and pro-caspase-8. c-FLIP can block the recruitment of pro-caspase-8 to the complex. The proximity of several pro-caspase-8 molecules results in its activation. Caspase-8 can proteolytically activate caspase-3, or it can cleave Bid to its truncated form t-Bid, which binds to Bax and gets integrated into the mitochondrial membrane to release cytochrome c. In response to various cellular stress-induced apoptotic stimuli, the intrinsic mitochondrial pathway is activated. This pathway involves the translocation of proapoptotic molecules such as Bax from the cytosol to the mitochondrial membrane. Bax can release cytochrome c from the mitochondria into the cytosol. Cytochrome c associates with Apaf-1 and caspase-9 to form the apoptosome and subsequent activation of caspase-3. Mitochondria also release AIF and Endo G, which may exert their effects on the nuclei. Mitochondria released Smac/Diablo and Omi/HtrA2 sequesters inhibitors of apoptosis (IAPs) to prevent them from inhibiting caspase-3. BNIP3 is a Bcl2 family member that is translocated and integrated into the mitochondria. Unlike other Bcl2 family members, BNIP3 can induce necrotic cell death in response to death stimuli. Activation of poly (ADP-ribose) polymerase (PARP) leads to NAD⁺ depletion and may induce mitochondrial depolarization to release AIF. ROS, reactive oxygen species. Legend and Figure from citation (320).

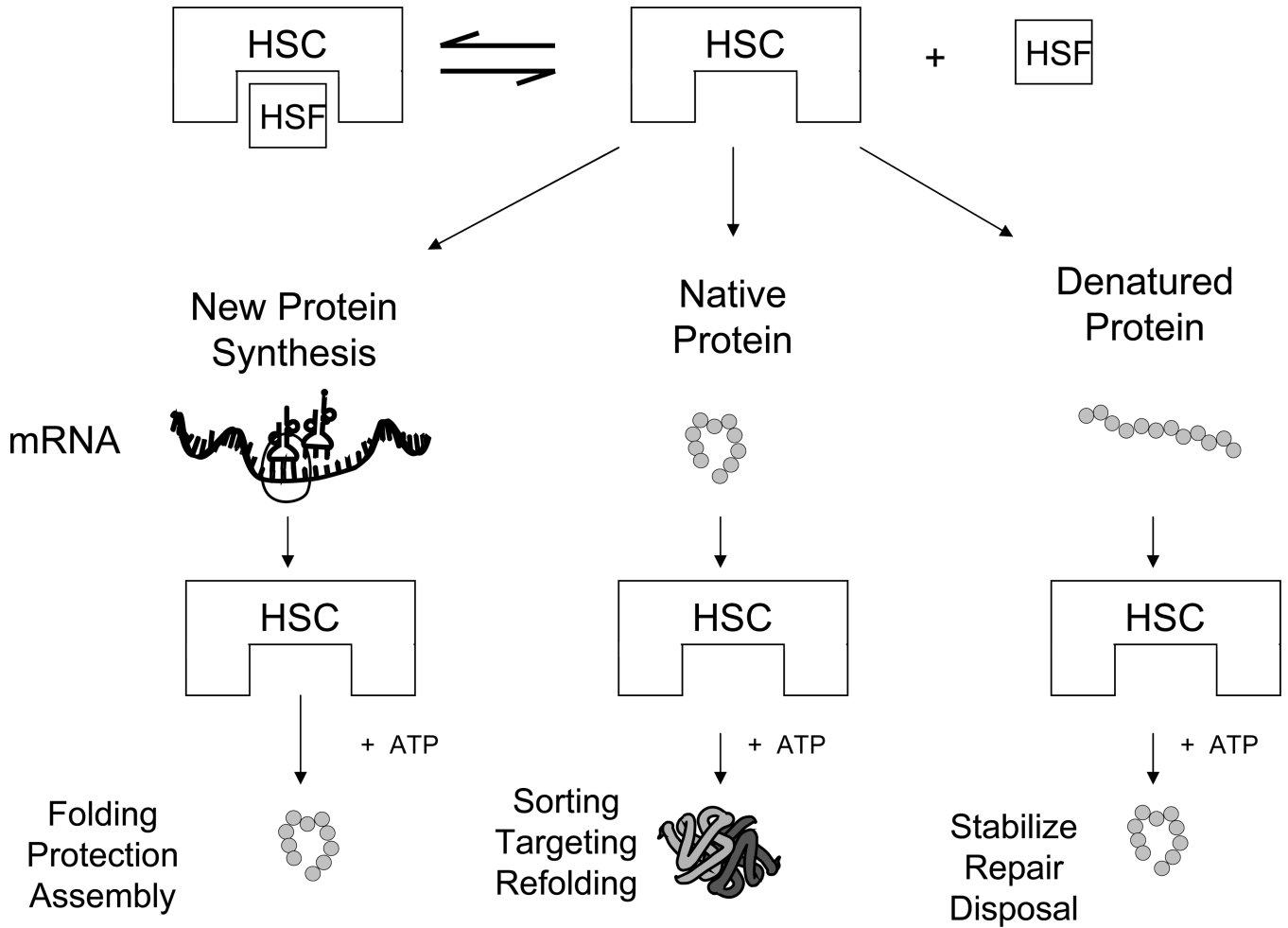


Figure 10. Control of HSP expression in response to cell stress

Shown are the known actions of the constitutively expressed Hsps, primarily of the Hsp70 family called heat shock cognates (HSC) in processing cellular functions. Cell stress increases denatured proteins increasing the demand for HSC. Heat shock transcription factor (HSF), reversibly binds to HSC and is released with the increased demand for HSC. HSF then rapidly initiates transcription for all inducible Hsps including Hsp 70 and Hsp 25/27. Figure from citation (521).

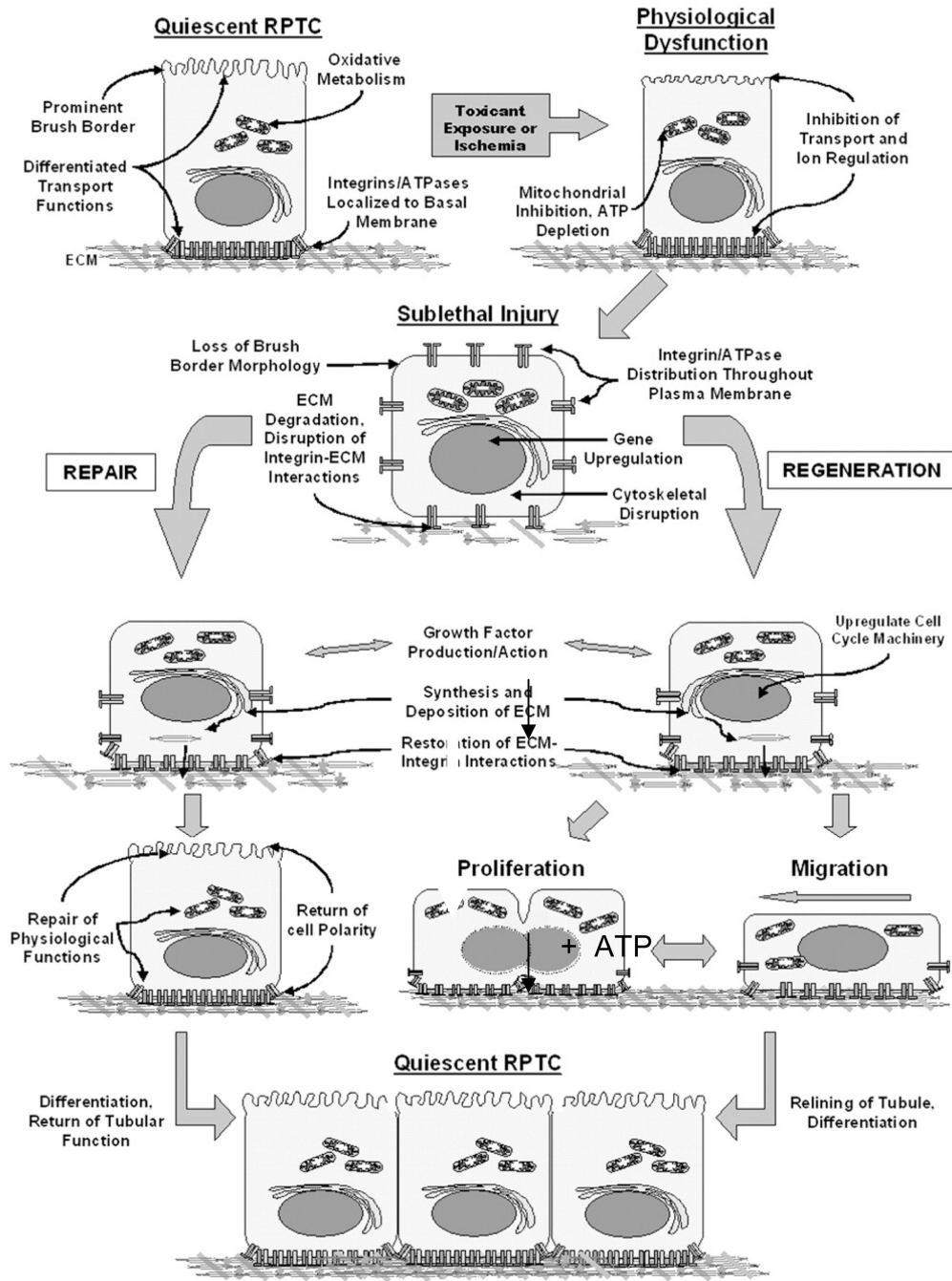


Figure 11. Repair and regeneration of renal proximal tubule cells following acute sublethal injury
 Sublethally injured RPTC either repair physiological functions and restore normal tubular function or dedifferentiate, migrate, and/or proliferate to replace lost cells, then differentiate and resume normal function. The processes of repair and regeneration work in concert to ensure relining of the damaged nephron and restoration of renal function. From citation (402).

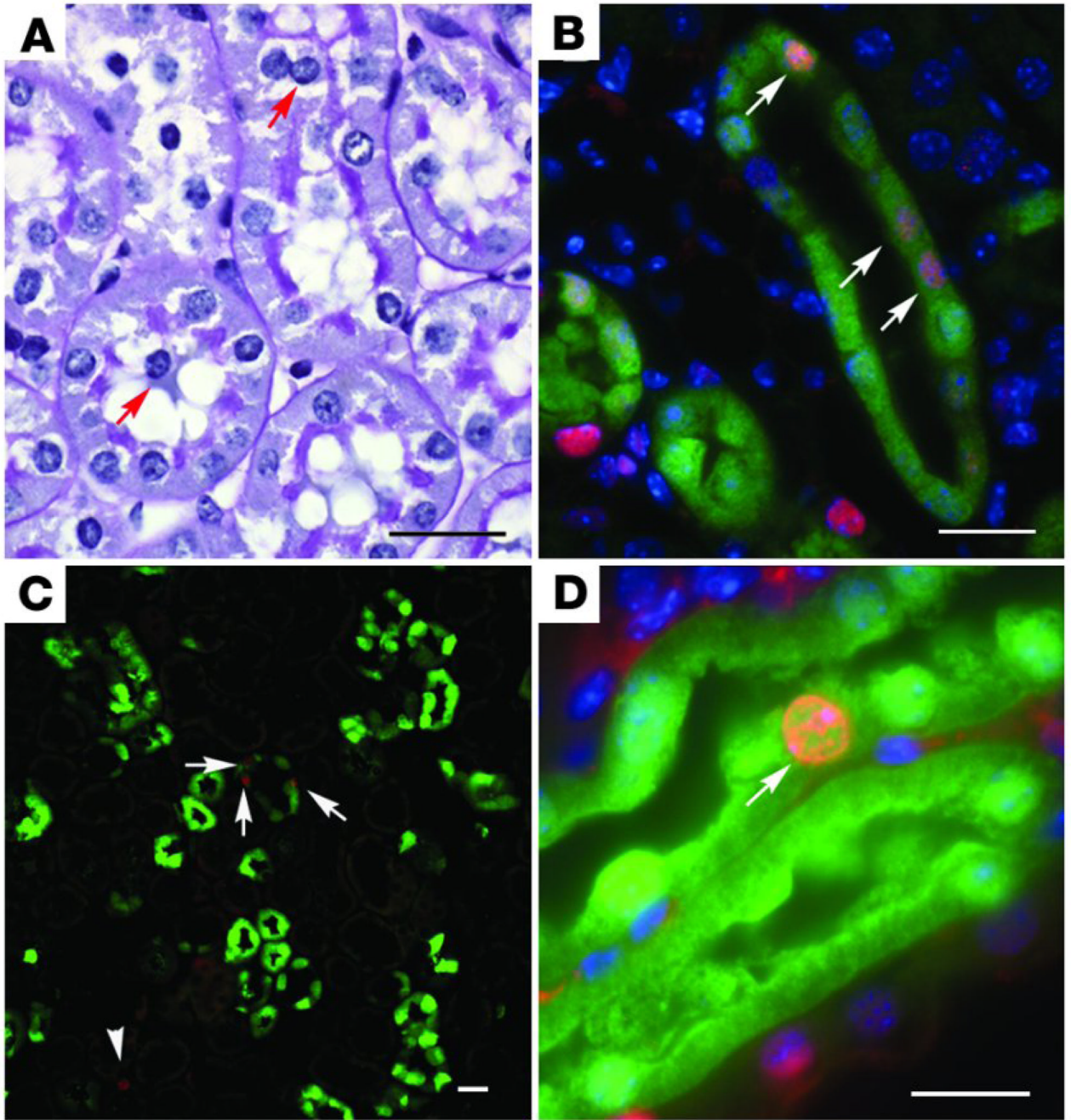


Figure 12. Evidence that sublethally damaged proximal tubules are the source of dividing cells during recovery from AKI

PAS staining and immunostaining were performed in the kidney sections of $cre^{ksp};Z/EG$ mice with renal IRI. (A) PAS staining of the kidney at 2 days after IRI. Tubular injury is shown by the loss of brush border membrane, cell detachment from the basement membrane, and nuclear condensation in some cells (arrows). (B) Expression of PCNA in tubular cells (red, arrows). (C) Low-power image of BrdU incorporation in renal tubules. Some BrdU-containing cells (red, arrows) colocalized with EGFP-expressing cells (green). The arrowhead indicates BrdU incorporation in an EGFP-negative cell. (D) BrdU incorporation (red, arrow) in the epithelial cells expressing EGFP (green). Note: the cre

transgene labels tubular cells and their progeny with EGFP. The nuclei were counterstained with DAPI, and images were merged (B–D). Scale bars: 20 μm . From citation (325).

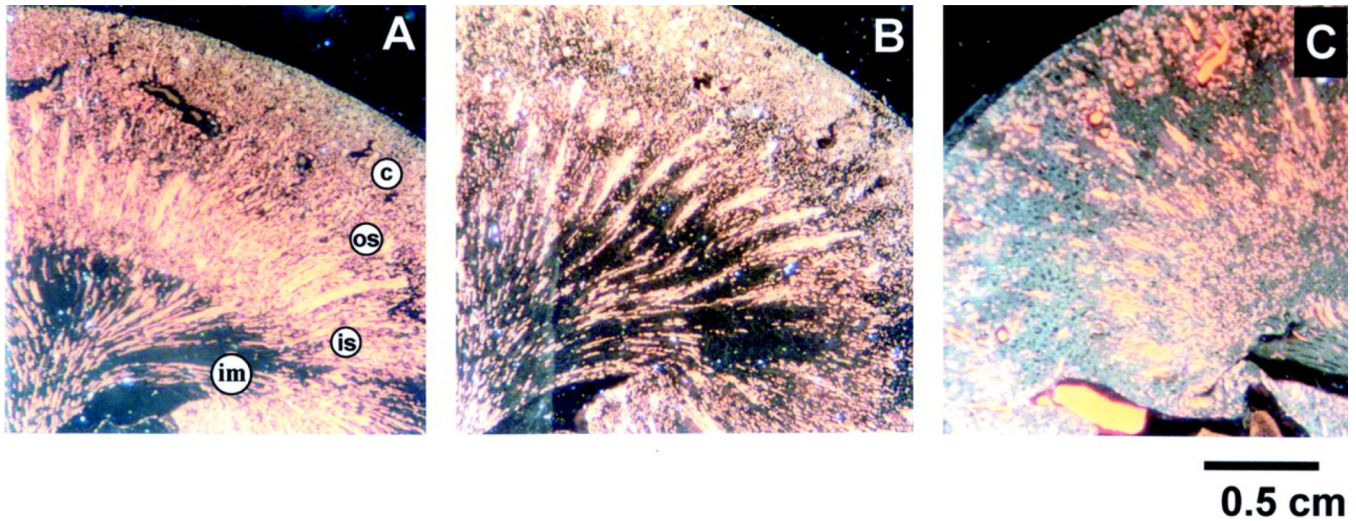


Figure 13. Gross renal morphology and capillary filling in normal and postischemic kidneys
 Representative stereoscopic views of 20- μ m Microfil-infused kidney section. Shown are Microfil-infused kidneys from a sham-operated rat (A) at 4 (B) and 8 wk (C) post-ischemic injury. In this stereoscopic view, Microfil appears as bright yellow against a dark background. A reduction in Microfil-infused structures in recovered postischemic kidneys is evident. c, cortex; os, outer stripe of the outer medulla; is, inner stripe of the outer medulla; im, inner medulla. Magnification is shown. From citation (32).

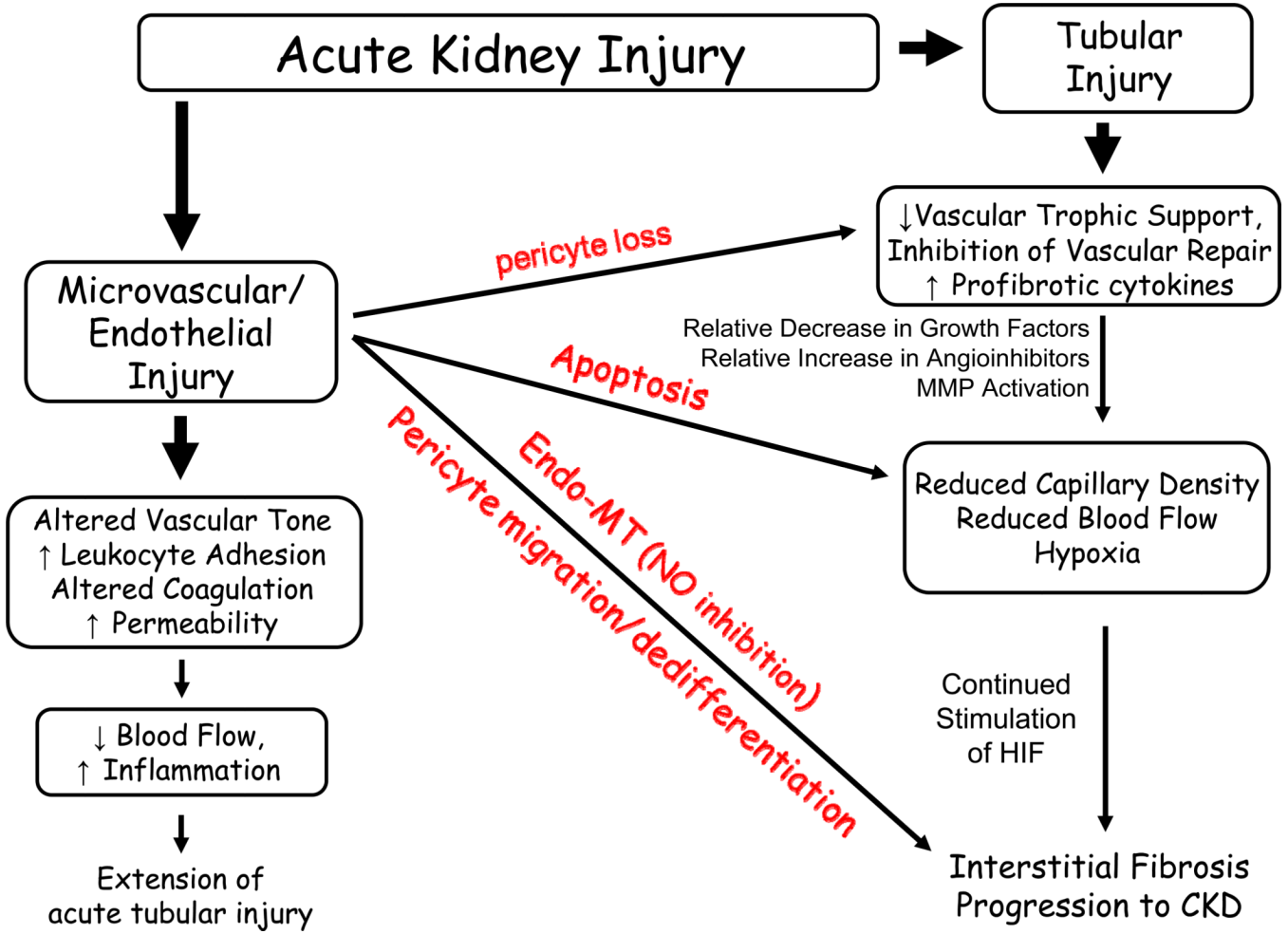


Figure 14. A potential role for vascular dropout in promoting the development of CKD following AKI

Acute injury has the potential to affect both tubular and vascular compartments. In addition to direct injury to the microvascular compartment, tubular injury may compromise normal vascular support, shifting the environment to one which promotes vascular impairment rather than vascular stability, including the loss of VEGF expression, the increase in TGF-beta expression as well as several other angio inhibitory compounds. The resultant decrease in capillary structures has a number of potential consequences on renal function including the exacerbation of hypoxia and the impairment of Na handling hemodynamic responses. Hypoxia, along with the potential endothelial mesenchymal transition are likely to participate in the development of fibrosis, which is also influenced by sustained immune/inflammatory activity. The figure modified from and earlier version published in citation (29).