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Role of renal hypoxia in the progression from acute kidney injury to chronic kidney disease

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Abstract

Over the last 20 years, there has been an increased appreciation of the long-term sequelae of acute kidney injury (AKI) and the potential development of chronic kidney disease (CKD). Several pathophysiological features have been proposed to mediate the AKI to CKD progression including maladaptive alterations in tubular, interstitial, inflammatory and vascular cells. These alterations likely interact to culminate in the progression to CKD. In this article we focus primarily on evidence of vascular rarefaction secondary to AKI, and the potential mechanisms by which rarefaction occurs in relation to other alterations in tubular and interstitial compartments. We further focus on the potential that rarefaction contributes to renal hypoxia. Consideration the role of hypoxia in the AKI to CKD transition focuses on experimental evidence of persistent renal hypoxia following AKI and experimental maneuvers to evaluate the influence of hypoxia, *per se*, in progressive disease. Finally, consideration of methods to evaluate hypoxia in patients is provided with the suggestion that non-invasive measurement of renal hypoxia may provide insight into progression in post AKI patients.

Keywords

Fibrosis; peritubular capillaries; inflammation; renal hypoxia

Background

Acute kidney injury (AKI) is characterized by an abrupt loss of glomerular filtration rate, often associated with altered renal blood flow and increased local and systemic inflammation.¹ The leading causes of AKI are ischemia reperfusion (I/R) injury,² which may occur secondary to major surgery such as cardiac surgery (particularly when requiring cardiopulmonary bypass),³ sepsis⁴ and exposure to nephrotoxic agents.⁵ It is estimated that approximately 5% to 10% of critically ill patients suffer from AKI ⁶ and every year around 600,000 new cases of AKI are reported in the United States.⁷

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While recovery of renal function is expected in the majority of surviving patients following AKI, epidemiological studies have shown that such patients are more likely to develop chronic kidney disease (CKD) compared with patients without a history of AKI.^{8,9} Using data from the U.S. Department of Veterans Affairs, Amdur et al. reported that patients developing AKI following myocardial infarction or pneumonia showed a decline in renal function and an increased risk of developing stage-4 CKD versus similar patients without AKI.⁹ In addition, Ishani et al. reported that AKI in U.S. Medicare patients was associated with increased risk of end-stage renal disease (ESRD), which was significantly enhanced in patients with a prior history of CKD.⁸ In a meta-analysis, Coca et al. reported that AKI was associated with an increased risk of CKD by 8.8-fold and ESRD by approximately 3.1-fold versus patients without prior AKI.¹⁰ These reports suggest that AKI predisposes the development of CKD. However, most of the information on pathophysiological mechanisms of the AKI-to-CKD transition derives from studies using animal models of renal injury.

The pathophysiology of AKI to CKD

Following AKI, CKD may progress because of incomplete or maladaptive tissue repair, setting in motion processes promoting the development of interstitial fibrosis.¹¹ Alterations in renal structure that may promote development of CKD following acute injury include nephron loss, incomplete tubular repair, changes in interstitial cell composition, inflammation and vascular rarefaction (Figure 1). These alterations do not occur in isolation and interact to direct progressive CKD. With regard to capillary rarefaction, we have suggested that limitations in renal perfusion may enhance renal hypoxia and fuel progression of CKD.¹² In this article, we summarize recent developments in our understanding of maladaptive repair responses that contribute to the AKI-CKD transition. We focus on the genesis of capillary rarefaction following both AKI and in progressive CKD. We consider the potential role of hypoxia, and consider how future efforts may be pursued to translate data from preclinical modes to the patient.

Adaptive and maladaptive epithelial repair following AKI

Recovery of renal function is expected following AKI in most in surviving patients, while subsets of patients may develop renal insufficiency and progress to CKD.¹ Generally, recovery from AKI is associated with the resolution of renal perfusion and repair of the damaged epithelial cells, typically the most prominent morphological feature following injury.¹³ Increasing the severity of AKI may tilt the balance toward maladaptive and inefficient repair while adaptive repair may become limited.¹¹ Not surprisingly, in models of ischemia reperfusion (I/R), increasing ischemic time results in both greater initial tubular damage as well as a greater degree of renal fibrosis one month following reperfusion.¹⁴

Efficient tubular repair results from recovery of sub-lethally damaged cells and/or proliferation of damaged de-differentiated cells to repopulate the tubular basement membrane.¹⁵ Re-differentiation and hypertrophy of recovered or new cells helps to re-establish normal tubular architecture. However, areas of tubular atrophy or evidence of persistent de-differentiation have been observed following I/R injury. In rats, the frequency of these structures is enhanced when injury occurs in the setting of reduced renal mass,

becoming more common following I/R with a 50% or 75% reduction in renal mass.¹⁶ While complete tubular atrophy reduces total nephron number, focal lesions around atrophic tubules are sharply demarcated and fibrosis does not progress from these areas, suggesting that atrophic nephrons do not directly contribute to progressive interstitial fibrosis.^{16,17}

In contrast, Venkatachalam and colleagues have proposed that small clusters of proximal tubular epithelial cells that fail to completely re-differentiate may promote interstitial fibrosis.¹⁸ These undifferentiated cells can be identified by expression of vimentin and are associated with increased expression of pro-fibrotic growth factors including: transforming growth factor- β 1 (TGF- β 1), platelet derived growth factor-B (PDGF-B) and connective tissue growth factor (CTGF), when compared with fully differentiated tubules.^{17,19,20} These secreted growth factors have been proposed to act in a paracrine fashion on interstitial cells *i.e.*, pericytes or fibroblasts, found in close proximity.¹⁸

The activation of cell cycle activity is important for effective recovery of the tubule, while persistent activation of the cell cycle, identified by activity of c-Jun kinase (JNK), has been associated with development of interstitial fibrosis.^{19,21} A potential role for cell cycle activity in fibrosis was addressed in studies using various murine models of renal injury including moderate reversible I/R injury, severe I/R injury, aristolochic acid nephropathy and unilateral ureteral obstruction (UUO).¹⁹ Interestingly, the models associated with a greater degree of fibrosis displayed a greater number of cells in G2/M cell-cycle arrest, identified by phosphorylation with histone H3. These cells produce higher levels of pro-inflammatory and pro-fibrotic cytokines, including TGF- β and CTGF when compared with cells in G0/G1 after I/R injury.¹⁹ Interestingly, blockade of JNK bypassed G2/M arrest and reduced fibrosis following bilateral renal I/R injury.^{19,22}

Taken together, persistent de-differentiation and/or G2/M cell cycle arrest are considered to be important drivers of renal fibrosis following injury. We suggest that physiological factors related to re-establishment of perfusion may influence the success or failure of tubular repair, and therefore the fibrotic nature of tubules following AKI. Specifically, peritubular capillaries, which provide nutrient and oxygen for tubules, are compromised following AKI and may lead to failed tubular recovery (Figure 1). Support for this idea comes from studies of rats subjected to renal micro-embolization injury following infusion of acrylic microspheres into the renal artery.²³ Tubules from rats at 4 weeks following injection demonstrated heterogeneous patchy areas of undifferentiated, vimentin-positive cells, which expressed PDGF-B, localized adjacent to expanded interstitium. In contrast, vimentinnegative cells, even in the same tubule, did not express PDGF-B.²³ Thus, disruption of capillary flow may impair efficient tubular recovery and promote a persistent pro-inflammatory/pro-fibrotic phenotype.

Capillary dropout after AKI

In contrast to the renal tubular system, the renal microvascular system has a poor regenerative capacity and impairments in microvascular structure have been proposed to contribute or hasten progressive CKD.^{12,24} Observations of Basile and colleagues demonstrated significant and essentially permanent reductions of peritubular capillary (PTC)

density in rats following bilateral I/R injury (Figure 2).²⁴ In their study, Microfil® silicon rubber was infused into rat kidneys between 4 to 40 weeks after renal I/R injury. Despite evidence of recovery of renal function and tubular structure, capillary density was reduced in both the cortex ($\sim 25\% - 30\%$) and the inner stripe of the outer medulla ($\sim 35\% - 40\%$).²⁴ The finding of reduced capillary density following AKI has been confirmed by other investigators using a variety of different methods (Table 1). For example, Kramann et al. utilized fluorescence micro-angiography and reported a reduction of the total perfused area in both renal cortex and medulla after severe I/R-induced AKI in mice.²⁵ The reduced perfusion was associated with a reduction in both the density of vessels as well as the diameter of individual surviving PTCs.²⁵ Several others have utilized immunohistochemical staining to detect changes in vascular density. Studies using antibodies against CD31, RECA (rat endothelial cell antigen), MECA32 (mouse endothelial cell antigen) and cablin, have all shown acute and sustained reductions in PTC density following AKI (Table 1). Labeling of endothelial cells using a transgenic approach by crossing Tie2-Cre mice with a floxed reporter line, has also been used to demonstrate an overall reduction in PTC following AKI. ²⁶ Capillary rarefaction has been demonstrated in other experimental animal models of AKI including, UUO and folic acid nephropathy.^{27,28} and has also been demonstrated in human patients after delayed graft function.^{29,30}

It should be recognized that all of these approaches have limitations and caution should be exercised in interpreting results from these studies. Measurements that rely on staining a putative endothelial marker are subject to changes in protein expression, which could be altered in response to sub-lethal injury and not reflective of actual loss of vessels. For example, CD31/platelet endothelial cell adhesion molecule, one of the most widely used endothelial cells (EC) markers, may be transiently down-regulated in response to interferon gamma or TNF-a.³¹ The loss of staining may therefore overestimate PTC loss, especially in the early post-AKI period. By extension, a subsequent increase in CD31 expression may not distinguish between recovery of sub-lethally damaged cells versus the generation of new endothelial cells due to repair. In contrast, techniques based on vascular filling (Microfil, micro computed tomography (microCT), microangiography) are fraught with technical and biological considerations. In particular, the ability to fill a vessel may be difficult if it is constricted or congested. Since both of these complications are present in the early stages of AKI,¹ reduced filling may not necessarily reflect rarefaction. Despite these experimental concerns, the abundance of reports using different experimental approaches strongly suggests that renal microvascular density is compromised following acute kidney injury.

While the majority of studies indicate persistent rarefaction following AKI, few studies have carefully evaluated the time course of this response. One notable exception provided by Ehling et al. used MECA32 staining to quantify renal microvessels following I/R in mice.³² Reduced vessel density was detectable as early as 1 day post I/R, but progressed steadily up to 14 days and remained stable thereafter. This timing is noteworthy as it suggests that renal capillary rarefaction develops slowly, simultaneous to the time normally associated with repair of renal tubules.³² Moreover, these data imply that a window of opportunity during the repair phase could be envisaged to attenuate vascular rarefaction. Several approaches, including treatment with vascular endothelial growth factor (VEGF)-121³³ or cartilage

oligomeric matrix protein/angiopoietin-1³⁴ provide protection against microvessel loss following AKI.

The mechanisms mediating PTC rarefaction are not completely understood. Surprisingly, evidence of apoptotic cell death within the endothelium is minimal. Horbelt et al., using a Tie2-GFP mouse, could not detect endothelial apoptosis by TUNEL staining following renal I/R injury, while injection of an anti-CD95 antibody abundantly labeled TUNEL+EC in kidney; thus EC apoptosis may be context dependent.⁴¹ Conversely, ECs express markers of mesenchymal cells for at least 7 days following I/R in rats, suggesting endothelial-mesenchymal transition (EndoMT).²⁶ The findings of fate-tracing studies using Tie2-Cre mice crossed with YFP reporter mice are consistent with activation of EndoMT in the setting of AKI.²⁶ Live multiphoton imaging studies 2 weeks following I/R demonstrated focal regions where peritubular vessels were devoid of blood flow.²⁶ The YFP-positive cells were thickened and lacked the normal endothelial appearance observed in sham-operated controls. ²⁶ Moreover, there was little evidence of endothelial proliferation following injury suggesting that EndoMT, combined with a lack of adequate endothelial repair mechanisms, contributes to PTC loss following AKI.

Renal microvascular rarefaction may be due in part to alterations in the balance of positive and negative trophic factors in the AKI microenvironment. Transient elevations of antiangiogenic factors have been reported following AKI.^{1,42} VEGF-A, an angiogenic factor, is expressed in renal tubular epithelial cells and might bind to its receptor located on endothelial cells of adjacent PTCs.⁴³ This tubulo-vascular crosstalk is crucial for maintenance of PTC density in kidney. In mice expressing the diphtheria toxin receptor in proximal tubules, administration of the toxin induces both proximal tubule injury and a reduction in capillary density.⁴⁰ Consistent with this, Dimke et al. demonstrated that genetic ablation of VEGF-A in mouse renal tubules also decreased PTC density.⁴³

Several studies have demonstrated that renal VEGF expression is compromised early after AKI in rats ^{33,44} and is associated with reduction of PTC density.³³ VEGF expression is reduced in other chronic injury models associated with fibrosis.^{45,46} The loss of VEGF expression is counter-intuitive since other HIF-related genes are prominently activated in response to I/R injury.⁴⁴ However, it was suggested that reduced VEGF expression reflects the undifferentiated phenotype of injured tubules.¹⁶ Regardless, the transient reduction in VEGF may participate in vascular rarefaction since treatment of rats with VEGF-121 during the early post-ischemic period ameliorates capillary rarefaction and subsequent CKD.³³

Vascular stability is also provided by the tight interplay between pericytes and endothelial cells.⁴⁷ Several recent studies have generated evidence that the distance of pericytes from ECs increases within hours of renal I/R injury. This detachment or disruption of the normal pericyte-EC structural relationship results in migration of pericytes from the perivascular to the interstitial space and activation of these cells to extracellular matrix producing myofibroblasts.⁴⁸⁻⁵¹ Evidence suggests that these activated myofibroblasts represent the primary cell type responsible for interstitial fibrosis in response to injury.^{48,49,52} Pericyte activation is also thought to destabilize the PTC network. When compared with pericytes from non-injured kidneys, injury-activated kidney pericytes/myofibroblasts express anti-

angiogenic factors such as ADAMTS-1 and these cells disrupt endothelial 3-D networks *in vitro*.⁵³ Evidence for a critical role for proper perictye-EC interaction in PTC structure was provided by a study in which ablation of Gli1+ pericytes promoted endothelial cell damage, a reduction in PTC number, focal hypoxia and subsequent evidence of damage to the adjacent tubular epithelium.⁵⁴

Capillary loss, hypoxia and progressive CKD

Renal distribution and consumption of oxygen is heterogeneous in nature.⁵⁵ The unique microvasculature network of the kidney creates a gradient of oxygen tension between cortex and medulla such that the partial pressure of oxygen in the cortex varies between 30-50 mmHg and approximately 10-20 mmHg in the renal medulla. Therefore, the renal medulla is generally considered to be on the brink of hypoxic injury.⁵⁶ In the kidney, peritubular capillaries run parallel to renal tubules and provide nutrients and oxygen to the adjacent tubules in a manner that fulfills metabolic demand.⁵⁷

Renal oxygen tension is determined by the balance between renal oxygen delivery and oxygen consumption.⁵⁸ Pathological processes that decrease renal blood flow and oxygen delivery, or processes that increase renal oxygen consumption (e.g., by increasing glomerular filtration rate), could impair renal oxygenation, and promote renal damage.⁵⁹ Clinical agents such as polyene antibiotics, nonsteroidal anti-inflammatory drugs, and radiographic contrast agents that reduce renal blood flow or increased tubular workload may exacerbate renal hypoxia and worsen the kidney injury.⁵⁶ Amelioration of renal hypoxia by increasing medullary blood flow or reducing metabolic demand is thought to provide protection to kidneys from hypoxic damage.^{12,60}

It has been two decades since Fine and colleagues proposed renal tissue hypoxia as a mediator of progressive kidney disease.⁶¹⁻⁶³ Reductions in PTC density could impair postglomerular perfusion and contribute to renal hypoxia.⁶⁴ Reductions in PTC density have been reported in virtually every model of renal disease associated with interstitial fibrosis, including, but not limited to, hypertension, aging, diabetes and reduced renal mass.¹² Capillary rarefaction was identified in biopsies from human patients with CKD of different etiologies and the degree of rarefaction strongly correlated with interstitial fibrosis and was predictive of organ failure.⁶⁵ Decreased renal oxygen tension has been proposed as a factor predisposing the development of tubulointerstitial fibrosis by activating various profibrotic signaling pathways (described further below). A vicious cycle of progressive tubulointerstitial fibrosis as the oxygen diffusion distance from capillaries to parenchyma expands.⁶¹⁻⁶³

Effect of hypoxia on fibrotic pathways

An essential feature of the chronic hypoxia hypothesis is the putative role that low oxygen levels play in activating pro-fibrogenic pathways. The activation of the hypoxia inducible factor (HIF-1 α and HIF-2 α) pathways appears central to the activities of tubular, interstitial and inflammatory cells. HIFs are oxygen sensitive transcription factors, which represent key

mediators of the cellular adaptive response to hypoxia, and much has been written on the effects of HIF on resistance and recovery from renal injury.⁶⁶–⁶⁸ In this pathway, molecular oxygen levels are sensed by prolyl-hydroxylase-2 (PHD-2), which facilitates modification of transcription factors HIF-1 α and HIF-2 α by von Hippel-Lindau tumor suppressor protein.⁶⁸ With adequate oxygen levels, PHD-2 targets the α -subunits of HIF toward ubiquination and

proteasomal degradation. However, under reduced oxygen conditions, HIF α -subunits are not degraded and so are able to form a dimer with the β -subunit and thus regulate specific gene responses.⁶⁸

Activated HIF binds to upstream promoters of pro-fibrogenic genes and induces the expression of matrix modifying factors including matrix metallopeptidase 2, tissue inhibitor of metalloproteinase 1, collagen I and CTGF leading to accumulation of extracellular matrix in the tubulointerstitum.^{68,69} Modulation of tubular epithelial cells to mesenchymal phenotype (EMT) may be mediated via HIF-1a dependent signaling pathways.^{70,71} In line with this notion, genetic ablation of HIF-1a in proximal tubular epithelial cells was shown to ameliorate the development of tubulointertitial fibrosis and inflammation in mice with UUO.⁷¹ In addition, there was a reduction in the number of S100A4-positive fibroblasts, possibly derived from epithelial to mesenchymal transition.^{71,72} In the endothelium, sustained HIF-1a expression, stimulated via targeted EC specific knockdown of PHD-2, resulted in EndoMT and development of renal fibrosis.⁷³ Thus, sustained HIF signaling under hypoxic conditions might play an important role in the development of renal fibrosis.

Additional signaling pathways including TGF- β , nuclear factor- κB (NF- κB), PI3K/Akt, angiotensin II /reactive oxygen species and the Notch pathway can be activated in response to renal hypoxia, which subsequently may stimulate myofibroblasts to produce extracellular proteins in the renal intertitium.⁷⁴ Inhibition of several of these pathways (angiotensin II, TGF- β) have been shown to contribute to the AKI-to-CKD transition.^{75,76}

Effects of hypoxia on inflammatory responses

It is now appreciated that renal hypoxia is associated with inflammation in models of progressive kidney disease. Yamaguchi et al. demonstrated that renal tissue hypoxia and inflammation co-exist and are reciprocally modulated in both acute and chronic kidney diseases.⁷⁷ Inflammation may be modulated by overexpression of the CCAAT/enhancerbinding protein δ (CEBPD), an inflammatory response gene expressed in proximal tubular cells in both acute and chronic hypoxic conditions. Data from both *in vitro* studies and *in vivo* models suggest that the effects of hypoxia on CEBPD expression are mediated through NF- κ B-dependent pathways.⁷⁷ CEBPD binds to the promotor of HIF-1 α and subsequently increases HIF-1 α expression in tubular epithelial cells, which in turn stimulates infiltration of inflammatory cells and production of inflammatory cytokines.⁷⁷

Infiltration of inflammatory cells in the renal interstitium contributes to development of renal fibrosis and plays a vital role in transition of AKI to CKD.^{1,78,79} Renal inflammation, characterized by infiltration of macrophages, neutrophils and lymphocytes is evident in experimental models of kidney disease⁸⁰ and in human renal biopsies.^{81,82} Moreover, the

number of interstitial macrophages positively correlates with interstitial fibrosis and negatively correlates with vascular density.⁶⁵

The role for inflammation in the AKI to CKD transition has received significant attention.⁸³ Inflammatory cells are thought to initiate phagocytosis and release inflammatory and profibrotic cytokines such as TGF-β. TGF-β, mainly secreted by macrophages,⁸⁴ but also by tubules ⁸⁵ promotes renal scar formation by activating myofibroblast proliferation and reducing the production of extracellular matrix-degrading proteinases.^{86,87} Thalidomide, an immunomodulatory drug, attenuated infiltration of macrophages and reduced the development of interstitial fibrosis in the kidney.⁸⁰ Specific ablation of the TGF-βRII from macrophages protected against the development of interstitial fibrosis following severe renal I/R injury.⁸⁸ Interleukin-1 receptor-associated kinase -M is, a kinase enzyme, associated with switching macrophages from an M1 inflammatory to an M2 pro-repair phenotype. Deletion of this kinase resulted in sustained M1 activation and increased severity of fibrosis following I/R injury.⁸⁹

In addition, increased IL17 expression from either T-cells or NK (natural killer) cells has been shown to contribute renal fibrosis in an AKI-to-CKD model.⁹⁰

An assessment of hypoxia in the AKI to CKD transition

Evaluation of the chronic hypoxia hypothesis in the AKI-to-CKD transition necessitates addressing fundamental features directly related to the hypothesis such as: a) Is renal hypoxia present following AKI? and b) Does alteration of renal oxygen tension alter progression of renal disease? In our opinion, available evidence in animal models is supportive but not definitive regarding the role of chronic hypoxia in the setting of AKI-to-CKD. Moreover, because of the lack of studies investigating renal hypoxia in patients, few clinical studies have helped to provide significant insight into potential roles of hypoxia in the AKI-to-CKD transition. Below we summarize some of the experimental evidence addressing these questions.

a) Is renal hypoxia present following AKI?

To date, multiple techniques have been used to detect renal tissue hypoxia in the pathogenesis of kidney disease. These techniques include pimonidazole adduct immunohistochemistry (PIM), assessment of expression of HIFs, Clark-type micro electrodes and blood oximetry, oxygen telemeter, and blood oxygen level dependent-magnetic resonance imaging (BOLD-MRI).⁹¹ Excellent reviews are available summarizing these methods in depth and illustrate advantages and limitations.⁹¹ Below, we categorize and summarize both qualitative and quantitative approaches that have been used and highlight where these have been applied in AKI and transition to CKD (Table 2).

Qualitative methods

PIM:

Pimonidazole is oxidized by molecular oxygen and reduced to a hydroxylamine intermediate in cells with oxygen tension lower than $\sim 10 \text{ mmHg}$.^{92,93} These hydroxylamine intermediates

bind to thiol containing proteins forming adducts. For *in vivo* assessment of hypoxia, pimonidazole is administered to animals prior to tissue harvest and the protein adducts are detected using immunohistochemistry, which provides excellent spatial resolution of hypoxic regions.⁹⁴ Positive staining for PIM adduct formation has been observed in the medulla and cortico-medullary region in kidneys of healthy rats, consistent with the notion that the mammalian renal medulla is hypoxic.⁹⁵ In a rat model of adenine-induced CKD, pimonidazole-adducts were confined to the tubular epithelium in a close spatial relationship with the interstitial fibrosis in both the cortex and medulla.^{95,96} This technique has also been used to detect cellular hypoxia in experimental models of diabetic nephropathy⁹⁷ and a rat model of glomerulonephritis.⁶⁴

Enhanced PIM-staining was found in post ischemic rat kidney 5 weeks following I/R despite the recovery of serum creatinine and a return of total renal blood flow.⁶⁰ Unilateral nephrectomy (UNX), a maneuver designed to increase tubular workload and thereby increased oxygen consumption, in combination with 1-kidney I/R injury prominently enhanced PIM-staining when compared to kidneys following bilateral I/R. In contrast, PIM staining was reduced by a maneuver which increased renal blood flow after induction of AKI, *i.e.*, oral supplementation with L-arginine.⁶⁰ The enhancement of hypoxia by UNX hastened the development of CKD as illustrated by fibrosis and proteinuria, while reduction of hypoxia by L-arginine sharply attenuated CKD progression.⁶⁰

The major limitation of this method is that it provides poor temporal resolution of hypoxia as an assessment can only be performed in processed tissue *i.e.*, paraffin-embedded sections or in frozen sections at a discrete time point.⁹¹ Caution should also be taken to ensure minimal false positive staining, particularly in cellular debris and casts.⁹⁶ In addition, hypoxia may be introduced during tissue harvest leading to elevated signal ⁹⁴ (and D. P. Basile, unpublished observations).

Hypoxia inducible factors (HIFs):

As described earlier, cellular adaptation to hypoxia is mediated by cellular accumulation of HIFs and therefore, increased HIF or HIF-target gene expression could be interpreted as evidence of local tissue hypoxia. Both *in vivo* ⁹⁸ and *in vitro* ⁹⁹¹⁰⁰ studies have demonstrated increased cellular levels of HIFs under hypoxic conditions. Increased expression of HIF-1a and HIF-2a were reported in various renal cell populations when rats were exposed to systemic hypoxia induced by either bleeding anemia, functional anemia (0.1% carbon monoxide), renal ischemia, or cobalt chloride (which is known to mimic hypoxia). HIF expression co-localized with HIF target-genes such as *HO-1, EPO and GLUT-1.*⁹⁸ Recently, Li et al. reported sustained hypoxia up to 28 days following severe I/R injury in mice using PIM and a corresponding expression of HIF-1a in renal tubules. The chronic induction of FoxO3, an anti-fibrotic factor, following I/R was dependent on the activity of HIF-1a fibrosis following AKI.¹⁰¹

One advantage of this approach is that it provides potential insight into activation of hypoxic signaling in renal biopsies. For example, Rosenberger et al. detected tubular HIF1 α 2 weeks following renal transplant, with higher levels being associated with longer cold-ischemic times.¹⁰² At 3 months post-transplant, HIF-1 α was not detectable in normal functioning

grafts, but was evident in grafts with evidence of clinical or subclinical rejection.¹⁰² In another study, HIF-1a expression was found in infiltrating cells and the number of HIF-1a-positive cells was associated with poor graft survival.¹⁰³

Quantitative methods

These aforementioned approaches provide no quantitative information regarding actual pO_2 levels in the kidney, making evaluation of the hypoxia hypothesis challenging. The Clarktype microelectrode is an oxygen sensor that measures soluble oxygen concentrations in a liquid medium or in tissues based on an oxidation-reduction reaction.⁹¹ The Clark-type microelectrode has been used to determine renal tissue pO_2 at various depths below the kidney surface, so can provide some degree of spatial resolution of renal tissue pO2.104,105 In 1997, Liss and colleagues introduced a modified Clark electrode with an outer tip diameter of $5.5\pm1.9 \,\mu\text{m}$ and measured renal tissue pO₂ in the cortex was $45\pm2 \,\text{mmHg}$ (meant standard error of mean), outer medulla $(32 \pm 2 \text{ mmHg})$ and inner medulla $(25\pm 2 \text{ mmHg})$ mmHg) of rats.¹⁰⁶ The approach has been used to demonstrate reductions in medullary oxygen tension in response to radiographic contrast agents or indomethacin in rats.^{106,107} Renal medullary hypoxia has also been detected in kidneys of diabetic rats, which was improved by administration of anti-oxidants such as a-tocopherol.¹⁰⁸ In addition, this approach has been used to evaluate reductions in pig renal medulla in response to graded levels of renal artery occlusion, a model associated with progressive fibrosis.¹⁰⁹ However, in a model of I/R induced AKI in rats, renal pO₂ levels measured with the Clark-electrodes were similar 5 days following reperfusion when compared to sham-operated control rats.¹¹⁰ A similar observation was also found in rats instrumented with radio-telemeter in renal medulla or post mortem assessment of cellular hypoxia by PIM.¹¹⁰ In that study, the marked deficit in renal oxygen consumption (49%) was observed in rats 5 days after reperfusion than after sham-ischemia which likely provided some protection against development of renal hypoxia in a rat model of I/R injury during subacute phase.

Recently, Koeners et al. developed a radio-telemeter based oxygen sensor for continuous monitoring of renal tissue pO_2 in freely moving rats.¹¹¹ This method has been shown to generate stable recordings of medullary or cortical tissue pO_2 for up to 3 weeks.^{111,112} In a model of angiotensin II induced hypertension and CKD, Emans et al. used this approach to detect cortical tissue hypoxia, within 15 hours of renin-angiotensin-aldosterone system activation.¹¹² These investigators also used this approach to detect renal medullary hypoxia in rats treated for 2 weeks with N- ω -nitro-L-arginine (a nitric oxide synthase inhibitor).¹¹³

Techniques with potential application in clinical practice

While oxygen sensing micro-electrodes methods provide quantitative information regarding renal tissue pO_2 , their application in human subjects is limited due to their invasive nature. The following represent less invasive approaches which are required to evaluate pO_2 in patients.

BOLD-MRI:

BOLD-MRI utilizes the paramagnetic properties of deoxyhemoglobin to generate images that detect changes in renal blood oxygenation and does not require administration of contrast agents for image acquisition.¹¹⁴ BOLD-MRI measures R2* values, defined as the time required for the MRI signal to decay in the transverse plane. This value is proportional to the blood content of deoxyhemoglobin. R2* values assigned in 3D voxels provides spatial information on regional oxygenation.¹¹⁴

Experimental evidence demonstrates a linear relationship between blood R2* values measured by BOLD-MRI and tissue pO_2 (ranging from 0.2 to 10 kPa).¹¹⁵ Studies by Prasad et al. using healthy young volunteer subjects, have shown a decrease in R2*, reflective of an increase in oxygenation in the renal medulla following furosemide administration.¹¹⁶ This observation is consistent with those of previous studies in which medullary tissue pO_2 , measured with Clark-electrodes was increased in anesthetized rats when furosemide was administered to inhibit transport in the thick ascending limb.¹¹⁷

Until recently BOLD-MRI has been extensively used to detect changes in renal blood oxygenation in both experimental animals and human subjects in response to diuretics, diabetic nephropathy, radiocontrast nephropathy and renal artery stenosis.¹¹⁸ However, recently, Prujim et al. conducted a three-year follow up study of patients with CKD, hypertensive patients without CKD, and normotensive controls. These investigators reported that low cortical oxygenation predicted a decline in yearly estimated glomerular filtration rate in patients with CKD.¹¹⁹ Although promising, a potential concern regarding BOLD-MRI is that it provides information on blood pO₂, rather than tissue pO₂ directly.⁹¹ In addition, multiple external and internal factors including hydration status,¹²⁰ low pH,¹²¹ body temperature,¹²¹ dietary sodium intake,¹²² vascular volume and vessel geometry¹²³ may influence BOLD-MRI signal thus R2* values.

Recently, Kodama and colleagues have employed a new method that utilizes a combination of electron paramagnetic resonance imaging and dynamic nuclear polarization-MRI technique in living animals in order to quantify renal oxygen tension in mmHg.¹²⁴ This double-resonance technique allows co-registration of tissue oxygen tension and renal structure non-invasively by using paramagnetic contrast agents such as oxygen-sensitive triarylmethyl radical 0X63.¹²⁴ 0X63, which is administered intravenously, interacts with dissolved paramagnetic oxygen and broadens the spectral line width of 0X63 and thereby provides quantitative information related to tissue oxygen tension. Kodama et al. have shown that renal oxygen tensions estimated by using this dynamic nuclear polarization-MRI technique were similar to those estimated with the "gold standard" microelectrode method in experimental mice models of type 1 and type 2 diabetes.¹²⁴

Urinary PO₂

Measurement of urinary pO_2 as a "physiological biomarker" for risk of AKI has received increasing attention for its potential usefulness in clinical settings.¹²⁵ Evidence demonstrates that urinary pO_2 reflects medullary tissue pO_2 .¹²⁶⁻¹²⁸ In 1965, Leonhardt et al. performed a preliminary study for measurement of pO_2 in human kidney by using oxygen micro-

electrode and showed excellent agreement between pelvic urinary pO_2 and medullary pO_2 in human patients under anesthetized conditions.¹²⁸ Therefore urinary pO_2 could represent a potential biomarker for risk of AKI.

Recently, Lankadeva et al. for the first time utilized fiber-optic probes for measurement of urinary pO_2 in an ovine model of acute kidney injury.¹²⁶ In that study, changes in medullary pO_2 closely paralleled changes in urinary pO_2 during development of AKI in conscious sheep. The fiber-optic probe, which works on the principle of fluorescence lifetime oximetry,¹²⁹ was inserted into a standard bladder catheter without blocking urine flow. Zhu et al. studied patients at high risk of AKI associated with cardiac surgery requiring cardiopulmonary bypass.¹²⁵ These investigators demonstrated that urinary pO_2 decreased progressively during cardiac surgery requiring cardiopulmonary bypass.¹²⁵ Strikingly, urinary pO_2 was reduced in patients who developed AKI relative to patients who did not develop AKI (18.7±9.0 mmHg vs 26.7±17.2, P=0.03) during post bypass epoch, and urinary hypoxia was an independent predictor for the development of postoperative AKI.¹²⁵

However, at the current time, no studies have sought to evaluate the degree to which urinary pO_2 may be persistent following an episode of AKI or whether this may correlate with the future development of CKD.

b) Does alteration of renal oxygen tension alter progression of renal disease?

Over the last two decades, an increasing amount of experimental evidence suggests the presence of renal tissue hypoxia in multiple forms of CKD, including diabetic nephropathy, ¹³²⁻¹³⁴ the remnant kidney model,¹³⁵ hypertension,¹³⁶ polycystic kidney disease, ^{137,138} CKD after recovery from I/R injury.¹³⁹ In many cases, such as in I/R injury⁶⁰ or in STZ induced diabetes,¹³³ evidence of hypoxia precedes evidence of progressive disease suggesting hypoxia may be a cause rather than an effect of the subsequent development of interstitial fibrosis.

However, whether hypoxia drives progression is not firmly established. As described, progression of CKD relies on multiple interacting pathways in the setting of renal injury (Figure 1). Much of the evidence regarding the role of hypoxia is largely indirect. It should be recognized that experimental interventions with the potential to influence renal hypoxia might also affect other pathophysiological pathways making it difficult to firmly establish a role for hypoxia per se. For example, studies from our group have shown that preservation of peritubular capillaries by VEGF-121 attenuates CKD progression following AKI.³³ While these results may be attributable to improved oxygen delivery to tissue, VEGF-121 prevents initial endothelial damage ³³ and therefore this treatment may also attenuate the activation of pericytes to myofibroblasts to reduce the population of this fibrotic cell type. Recent studies also have shown that when modified VEGF-121 is administered to pigs with renal artery stenosis, there is a shift from pro-inflammatory M1 macrophages to pro-repair M2 macrophages with sustained VEGF expression.¹⁴⁰

In addition, losartan was shown to increase renal blood flow and attenuate renal fibrosis in post-ischemic rats. ¹⁴¹ This result is consistent with a potential role for Ang II blockade to improve oxygenation as a means of suppressing progression. However, blockade of Ang II

signaling is also known to affect glomerular hydrostatic pressure, influence macrophage and lymphocyte activity, as well as reduce tubular expression of pro-fibrotic growth factors such as TGF- β . Finally, as described above, L-arginine increased RBF and attenuated hypoxia and progression to CKD in post AKI rats,⁶⁰ but the putative activation of NO-related events could easily be thought to influence inflammatory processes.

Other investigators have attempted to evaluate whether exacerbation of hypoxia might hasten the development of CKD. For example, reduced renal mass results in a greater degree of renal hypoxia and accelerates the AKI-to-CKD progression.²⁴ This approach would also tend to support a role for hypoxia in CKD progression. However, other studies have shown that reduction in renal mass results in a greater percentage of tubules which fail to redifferentiate completely.¹⁶ As described earlier, undifferentiated tubules may provide a strong driving force for the development of fibrosis. Taken together, while a number of studies seem to support a likely role for hypoxia in progression, experimental maneuvers designed to increase or decrease hypoxia may affect other processes making firm conclusions difficult.

Despite these complications, an interesting study by Friedrich-Persson and colleagues attempted to evaluate if renal tissue hypoxia *per se* may be sufficient to initiate the development of nephropathy, even in the absence of other confounding factors like hyperglycemia, hypertension, oxidative stress or prior tissue damage.¹⁴² In their study, normoglycemic rats received 2, 4-dinitrophenol, a mitochondrial uncoupler that increases oxygen consumption, for 30 consecutive days by oral gavage. This resulted in increased oxygen consumption and a significant reduction of oxygen tension in both the renal cortex and medulla. This treatment resulted in increased proteinuria, infiltration of inflammatory cells and tubular injury as indicated by increased expression of vimentin positive cells. In contrast, renal markers of oxidative stress, mean arterial pressure, renal blood flow, glomerular filtration rate and urinary excretion of Na+ remained unaffected in rats that received chronic dinitrophenol treatment.¹⁴² These studies suggest that renal tissue hypoxia, by itself, has the potential to act as a pathogenic factor in the kidney.

A developing area of interest that may impinge on this conversation may come from information of patients residing in high altitudes. As reviewed by Luks et al. reductions in inspired pO_2 are associated with reduced renal pO_2 .¹⁴³ There are limited studies demonstrating that Navajo Indians living at high altitude have higher rates of ESRD compared with other Native Americans living at low altitudes.¹⁴⁴ Another study compared patients with type 2 diabetes living near sea level or above 1700 meter.¹⁴⁵ Despite a similar level of glycemic control, it was reported that patients at high altitude had increased levels of proteinuria and serum creatinine versus patient near sea level.¹⁴⁵

Future Directions

The last 20 years has seen a dramatic increase in the understanding of the pathophysiology of the AKI to CKD transition and the identification of several molecular pathways mediated by hypoxia. However, it remains to be determined whether hypoxia represents an important mediator of the AKI-to-CKD transition. Few longitudinal studies have been conducted in an

effort to evaluate patients with AKI at risk for progression of CKD. We propose that longterm follow up studies should consider evaluation of renal hypoxia as one of the potential risk factors which may predict future progression. Given the improvement in non-invasive methods for measurement of renal hypoxia such as a newly developed dynamic nuclear polarization-MRI technique would be useful for understanding the role of hypoxia in pathogenesis of AKI-CKD transition. This information might therefore be helpful in identifying whether hypoxia in patients with AKI represents potential risk factor for progression of CKD, and perhaps such approaches could identify which patients with AKI that may require long term follow up care following discharge.

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Figure 1. Hypothesized role of renal hypoxia in the progression of acute kidney injury to chronic kidney disease.

Acute kidney injury is associated with damage to both tubular and vascular compartments. Tubular injury results in increased expression of profibrotic cytokines and growth factors and decreased expression of vascular factors such as VEGF in tubular cells may lead to renal microvascular loss. The reduction of peritubular capillaries restricts downstream blood flow from glomerular capillaries and contributes to the development of renal hypoxia, while also being associated with activation of pericytes toward a fibrogenic myofibroblast phenotype. Hypoxia influences pro-fibrotic and inflammatory activity of tubular and inflammatory cells.

Myofibroblast profibrotic signaling is influenced by factors produced from hypoxic tubules or inflammatory cells or may be influenced directly by hypoxia (?). Progressive interstitial fibrosis exacerbates renal hypoxia by increasing oxygen diffusion distance between peritubular capillaries and parenchyma forming a pathological cycle promoting progressive chronic kidney disease.



Figure 2. Micrographs of Microfil®- perfused kidney sections processed from rats kidney at 4 weeks after sham-operation (A) or post ischemia reperfusion injury (B). Note, renal capillaries are filled with Microfil which appears as bright yellow against a dark background. In post ischemic kidneys, capillary density is reduced in both the cortex and the medulla. Cortex (c), outer stripe of the outer medulla (os), inner stripe of the outer medulla (is), and inner medulla (im) are shown. Images were reproduced from Basile et al. with permission.²⁴

Table 1.

Evidence of capillary drop out following AKI

Model/Condition	Method	Observation	Citation	
Rat I/R	Microfil	30-50% reduction	(Basile et al 2001) ²⁴	
Rat I/R	CD31/PECAM staining Cablin staining	~30% reduction by Cablin staining, preserved by VEGF 121	g, preserved by (Basile et al 2011) ²⁶ , (Leonard et al 2008) ³³	
Rat I/R	CD31 staining	90% loss of CD31 staining SS31	(Liu et al 2014) ³⁵	
Rat I/R	RECA staining	Reduced by $> 50\%$, preserved by EPC microvesicles	(Cantaluppi et al 2012) ³⁶	
Rat I/R	JG12 staining	~50% reduction; exacerbated by reduced renal mass	(Polichnowski et al 2014) ¹⁶	
Mouse I/R	Flurescence micro- angiography	40% reduction in capillaries at 8 weeks; reduced size of remaining vessels. ~60% decrease in total perfusion area	(Kramann et al 2014) ²⁵	
Mouse I/R	Meca-32 staining and Micro-CT	20-43% reductions; tight correlation with Meca-32 and Micro-CT; inversely correlated with collagen I staining	(Ehling et al 2016) ³²	
Mouse I/R	Cablin staining	30-40% after 4 weeks MMP9 loss prevented loss of cablin staining following IR	(Lee et al 2011) ³⁷	
Mouse I/R	CD31 staining	Reduction in staining by 20%, preserved by MSC	(Xing et al 2014) ³⁸	
Mouse I/R	CD31 staining	15-20% rarefaction; some evidence that endothelial proliferation is enhanced in Id1 overexpressing mice	(Lee et al 2011) ³⁹	
DTR transgenic mouse	CD31 staining	Dramatic reduction in CD31 staining associated with fibrosis	(Grgic et al 2012) ⁴⁰	
Human DGF	CD31/CD34 co- staining	Reduced peritubular capillaries in biopsy 3 months post DGF	(Steegh et al 2011) ³⁰	
Human DGF	vWF staining	Evidence of peritubular capillary endothelial cells damage in sustained acute renal failure	(Kwon et al 2008) ²⁹	

Abbreviation: I/R, ischemia reperfusion; PECAM, platelet endothelial cell adhesion molecule; VEGF, vascular endothelial growth factor; RECA, rat endothelial cell antigen; EPC, endothelial progenitor cells; JG12, endothelial aminopeptidase P in renal tissue; Meca-32, endothelial cell antigen; Micro-CT, micro computed tomography; MMP-9, matrix metalloproteinase-9; MSC, mesenchymal stem cells; Id1, inhibitor of DNA-binding/differentiation proteins; DTR, diphtheria toxin receptor; DGF, delayed graft function, Vwf, von Willebrand Factor.

Table 2.

Methods for assessment of renal tissue pO2 in the setting of AKI and AKI-to-CKD transition

Methods	<i>In vivo/</i> processed tissue	Data type	Tissue pO_2 in AKI/ AKI-to-CKD transition	References
Pimonidazole adduct immune histochemistry	paraffin- embedded sections	Qualitative	~5 weeks following recovery from I/R injury, positive pimonidizole staining was observed in the renal outer medulla of rats, positive pimonidizole staining was largely absent in kidney tissue at both 24 h and 5 days following I/R in rats	(Basile et al) ⁶⁰ (Ow et al 2018) ¹¹⁰
HIF immune histochemistry	paraffin- embedded sections/frozen sections	Qualitative	Increased expression of HIF-1a in renal tubules 28 days following sever I/R injury in mice	(Li et al 2019) ¹⁰¹
Clark-type micro electrodes	Anesthetized rat	Quantitative	Evidence of marked outer medullary hypoxia 20 minutes after injection of indomethacin or 10 minutes after injection of sodium iothalamate in rat, Renal tissue pO_2 was consistently greater in rats subjected to ischemia than sham ischemia after 24 h post I/R	(Heyman et al 1991) ¹⁰⁷ (Ow et al 2018) ¹¹⁰
Radio-telemeter based oxygen sensor	Conscious rat	Quantitative	Inner medullary tissue pO_2 did not differ significantly between sham rats and ischemic rats across 5 days after I/R	(Ow et al 2018) ¹¹⁰
BOLD-MRI	Human, rat, mice	Qualitative	Increased R2* values observed 24 hours post I/R in mice and in renal medulla 20 minutes after administration of iodinated contrast media in human	(Oostendorp et al 2011) ¹³⁰ (Hofmann et al 2006) ¹³¹
Urinary oxygen tension	Human, sheep	Quantitative	Urinary pO_2 decreased during 24 hours period of development of sepsis in sheep and in human urinary pO_2 decreased across the course of the cardiac surgery	(Lankadeva et al 2016) ¹²⁶ (Zhu et al 2018) ¹²⁵

Abbreviation: AKI, Acute kidney injury; CKD, Chronic kidney disease; I/R, Ischemia reprfusion; HIF,_Hypoxia inducible factor; BOLD-MRI, Blood oxygen level dependent-magnetic resonance imaging; I/R, Ischemia reperfusion.