

Oxidative Stress in Diabetes and Hypertension

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Oxidative stress implies an increased production of reactive oxygen species (ROS) or a decreased capacity to metabolize them. Recent studies suggested that oxidative stress is implicated in the pathogenesis of tissue injury and dysfunction in diabetes and hypertension, which are two major causes of ESRD. In these diseases, oxygen radicals are increased and contribute to diabetic nephropathy and hypertension by enhancing renal vascular tone, sensitivity to vasoconstrictors, inflammatory cell infiltration and tubuloglomerular feedback. ROS induced nitric oxide inactivation and induction of stress activated signaling pathways are main underlying mechanisms for ROS induced tissue injury and dysfunction. Treatment with pharmacological antioxidant agents such as TEMPOL (superoxide dismutase mimetic) reverses many of these injury and dysfunction underlying the role of oxidative stress in the pathogenesis of diabetic nephropathy and hypertension.

Key Words : Reactive oxygen species, Pathogenesis, Diabetes mellitus, Hypertension

Introduction

Production of the high-energy compounds that fuel the biochemical, biophysical and mechanical functions of the body is coupled with ongoing generation of potentially cytotoxic reactive oxygen species (ROS).

ROS can attack, denature or modify structural and functional molecules and thereby cause cytotoxicity, tissue injury and dysfunction. Oxidative stress, arising as a result of an imbalance between ROS production and antioxidant defenses has been implicated in the pathogenesis of tissue injury and dysfunction in a wide range of human diseases including atherosclerosis, infection, inflammation, neoplasm, degenerative disorders, metabolic diseases and radiation injury^{1, 2)}. There is now accumulating evidence that ROS may participate in renal vascular dysfunction in cardiovascular-renal diseases. Renal dysfunction is a cen-

tral cause of hypertension and a common consequence of diabetes mellitus. These pathophysiological conditions set up a vicious cycle of repeated renal injury and are the two leading causes of end-stage renal failure.

Oxidative stress is associated with both diabetes and hypertension in humans and in experimental animal models. This review will briefly discuss the production, metabolism and targets of ROS, the role of ROS in renal dysfunction during diabetes mellitus and, finally, highlight recent studies that suggest a role for oxygen radicals in the renal vasculature during hypertension.

ROS production and antioxidant defense

1. Superoxide production

A free radical can be defined as any molecular species capable of independent existence that contains an unpaired electron in an atomic orbital¹²⁾. Many radicals are highly reactive and can either donate an electron to or extract an electron from other molecules, therefore behaving as oxidants or reductants.

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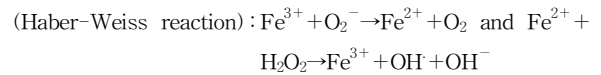
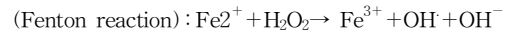
The most important free radicals in many disease states are oxygen derivatives, particularly superoxide (O_2^-) and the hydroxyl radical (OH). Superoxide is produced endogenously during mitochondrial respiration and by NADPH oxidase, xanthine oxidase, cyclooxygenase and lipoxygenase, nitric oxide synthase (NOS) and cytochrome P450³⁾. Several molecules, including adrenaline, flavine nucleotides, thiol compounds, and glucose, can oxidise in the presence of oxygen to produce superoxide, and these reactions are greatly accelerated by the presence of transition metals such as iron or copper²⁾. The electron transport chain in the inner mitochondrial membrane performs the reduction of oxygen to water. During this process free radical intermediates are generated, which are generally tightly bound to the components of the transport chain.

However, there is a constant leak of a few electrons into the mitochondrial matrix and this results in the formation of superoxide. The activity of several other enzymes, such as cytochrome p450 oxidase in the liver and enzymes involved in the synthesis of adrenal hormones, also results in the leakage of a few electrons into the surrounding cytoplasm and hence superoxide formation. There might also be continuous production of superoxide by vascular endothelium to neutralize nitric oxide, production of superoxide by other cells to regulate cell growth and differentiation, and the production of superoxide by phagocytic cells during the respiratory burst.

2. ROS metabolism and antioxidant system

Superoxide spontaneously gains an electron to form hydrogen peroxide; however, three isoforms of superoxide dismutase (SOD) also catalyze this reaction. Mn-SOD is located in mitochondria and two isoforms of Cu,Zn-SOD are located either extracellularly or intracellularly. Once produced, hydrogen peroxide can be scavenged to water by catalase or by glutathione (GSH) peroxidase in the presence of GSH.

Decomposition of hydrogen peroxide in the presence of Fe^{2+} produces a hydroxyl radical, also known as the Fenton reaction²⁾. Superoxide and hydrogen peroxide also can react together directly to produce the hydroxyl radical under the presence of transition metal ion, also known as Haber-Weiss reaction²⁾.



The hydroxyl radical (OH \cdot), or a closely related species, is probably the final mediator of most free radical induced tissue damage. All of the ROS described above exert most of their pathological effects by giving rise to hydroxyl radical formation. The reason for this is that the hydroxyl radical reacts, with extremely high rate constants, with almost every type of molecule found in living cells including sugars, amino acids, lipids, and nucleotides.

3. Targets of ROS

The three main cellular targets of ROS are lipids, proteins, and DNA. Extensive lipid peroxidation in biological membranes causes alterations in fluidity, permeability and membrane potential, and eventual rupture of the cell. Oxidation of proteins changes their primary structure, including the overall charge, folding and hydrophobicity, which can lead to increased aggregation and degradation. Oxygen-radical-induced damage of DNA includes changes in both DNA structure and chemistry with the result being strand breakage.

In addition to their ability to directly inflict damage upon cellular macromolecules, ROS play a significant role in activating stress-sensitive signaling pathways that regulate gene expression resulting in cellular damage⁴⁾. Those genes involved in stress-activated signaling pathways can lead to the development of microvascular complications of diabetes, insulin resistance, inflammation and hypertension⁵⁾.

Whether oxygen radicals attack those targets depends on the delicate balance between levels of ROS and antioxidants. Under many conditions an increase in oxygen radical formation signals the activation of antioxidant enzymes to aid in the increased metabolism necessary to achieve redox balance. However, when the amount of radicals produced exceeds the resources for metabolism, oxidative stress results.

Diabetes mellitus

1. The importance of oxidative stress in the pathogenesis of diabetic complication

1) Oxygen radicals reduce nitric oxide (NO) function in diabetic vessels

Endothelial dysfunction in peripheral and renal vessels is a common sequela of diabetes mellitus⁶. Some observations indicate that the tonic influence of NO is suppressed and contributes to the impaired endothelium-dependent relaxation in the renal vasculature during diabetes⁷. Because superoxide rapidly binds and inactivates NO⁸, one possible explanation for the lack of NO function in the kidney during diabetes is excessive superoxide. Indeed, renal cortical tissue from diabetic rats has increased superoxide production⁹. Several other studies^{10, 11} indicate that in-

creased superoxide in diabetes reduces NO-dependent modulation of basal renal vascular tone, which can be restored by antioxidant treatment.

NO modulates and buffers the renal and peripheral vasoconstriction caused by several endogenous agents, including angiotensin II, thromboxane A₂ and endothelin-1. Because superoxide limits the bioavailability of NO, the buffering capability of NO during agonist-induced vasoconstriction may be decreased in diabetes.

Indeed, Schoonmaker et al.¹² have shown that the renal afferent arteriolar responsiveness to angiotensin II is enhanced in juxtamedullary nephrons from diabetic rats and that L-NNA (NO synthase inhibitor N^w-nitro-L-arginine) did not alter the response. However, treatment with SOD restored the ability of L-NNA to enhance the vascular response to angiotensin II. These data suggest that excess superoxide is responsible for the increased sensitivity of renal microvessels to angiotensin II in diabetes.

In addition to functional NO deficiency, avid reaction of superoxide with NO leads to formation of highly reactive nitrogen species such as peroxynitrite¹³ or peroxynitrous acid¹⁴. The latter agents can, in

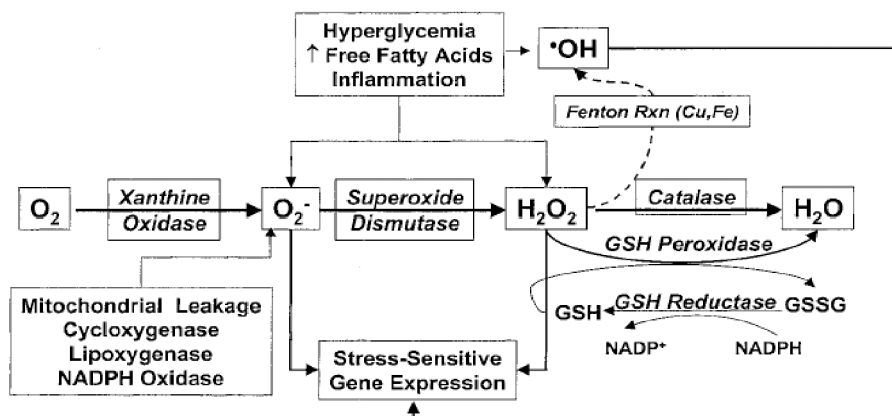


Fig. 1. Endogenous stimuli leading to ROS generation and activation of stress-sensitive gene expression. The endogenous antioxidant enzymes including GSH, superoxide dismutase, GSH peroxidase, and catalase function to maintain redox equilibrium. However, in situations such as chronic hyperglycemia, the compensatory response is inadequate, leading to both ROS formation and activation of stress- and redox-sensitive gene expression (e.g., via the redox-sensitive transcription factor NF- κ B). Catalase is localized primarily in peroxisomes, whereas GSH peroxidase is the major peroxidase in mitochondria [Derived from Ref. 5].

turn, attack, denature or modify various structural and functional molecules^{15, 16}. For instance peroxynitrite can react with tyrosine or cysteine residues of the proteins producing nitrotyrosine or nitrocysteine which are considered as footprint of ROS interaction with NO¹⁷.

2) Oxidative stress and stress-activated signaling pathways

Recent evidence suggests that common stress-activated signaling pathways such as nuclear factor- κ B (NF- κ B), p38 MAPK, NH₂-terminal Jun kinases/stress-activated protein kinases (JNK/SAPK) are activated by oxidative stress and underlies the development of late diabetic complications⁵. Among these, transcription factor NF- κ B is one major intracellular target of hyperglycemia and oxidative stress. The consequence of these signaling pathways is the production of gene products, such as VEGF and others, which cause cellular damage and are ultimately responsible for the long-term complications of diabetes.

3) Formation of glycoxidative advanced glycation end products (AGEs)

Possible mediators of untoward effects of hyperglycemia include AGEs known to accumulate in diabetic subjects. AGEs comprise a variety of molecular structures, such as N^ε-(carboxymethyl)lysine (CML), pentosidine, and pyrroline¹⁸. They are generated by the Maillard reaction through nonenzymatic glycation of protein amino groups and characterized by different formation mechanisms¹⁸. Among them, CML and pentosidine requires both glycation and oxidation for their formation (thereby termed glycoxidation)¹⁹.

Supporting the role of glycoxidative AGE in the pathogenesis of diabetic nephropathy, it was recently demonstrated that among AGE, glycoxidation products such as CML and pentosidine, accumulate in expanded mesangial matrix and nodular lesions in DN, in colocalization with malondialdehyde-lysine (MDA-lysine), a lipoxidation product, whereas pyrroline, another AGE structure whose deposition is rather independent from oxidative stress, was not found

within diabetic glomeruli²⁰. Because CML, pentosidine, and MDA-lysine are all formed under oxidative stress by carbonyl amine chemistry between protein amino group and carbonyl compounds, their colocalization suggests a local oxidative stress and increased protein carbonyl modification in diabetic glomerular lesions.

4) Oxidative stress as a common linking mechanism of hyperglycemia induced damage

It has been suggested that following four pathways are mainly involved in the pathogenesis of long-term diabetic complications: ① increased polyol pathway flux; ② increased AGE formation; ③ activation of protein kinase C (PKC) isoforms; ④ increased hexosamine pathway flux²¹. Recent study suggested that excess ROS produced during mitochondrial respiration is a unifying hypothesis linking these seemingly independent four mechanisms²².

Potential mechanisms by which hyperglycemia-induced ROS (superoxide) overproduction activates four pathways of hyperglycaemic damage are as follows (Fig. 2). Excess superoxide partially inhibits the glycolytic enzyme glyceraldehyde-3-phosphate dehydrogenase (GAPDH), thereby diverting upstream metabolites from glycolysis into pathways of glucose overutilization. This results in increased flux of dihydroxyacetone phosphate (DHAP) to diacylglycerol, an activator of PKC, and of triose phosphates to methylglyoxal, the main intracellular AGE precursor. Increased flux of fructose-6-phosphate to UDP-N-acetylglucosamine increases modification of proteins by O-linked N-acetylglucosamine (GlcNAc) and increased glucose flux through the polyol pathway consumes NADPH and depletes GSH as described below.

2. Sources of oxygen radicals in diabetes mellitus

For hyperglycemia, increases in oxidant productions are due to multiple processes. Simply, glucose can undergo non-enzymatic reactions forming gluco-oxi-

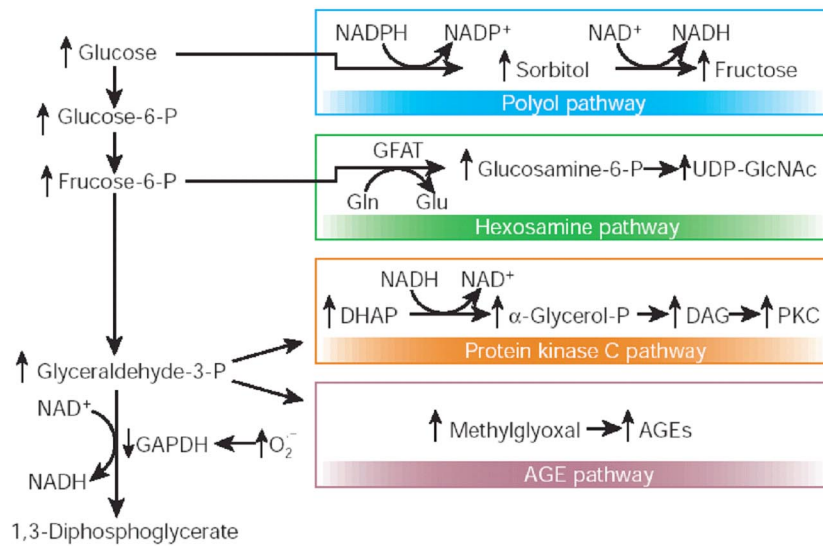


Fig. 2. Potential mechanism by which hyperglycemia-induced mitochondrial superoxide overproduction activates four pathways of hyperglycemic damage.

dants and glycated products (AGEs), which themselves can be pro-oxidants^{23, 24}. In addition, high glucose-induced ROS production in renal mesangial cell can be effectively blocked by inhibition of PKC, NADPH oxidase, and mitochondrial electron transfer chain complex, suggesting that PKC, NADPH oxidase, and mitochondrial metabolism all play a role in high glucose-induced ROS generation²⁵.

Among those possibilities, recent focus has been on mitochondrial metabolism and activation of NADPH oxidases²⁴. Suggestions have been made that most glucose-induced oxidants are derived from glycolysis and mitochondrial oxidative phosphorylation with the productions of superoxide²¹. Electron transfer through mitochondrial enzyme complexes I, III and IV generates a proton gradient that drives ATP synthase (complex V). When the electrochemical potential difference generated by this proton gradient is high, the life of superoxide-generating electron transport intermediates is prolonged resulting in leakage of superoxide.

There appears to be a threshold value above which superoxide production is markedly increased.

Hyperglycemia increases the proton gradient above this threshold value as a result of overproduction of

electron donors by the Krebs cycle, thereby increasing superoxide production²¹.

Metabolism of high glucose levels can activate NADPH oxidase in the vascular cells independent of mitochondrial metabolisms²⁶. One mechanism that can increase NADPH oxidase activity is the activation of PKC, which is elevated by glucose-induced elevation of diacylglycerol²⁶. It has been also established that angiotensin II stimulates vascular superoxide formation through activation of NADPH oxidase²⁷ and angiotensin-converting enzyme (ACE) inhibition attenuates the oxidative stress in the diabetic kidney²⁸.

These studies intimate that hyperglycemia and angiotensin II induced oxygen radical formation mediated by NADPH oxidase play a role in the oxidative stress of the diabetic renal microvasculature.

Among other possibilities, it has been suggested that activation of the polyol pathway could be another source of oxidative stress²¹. In polyol pathway reduction of glucose to sorbitol by NADPH consumes NADPH. As NADPH is required for regenerating reduced GSH, this could induce or exacerbate intracellular oxidative stress.

Koo et al.²⁹ showed that insulin therapy in streptozotocin diabetic rat resulted in a significant increase

in renal NOS expression. This is consistent with the known effect of insulin on NO production and endothelial NOS gene expression in endothelial cells via activation of phosphatidylinositol-3 kinase³⁰⁾. In their study²⁹⁾, tissue nitrotyrosine abundance, which is a marker of highly toxic and reactive nitrogen species was paradoxically increased in insulin treated diabetic rat (Fig. 3). This paradoxical result was partially explained by the upregulation of NOS isoforms in all tested tissues and the expected rise in NO production capacity.

This coupled with the residual oxidative stress with incomplete glycemia control in insulin-treated animals can account for the observed elevation of tissue nitrotyrosine abundance. In this regard, it has been suggested that inadequate glucose control with insulin treatment could be another mechanism of oxidative stress in diabetic kidney²⁹⁾.

Hypertension

Oxidative stress in the vasculature has been associated with human essential hypertension, pre-eclampsia and several hypertensive animal models, including the spontaneously hypertensive rat (SHR), angiotensin II-induced hypertension, Dahl salt-sensitive hyperten-

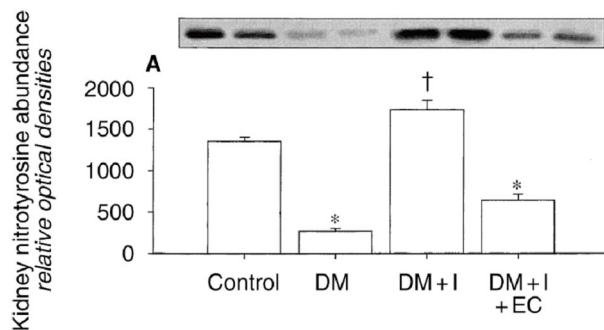


Fig. 3. Representative Western blots and corresponding group data illustrating nitrotyrosine abundance of kidney cortex in the normal control rats fed regular diet ($N=6$), untreated diabetic rats (DM, $N=6$), diabetic rats treated with once-daily ultralente insulin alone (DM+I, $N=5$) or the combination of insulin and vitamin E and C-fortified diet (DM+I+EC, $N=5$). * $P<0.005$ versus control and DM+I groups; † $P<0.05$ versus other groups [Derived from Ref. 29].

sion, lead-induced hypertension, obesity-induced hypertension, mineralocorticoid hypertension and hyperglycemia-induced hypertension^{3,31)}. Most studies indicate that antioxidant treatment lowers blood pressure and improves endothelial function in large conduit vessels^{3,31)}. Considering the major role of kidney in the pathogenesis of hypertension, above studies strongly suggested that increased oxidative stress is important in the renal vascular dysfunction in hypertension.

1. Oxygen radicals reduce NO function in hypertension

SHR has increased blood pressure and renal vascular resistance and an enhanced tubuloglomerular feedback response. Acute and long-term studies suggest that oxygen radicals may play an important role in these characteristics. The SOD mimetic TEMPOL (2,2,6,6-tetramethyl-1-piperidinoxy) normalizes the blood pressure, renal vascular resistance and renal excretion of the oxidative stress marker 8-iso prostaglandin $F_{2\alpha}$ in the SHR^{32,33)}. Furthermore, TEMPOL increases the basal luminal diameter of *in vitro* perfused afferent arterioles of juxtamedullary nephrons in SHR but has no effect in the Wistar-Kyoto rat (WKY)³⁴⁾. These studies suggest that oxygen radicals may contribute to the increased blood pressure and renal vasculature resistance in the SHR.

1) Redox regulation of the afferent arteriole and tubuloglomerular feedback

One of the potential mechanisms for the acute reduction in blood pressure and renal vascular resistance in SHR treated with antioxidants, is via enhancing NO action in the renal vasculature or in the juxtaglomerular apparatus. NOS type 1 or neuronal NOS is expressed in the macula densa. Renal macula densa derived NO counteracts afferent arteriolar vasoconstriction mediated by the tubuloglomerular feedback mechanism and thus control intraglomerular pressure and sodium and water excretion^{35,36)}. Systemic inhibition of NOS with N^{ω} -nitro-L-arginine

methyl ester (L-NAME, NOS inhibitor) blocks the acute anti-hypertensive actions of TEMPOL in SHR³²⁾. This study implies that NO-mediated vasodilation may be restored after scavenging of oxygen radicals in the SHR. The afferent arteriole and macula densa cell both contain a full complement of components of NADPH oxidase that generates superoxide^{35, 37)}. Several recent studies have shown extensive interaction between superoxide and NO in macula densa to regulate afferent arteriolar tone mediated by the tubuloglomerular feedback response^{35, 38)}. Superoxide in the macula densa and afferent arterioles will inactivate NO and enhance preglomerular resistance in animal models of oxidative stress³⁵⁾. As an increase in afferent arteriolar resistance can precede hypertension, oxidative stress could be important in determining the long-term blood pressure and thereby contribute to hypertension.

2) Role of oxidative stress in the angiotensin II induced hypertension

In recent studies, Welch and Wilcox³⁹⁾ showed that two weeks of treatment with the selective angiotensin AT₁ receptor antagonist candesartan in SHR restored a normal tubuloglomerular feedback response to NOS inhibitor 7-nitroindazole but had no effect on the response to 7-nitroindazole during TEMPOL microperfusion. These data suggest that angiotensin II stimulation of oxygen radical formation in the juxtaglomerular apparatus, mediated via AT₁ receptors, has an important role in elevating oxygen radicals and thereby diminishing NO signaling in the juxtaglomerular apparatus of SHR.

Angiotensin II-induced stimulation of oxygen radical formation, which can diminish NO action, is not specific to the SHR. Nishiyama et al.⁴⁰⁾ demonstrated that the increased blood pressure and renal vascular resistance in angiotensin II-infused hypertensive rats was ameliorated by TEMPOL. NOS inhibition markedly attenuated the hemodynamic response to TEMPOL. These studies strongly implicate angiotensin II as a major source of oxygen radical formation

in forms of hypertension that are dependent on or sensitive to angiotensin II. More importantly, Higash et al.⁴¹⁾ showed that angiotensin II mediated excessive oxidative stress is involved, at least in part, in impaired endothelium-dependent vasodilatation in patients with renovascular hypertension. In their study⁴¹⁾, correction of renal artery stenosis decreases oxidative stress with normalization of endothelial dysfunction. These results further support the role of oxidative stress in angiotensin II dependent hypertension.

2. Dys-regulation of antioxidant defenses in hypertension

In study using DNA microarray analysis to explore differentially-expressed genes in kidneys between SHR and WKY rat, several genes involved in antioxidant system such as gamma glutamylcysteine synthetase, GSH S-transferase and heme oxygenase 3 genes were down-regulated in SHR as compared to WKY rat⁴²⁾.

The changes in the expression of these 3 genes were confirmed by real time RT-PCR (Table 1). Gamma glutamylcysteine synthetase is the rate-limiting enzyme for GSH synthesis. GSH S-transferase adds GSH to electrophiles with a variety of chemical structures and protects cells from oxidative stress. Thus, down-regulations of gamma glutamylcysteine synthetase and GSH S-transferase can lead to GSH deficiency and defective protection against oxidative injury. These events can, in turn contribute to oxidative stress and hypertension in SHR. The latter proposition is supported by the result of previous study in which induction of oxidative stress by GSH depletion caused severe hypertension in genetically normal rats⁴³⁾. Heme oxygenase which exists in constitutive (heme oxygenase-2 and heme oxygenase-3) and inducible (heme oxygenase-1) isoforms degrades pro-oxidant heme to produce carbon monoxide and bilirubin, which is a potent antioxidant⁴⁴⁾. Endogenous carbon monoxide is a potent vasodilator and pos-

esses anti-inflammatory and anti-apoptotic properties. Therefore, down-regulation of constitutive heme oxygenase 3 isoform as shown in this study could potentially contribute to oxidative stress, inflammation and hypertension in SHR.

3. Oxygen radicals and inflammation in hypertension

Another possible mechanism for the renal protective actions of antioxidants in hypertension is through a reduction in the immune and inflammatory responses. ROS can act as second messengers for several transcription factors, including NF- κ B, which plays a critical role in the activation of multiple genes that contribute to the inflammatory response and end-organ damage³⁾. In the angiotensin II-dependent hypertension⁴⁵⁾ and deoxycorticosterone (DOCA) salt hypertension⁴⁶⁾ animal models, NF- κ B activation is increased in the kidney and renal monocyte/macrophage infiltration is increased. In association with a reduction in vascular oxygen radical formation, TEMPOL also reduces the blood pressure, NF- κ B activation and monocyte/macrophage infiltration in the kidneys of DOCA salt hypertensive rats⁴⁶⁾.

The transient administration of angiotensin II or NOS inhibitor, L-NAME to rats result in salt-sensitive hypertension that persists after removal of the inciting stimulus, and this is associated with the infiltration of immunocompetent cells and tissue injury in the renal interstitium^{47, 48)}. Many of the mononuclear cells, including some T cells, were shown to express angiotensin II, and there was also local gen-

eration of oxidants^{47, 48)}. Tubulointerstitial injury resulting from angiotensin II infusion was significantly reduced by anti-inflammatory treatment using mycophenolate mofetil, as were proliferative activity, T-cell infiltration and activation, superoxide-producing cells, and urinary excretion of marker of oxidative stress, MDA.

Similar results was obtained in SHR, in which mycophenolate mofetil treatment reduce blood pressure, oxidative stress and inflammatory cell infiltration in kidney⁴⁹⁾. In DNA microarray analysis of SHR, the complement regulatory membrane protein CD59 gene was down-regulated in SHR as compared to WKY rat⁴²⁾ (Table 1). Because the complement regulatory membrane protein, CD59, restricts membrane attack complex formation, it is likely that down-regulation of CD59 gene can potentially facilitate infiltration of immunocompetent cells in the kidney and, thereby, aggravate hypertension, tissue injury and vascular complication in SHR. These results signify that some of the renal vascular damage in hypertension may be due to the proinflammatory actions of oxygen radicals and inflammation with oxidative stress could be a possible mechanism of hypertension.

Conclusions

Oxidative stress is increased with diabetes mellitus and hypertension. In diabetes, hyperglycemia-induced activation of PKC, NADPH oxidase and mitochondrial metabolism play major role in ROS generation. ROS induced inactivation of NO and induction of stress-

Table 1. Comparison of Expression of mRNA Involved in Antioxidant and Anti-inflammatory System between SHR and WKY Rat

Gene	Normalized signal intensity from microarray analysis		Ratio (SHR/WKY) from real time RT PCR
	WKY	SHR	
Heme oxygenase-3	2.39	0.45	0.25
Glutathione S-transferase	1.65	0.51	0.34
Gamma glutamylcysteine synthetase	1.47	0.57	0.53
CD 59 antigen	1.39	0.66	

activated signaling pathway seems to be important mechanisms for the chronic diabetic complications including diabetic nephropathy. Recent studies suggest that increased production of superoxide by the mitochondrial electron transport chain is a causal link between elevated glucose and each of the four main pathways responsible for hyperglycaemic damage i.e. increased polyol pathway flux; increased AGE formation; activation of PKC; increased hexosamine pathway flux. In experimental hypertension, antioxidant defenses systems are down regulated at least in SHR, an animal model of human essential hypertension. ROS induced inactivation of NO seems to contribute to the increased basal vascular tone and sensitivity of the vasculature to angiotensin II and enhanced tubuloglomerular feedback shown in hypertensive animal model. It is also suggested that some of the renal vascular damage in hypertension may be due to the proinflammatory actions of oxygen radicals and inflammation with oxidative stress could be another possible mechanism of hypertension. Treatment with pharmacological agents that mimic SOD, such as TEMPOL, reduces vascular dysfunction in the kidney during diabetes. In animal models of hypertension, TEMPOL decreases markers of oxidative stress, improves renal vascular function and reduces blood pressure. These studies provide a rationale for the development of new pharmacological agents that target oxygen radicals for the treatment of renal dysfunction in hypertension and diabetes mellitus.

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