The Role of Oxidative Stress in the Metabolic Syndrome

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Loss of reduction-oxidation (redox) homeostasis and generation of excess free oxygen radicals play an important role in the pathogenesis of diabetes, hypertension, and consequent cardiovascular disease. Reactive oxygen species are integral in routine in physiologic mechanisms. However, loss of redox homeostasis contributes to proinflammatory and profibrotic pathways that promote impairments in insulin metabolic signaling, reduced endothelial-mediated vasorelaxation, and associated cardiovascular and renal structural and functional abnormalities. Redox control of metabolic function is a dynamic process with reversible pro- and anti-free radical processes. Labile iron is necessary for the catalysis of superoxide anion, hydrogen peroxide, and the generation of the damaging hydroxyl radical. Acute hypoxia and cellular damage in cardiovascular tissue liberate larger amounts of cytosolic and extracellular iron that is poorly liganded; thus, large increases in the generation of oxygen free radicals are possible, causing tissue damage. The understanding of iron and the imbalance of redox homeostasis within the vasculature is integral in hypertension and progression of metabolic dysregulation that contributes to insulin resistance, endothelial dysfunction, and cardiovascular and kidney disease. [Rev Cardiovasc Med. 2011;12(1):21-29 doi: 10.3909/ricm0555]

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The prevalence of cardiovascular disease (CVD) and chronic kidney disease (CKD) is progressively increasing in the United States, a phenomenon that closely parallels the epidemics of obesity and the metabolic syndrome.¹⁻¹² Approximately 70 million adults in the United States are obese and another 70 million have hypertension.¹⁻⁵ Data from the National Health and Nutrition Examination Survey indicate that the prevalence of hypertension increases progressively with increasing body mass index (BMI).³⁻⁵ Recent work supports the notion that obese patients exhibit more frequent impairments of insulin metabolic signaling, dyslipidemia, microalbuminuria, and hypertension—all components of the metabolic syndrome.⁶⁻⁹

The metabolic syndrome is a complex clinical syndrome that is associated with a substantial risk for CVD events such as stroke, congestive heart failure, CKD, and overall mortality.¹⁻⁹ CVD in the United States represents a continuing crisis of epidemic proportion. There are an estimated 64.4 million people in the United States who suffer from CVD.¹⁰⁻¹² CVD is often perceived as the disease of the elderly, but recent data suggest approximately 50% of CVD diagnoses and 15% of CVD deaths are in patients younger than 65 years. In addition, the metabolic syndrome has been shown to be independently related to CKD and to the development of acute kidney injury (AKI). CKD is an accepted CVD risk state, meaning that CKD contributes to incident and prevalent CVD. In almost every case, CKD renders the treatment of CVD more difficult and more hazardous. Thus, the metabolic syndrome poses a unique cardiorenal risk on a chronic basis, which is the set-up for acute CVD and AKI events that lead to excesses in end-stage renal disease and premature mortality.¹³ The growing incidence of CKD and CVD among the younger population can be attributed to the increasing incidence of obesity in the United States.

Obesity is characterized by the presence of insulin resistance and is now known to herald even greater CVD risk.⁶⁻¹³ As a result of excess adiposity, insulin resistance is central to the metabolic syndrome and implies impaired biologic and physiologic response to insulin. Insulin resistance is often accompanied by decreased insulin-dependent metabolic signaling and impaired glucose transport and utilization in skeletal muscle, liver, and fat.^{14,15} Fatty steatosis (or excesses in intracellular fatty acids and triglycerides), as well as the presence of visceral adipose

tissue, are cardinal features of insulin resistance. Thus, accentuation of some actions of insulin metabolic effects with concurrent resistance gives rise to diverse clinical manifestations and sequelae of the metabolic syndrome.

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Herein we describe the important mechanisms that contribute to the generation of oxidative stress in the vasculature and kidney that promote endothelial dysfunction and metabolic dysregulation, which contribute to the metabolic syndrome.

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result of excess adiposity, subclinical elevations of proinflammatory markers from the liver and adipose tissue (including interleukin-6, C-reactive protein, plasminogen activator inhibitor-1, and fibrinogen) are linked with the development of impaired insulin-dependent glucose utilization.¹⁵ The presence of insulin resistance is also associated with inappropriate activation of the reninangiotensin-aldosterone system (RAAS) due to the presence of angiotensinogen from adipose tissue.¹⁶ Mounting evidence supports the concept that angiotensin II (Ang II), acting through its type 1 receptor (AT_1R) , induces insulin resistance by inhibiting the actions of insulin in vascular and skeletal muscle tissue.¹⁷⁻²⁰ It does so by promoting oxidative stress and endothelial dysfunction through increased nicotinamide adenine dinucleotide H (NADH) and nicotinamide adenine dinucleotide phosphate (NADPH) oxidase activity and dependent generation of labile irondependent reactive oxygen species (ROS). Thereby, oxidative stress contributes to alterations in redox-sensitive insulin-dependent signaling through phosphatidylinositol-3-kinase and protein kinase B signaling. These changes promote metabolic dysregulation observed in obesity that manifests as insulin resistance and

The Importance of Insulin Resistance

Obese subjects are noted to have elevated levels of insulin, which is required to maintain glucose and fatty acid metabolism, and often have insulin resistance in traditional insulin-sensitive tissues such as skeletal muscle, liver, and adipose tissue, but also in nontraditional insulin-sensitive tissues such as heart, aorta, and kidney.²¹⁻²⁴ In the insulin-resistant state there is impaired biologic and physiologic response to insulin in tissue. However, this resistance to insulin is not uniform in all tissues. This altered response is often accompanied by decreased insulin-mediated metabolic signaling and impaired glucose transport/utilization, as well as impaired nitric oxide (NO)induced vasodilatation in skeletal muscle and cardiovascular tissue.²⁵⁻²⁸ Thus, accentuation of some actions of insulin metabolic effects, with parallel resistance to other actions, gives rise to diverse clinical manifestations associated with the metabolic syndrome. Sequelae that result from insulin actions on activation of the sympathetic nervous system include salt sensitivity, inappropriate activation of the RAAS, and increases in inflammation and oxidative stress.

There is also mounting evidence that supports the idea that excess

insulin (eg, hyperinsulinemia) as a compensatory mechanism in insulin resistance contributes to further adipose tissue deposition. In several models, excesses of insulin promote available glucose and fatty acid transport into adipocytes, therefore increasing cell size and fat mass. In addition, insulin and stimulated insulin-like growth factors drive the conversion of preadipocytes to mature and growing adipocytes, ready to accumulate more substrate and begin producing adipokines, which worsen the insulin resistance and proinflammatory state.²⁹

Role of RAAS in Oxidative Stress and the Metabolic Syndrome

Despite a state of relative volume expansion and salt sensitivity, the RAAS is inappropriately activated in obese individuals who display insulin resistance. Recent work suggests this effect is due to elevated levels of circulating aldosterone associated with increases in visceral adiposity.³⁰⁻³⁵ Preclinical and clinical reports support this relationship between BMI and elevations in RAAS markers such as plasma renin activity, plasma angiotensin-converting enzyme (ACE), and circulating Ang II.³⁴⁻³⁹

Mounting data support that adipose tissue displays a local RAAS. Data support that visceral tissue has higher levels of angiotensinogen (AOGEN) than subcutaneous tissue, and also expresses the AT₁R, which suggests an autocrine-paracrine role for visceral adipose tissue.⁴⁰ Some investigators postulate that adipose tissue RAAS may actually regulate systemic blood pressure to some degree. This is best illustrated in studies in transgenic AOGEN mice in which production is limited to adipose tissue-support that AOGEN produced in fat cells can enter the circulation and further regulate blood pressure and sodium homeostasis.^{40,41}

Traditional doctrine has been that Ang II is the main effector peptide of the RAAS and signals through the AT₁R and angiotensin type 2 receptor (AT₂R) with second messenger signaling that classically oppose each other. Ang II signaling through the AT₁R contributes to the deleterious effects and vasoconstrictor actions of Ang II that are opposed by binding of the AT₂R under routine physiologic conditions. It should also be noted that there are many other duction of oxidative stress through the enzyme complex NADPH oxidase. Aldosterone has been shown to act both directly and indirectly through potentiation of an Ang II effect that contributes to impairments in endothelium-dependent relaxation.^{15,40,41} This is associated with increased vascular oxidative stress resulting in reductions in bioavailable NO.

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effects of Ang II generally considered have endocrine, exocrine, to paracrine, and autocrine effects. Ligand receptor binding of the Ang receptors also leads to G protein versus non-G protein-mediated effects. G protein-mediated effects may be mediated through phospholipase C with formation of 1,4,5-inositol and diacylglycerol, whereas non-G protein-mediated effects happen through stimulation of tyrosine kinases. Both of these pathways eventually contribute to activation of components of NADPH oxidase and other metabolic oxidases, and generation of superoxide anion $(\cdot O_2^{-})$ and other free radicals observed in the metabolic syndrome.34-38

Although Ang II is classically thought to exert many of the profibrotic, proinflammatory, and prooxidative effects, recent work highlights what may be more important—direct actions mediated through Ang II stimulation of aldosterone production. It is now increasingly recognized that aldosterone has actions that are independent of Ang II stimulation. Most notably are aldosterone actions that augment AT₁R signaling that promote vascular proincreases in collagen synthesis and fibrosis, resulting in arterial stiffness, left ventricular hypertrophy, and renal fibrosis.⁴⁰⁻⁴⁶ As alluded to above, these maladaptive changes are augmented through a relative cross-talk between aldosterone and Ang II. Analogous to Ang II, aldosterone acts via rapid, nongenomic actions, mediating endothelial dysfunction and cardiac remodeling in animal models and in humans.^{42,43} Aldosterone stimulates pericytes and macrophages to secrete galectin-3, an important paracrine factor that signals local fibroblasts to secrete procollagen, which is then cross-linked to collagen in the extracellular matrix.44 In rodent models of both tissue Ang II and aldosterone excess, antagonism of the target of aldosterone (eg, the mineralocorticoid receptor) has been shown to reduce cardiac tissue oxidative stress through reductions in NADPH oxidase with improvements in interstitial fibrosis, remodeling, hypertrophy, and, ultimately, diastolic function.45,46

Vascular NADPH Oxidase

The vascular enzyme complex NADPH oxidase, which is induced in

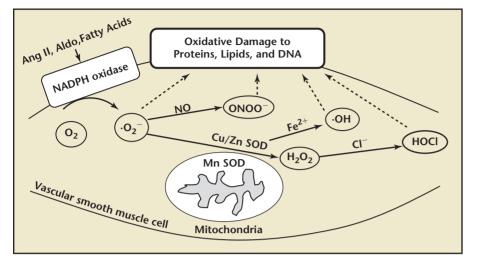


Figure 1. The role of oxidant stress in endothelial dysfunction and vascular tissue injury. Ang II, angiotensin II; Aldo, aldosterone; GSH, glutathione; NADPH, nicotinamide adenine dinucleotide phosphate; NO, nitric oxide; SOD, superoxide dismutase.

part by fatty acids, as well as Ang II and aldosterone, is largely considered the major source of ROS production in cardiovascular and kidney tissue (Figure 1).²³⁻²⁵ NADPH oxidase represents the enzyme responsible for much of the generation of $\cdot O_2^{-}$ in cardiovascular tissue that contributes to endothelial dysfunction, in addition to cardiac and renal tissue remodeling.¹³ The enzyme complex is composed of several membrane and cytosolic subunits that mobilize and activate under various agonists, such as fatty acids, Ang II, and aldosterone.47,48 The membrane complex consists of two subunits, gp91^{phox} (now referred to as NOX2) and $p22^{phox}$, which together form a heterodimer known as flavocytochrome b₅₅₈. The cytosolic subunits include p47^{phox}, which is considered the strategic organizer subunit when activated.49 Another cytosolic subunit, p67^{phox}, facilitates the transfer of the hydride ion from NADPH to flavin adenine dinucleotide upon interaction with the small G-protein Rac through the tetratricopeptide domain. Rac exists in an inactive state when bound to guanosine diphosphate; however, phospho-

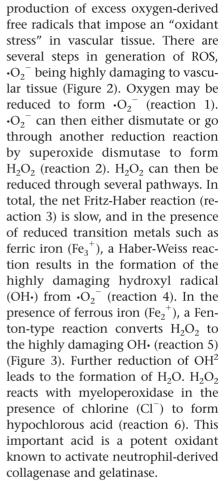
rylation results in activation (eg, Racguanosine triphosphate [GTP]) and translocation to the membrane complex. $p40^{phox}$ then binds to $p67^{phox}$ and facilitates the assembly of $p47^{phox}$ $p67^{phox}$ at the membrane. The final transfer of an electron to form $\cdot O_2^-$ is facilitated by two heme groups present within the NADPH complex.

Reactive Oxygen Species

ROS are important to routine physiologic and cellular functions.^{13,14} However, it is the imbalance in the metabolism of oxygen and resultant

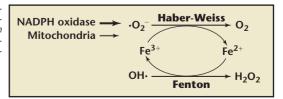
Figure 2. Pathways of generation of reactive oxygen species.

Figure 3. The Haber-Weiss reaction generates hydroxyl radicals from hydrogen peroxide and superoxide as a source of reaction oxygen species and involves reduction of ferric ion to ferrous. NADPH, nicotinamide adenine dinucleotide phosphate.



The generation of ROS and free oxygen radicals has been shown to regulate pathways involved in cell signaling, apoptotic pathways, cell and tissue growth, and salt and

Reaction $1 \rightarrow O_2 + e^- \rightarrow \cdot O_2^-$
Reaction $2 \rightarrow \cdot O_2^- + \cdot O_2^- + 2H^+ \rightarrow O_2^- + H_2O_2$
Reaction $3 \rightarrow H_2O_2 + \cdot O_2^- \rightarrow OH^- + OH \cdot + O_2$
Reaction 4 \rightarrow Fe ³⁺ + •O ₂ ⁻ \rightarrow Fe ²⁺ + O ₂
Reaction 5 \rightarrow H ₂ O ₂ + Fe ₂ ⁺ \rightarrow Fe ₃ ⁺ + OH ⁻ + OH·
Reaction $6 \rightarrow H_2O_2 + CI^- + H^+ \Leftrightarrow H_2O + HOCI$



fluid homeostasis. Free radicals such as $\cdot O_2^-$, hydroxyl moiety ($\cdot OH$), hypochlorite (ClO⁻), peroxynitrite (ONOO⁻), and protein and lipid species, are short lived and scavenged by a series of antioxidant enzymes. However, repeated agonist exposure generates an imbalance of redox reactions and thereby contributes to an excess of ROS and results in oxidative stress.

In the context of insulin sensitivity and obesity, Ang II, aldosterone, fatty acids, and even high salt levels contribute to increased production of $\cdot O_2^{-}$ and H_2O_2 . ROS are known promoters of downstream signaling pathways that include transcription factors, tyrosine kinases/phosphatases, ion channels, and mitogenic factors. The impact of ROS generated within various components of the vasculature (eg, heart, aorta, or kidney) is subject to local concentrations and the balance of pro- and antioxidant mechanisms. This is especially pertinent with the increasing interest in the availability of catalytic iron as a potential source for vascular ROS as described above.⁵⁰ Although there are a variety of normal molecular mechanisms to manage H_2O_2 , there are fewer that can manage the hydroxyl radical, which is generated from H_2O_2 by the catalysis of Fe_2^+ to Fe_3^+ . This is especially pertinent in chronic disease, including the various risk components of CMS for CVD and CKD.¹³⁻¹⁶ It is interesting to note the recent appreciation of siderocalin, otherwise known as neutrophil gelatinase lipocalin, as a marker of both chronic CVD and CKD.⁵¹ This protein attempts to manage catalytic iron and works as a protective factor against labile iron-associated oxidative stress mechanisms.⁵²

Redox Control of Vascular Function

Free radicals are generally divided into ROS and reactive nitrogen

species (RNS) generated for specific cellular processes: smooth muscle relaxation, inhibition of platelet adhesion, cell growth and differentiation, and numerous second messenger systems that promote impairments in insulin metabolic signaling.

Although the predominant form of ROS is $\cdot O_2^{-}$, critical to the metabolic phenotype is the NO radical (NO•).¹³ NO• is produced by oxidation of one of the terminal nitrogen atoms of L-arginine, a reaction that is then catalyzed by nitric oxide synthase (NOS). Levels of bioavailable NO, a potent vasodilator, diminish as NO is then consumed by ROS. NO is converted to other RNS; however, it is the predominance of $\cdot O_2^{-}$ that rapidly reacts with NO to yield ONOO⁻. Importantly, ONOO⁻ oxidizes NO synthase, which leads to decreased production of NO. Another notable reaction between ROS and NO is the oxidation of tetrahydrobiopterin (BH₄). In the absence of BH₄, NOS forms $\cdot O_2^{-}$ instead of NO, a process referred to as NO uncoupling, which is another critical source of oxidative stress that promotes endothelial dysfunction observed in the metabolic syndrome.

ROS also regulate various signaling mechanisms, and in excess promote derangements in cell cycle progression, growth, and proliferation, as well as insulin metabolic signaling. Oxidative stress is thought to contribute to the aging process and chronic disease states due to their electrophilic character, which promotes oxidization of cell constituents such as proteins, lipids, and DNA. The paradoxical nature of ROS can be resolved by understanding that only excessive production of these radicals results in damage, whereas their roles as mediators of cell signaling are temporally and spatially controlled. This is illustrated by the fact that ROS are known to

oxidize/reduce cysteine residues, a mechanism critical for mitogenactivated protein kinase, protein tyrosine phosphatases, protein tyrosine kinase, transcription factors, and other enzymes that are redox regulated. Thereby, redox-sensitive pathways regulate the cells to activate or inhibit signaling proteins and dynamically alter gene expression according to various agonist/antagonist conditions.

Therapeutic Interventions That Improve the Redox State

There is strong experimental evidence for targeting the RAAS to improve oxidative stress, particularly in the treatment of hypertension.^{15,16} The impact of the RAAS has classically been illustrated through the maladaptive effects of Ang II signaling through the AT_1R_1 as described above. Accordingly, there is sufficient evidence for a beneficial effect inhibition of ACE and blockade of AT₁R on oxidative stress.^{10,13,15} Preclinical data from our laboratory and others, in various metabolic rodent models (including transgenic rats that over-express the RAAS), suggest that inhibition of the AT₁R or the mineralocorticoid receptor (MR), and recently with renin inhibition, improves indices of oxidative stress that result in improved cardiovascular remolding and metabolic parameters. On a clinical level, RAAS inhibition through ACE inhibition and Ang II receptor blockers has been studied extensively in patients with hypertension, congestive heart failure, coronary artery disease, and CKD, and is recommended to prevent CVD and nephropathy in patients with type 2 diabetes mellitus.53,54 Recent data with blockade of the MR have shown promise in patients with heart failure, CKD, and resistant hypertension.¹⁶

In the context of the metabolic syndrome, some studies suggest a beneficial effect of RAAS blockade on insulin resistance and glucose home-ostasis.^{55,56} Possible mechanisms responsible for the reduced incidence of diabetes in these trials include

concentrations are greater than aldosterone. MR activation by glucocorticoids further potentiates oxidative stress observed in the metabolic syndrome.

Evidence also suggests that statins may have pleiotropic effects beyond

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improvement in insulin-mediated glucose uptake, enhanced endothelial function, increased NO activation, reduced inflammatory response, and increased bradykinin levels.¹³ These effects have largely been described in the context of ACE inhibition and AT_1R blockade. However, the impact of MR antagonism or direct renin inhibition on insulin resistance and glucose homeostasis remains to be fully elucidated.^{57,58}

From an experimental perspective, in vivo MR antagonism results in decreased NADPH oxidase activity and oxidative stress in concert with improved insulin-dependent glucose uptake, as well as attenuated whole body insulin resistance in skeletal muscle.57 Nonetheless, there have not been definitive clinical studies to confirm previous experimental and clinical observations suggesting improved insulin sensitivity through MR blockade. The discrepancy may be due to several factors. Most importantly, in the context of the metabolic syndrome, the MR has a high affinity for both aldosterone and 11β-hydroxy glucocorticoids. The presence of much lower levels of this enzyme in nonepithelial tissues allows for glucocorticoids to signal through the MR in cardiovascular and metabolic tissue such as skeletal muscle, liver, and fat. This is of particular importance in individuals with the metabolic syndrome, in whom circulating glucocorticoid their ability to reduce lipids through modulatory effects on oxidant stress. Statins act by blocking 3-hydroxy-3methylglutaryl coenzyme A reductase and inhibit synthesis of mevalonic acid, a precursor of many nonsteroidal isoprenoid compounds involved in intracellular trafficking of proteins involved in oxidative stress injury (ρ , Ras, Rac, Rab, Ral, and Rap).⁵⁹ Statins inhibit the activation of Rac1, critical to activation of NADPH oxidase, by preventing the geranylgeranyl-dependent translocaoxidized LDL.^{63,64} Regardless of the mechanisms, there is convincing evidence to support that both inhibition of the RAAS and statin therapy in humans ameliorates oxidative stress.

Is There a Role for Chelation Therapy?

There is increasing interest in chelation therapy, especially in the context of targeting poorly liganded iron in the generation of oxidative stress. Iron chelators have been used in various studies that examined therapeutic interventions in the management of coronary disease,65,66 as well as the investigation of kidney disease.⁶⁷ Recently, there have been several reports targeting pharmacologic interventions,^{68,69} such as desferrioxamine and the aforementioned statins, on poorly liganded iron and improvements in endothelial function.^{66,70} Although there is increasing interest in pharmacologic intervention, there is

Although there is increasing interest in pharmacologic intervention, there is also a considerable role for dietary nutrients in the chelation of iron and mitigation of the effects of oxidative stress.

tion of Rac1 from the cytosol to the cell membrane, thereby reducing ROS generation.⁶⁰⁻⁶² Statins have also been shown to block expression of protein subunits of G-proteins (p22^{phox} and NOX2), which determine oxidase activity of NADPH oxidase and expression of GTP-ase (NADPH activator).⁶² This then contributes to suppression of prooxidant enzyme systems. Statins have also been shown to improve bioavailable NO and interfere with low-density lipoprotein (LDL) oxidation by several mechanisms, including reducing serum levels of LDL available for oxidation and the expression of scavenger receptor CD36 on monocytes that inhibit uptake of also a considerable role for dietary nutrients in the chelation of iron and mitigation of the effects of oxidative stress. The best evidence exists with polyphenolic compounds, many of which are widely used as dietary antioxidants.^{71,72} This has classically been attributed to a Mediterranean diet, wherein the beneficial effects of the polyphenols are thought to be derived from their antioxidant capabilities.73-76 Many of the polyphenolic compounds, including flavones, isoflavones, stilbenes, and flavanones, are implicated for their antioxidant effects because they have been shown to chelate iron as well.77-80 The flavonoid compounds have a catechol moiety that are known iron-binding elements of microbial siderophores.⁸¹ It is no surprise, then, that a considerable number of studies on chronic disease states wherein oxidative stress has been implicated show promise with interventions containing polyphenolic components. However, the uptake of these compounds in the routine management of chronic diseases has been hampered by significant problems in determining absorption, bioavailability, and measures of appropriate markers.

Conclusions

Overfeeding and excess adiposity directly result in insulin resistance. As a response, excesses in insulin potentiate the growth and division of additional adipocytes. These cells-in concert with hepatocytes-produce adipokines and cytokines, which are responsible for systemic inflammation. In the setting of inflammation, poorly liganded or labile iron available within cells through catalysis allows the Haber-Weiss and Fenton equations to generate ROS and the particularly deleterious hydroxyl radical. These processes are fundamental to tissue fibrosis and potentially to instability of atherosclerotic plaque and incident CVD events. Very importantly, chronic CVD and

CKD are the important base conditions upon which acute events occur. These events, mediated by tissue hypoxia, generate a much larger quantity of labile iron and allow an explosion of local oxidative stress that results in tissue injury even after the hypoxia has been relieved. Understanding the role of iron and oxidative stress in the setting of obesity and insulin resistance as observed in the metabolic syndrome is critical to advancing diagnostic and therapeutic strategies in the future.

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Main Points

- Loss of reduction-oxidation (redox) homeostasis and generation of excess free oxygen radicals play an important role in the pathogenesis of diabetes, hypertension, and consequent cardiovascular disease.
- Reactive oxygen species are integral in routine in physiologic mechanisms. However, loss of redox homeostasis contributes to proinflammatory and profibrotic pathways that promote impairments in insulin metabolic signaling, reduced endothelial-mediated vasorelaxation, and associated cardiovascular and renal structural and functional abnormalities.
- Redox control of metabolic function is a dynamic process with reversible pro- and anti-free radical processes. Labile iron is necessary for the catalysis of superoxide anion, hydrogen peroxide, and the generation of the damaging hydroxyl radical. Acute hypoxia and cellular damage in cardiovascular tissue liberate larger amounts of cytosolic and extracellular iron that is poorly liganded; thus, large increases in the generation of oxygen free radicals are possible, causing tissue damage.
- The understanding of iron and the imbalance of redox homeostasis within the vasculature is integral in hypertension and progression of metabolic dysregulation that contributes to insulin resistance, endothelial dysfunction, and cardiovascular and kidney disease.

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