

Review Article

Oxidative Stress-Related Biomarkers in Essential Hypertension and Ischemia-Reperfusion Myocardial Damage

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Cardiovascular diseases are a leading cause of mortality and morbidity worldwide, with hypertension being a major risk factor. Numerous studies support the contribution of reactive oxygen and nitrogen species in the pathogenesis of hypertension, as well as other pathologies associated with ischemia/reperfusion. However, the validation of oxidative stress-related biomarkers in these settings is still lacking and novel association of these biomarkers and other biomarkers such as endothelial progenitor cells, endothelial microparticles, and ischemia modified albumin, is just emerging. Oxidative stress has been suggested as a pathogenic factor and therapeutic target in early stages of essential hypertension. Systolic and diastolic blood pressure correlated positively with plasma F2-isoprostane levels and negatively with total antioxidant capacity of plasma in hypertensive and normotensive patients. Cardiac surgery with extracorporeal circulation causes an ischemia/reperfusion event associated with increased lipid peroxidation and protein carbonylation, two biomarkers associated with oxidative damage of cardiac tissue. An enhancement of the antioxidant defense system should contribute to ameliorating functional and structural abnormalities derived from this metabolic impairment. However, data have to be validated with the analysis of the appropriate oxidative stress and/or nitrosative stress biomarkers.

1. Introduction

Cardiovascular diseases are a leading cause of mortality and morbidity worldwide [1], with hypertension being a major risk factor [2]. These diseases affect more than 600 million people and it is estimated that there will be 29% of the world adult population suffering hypertension by 2025 [3]. The pathophysiology of cardiovascular diseases is complex due to the multiple biological pathways that have been implicated. Accordingly, the absolute risk of future cardiovascular events cannot be estimated from a single biomarker. These diseases are originated in the vascular endothelium, a target for the classical cardiovascular risk factors. Following endothelial activation oxidative stress has an important role in the development of atherosclerosis [4] and hypertension [5, 6], thereby contributing to the progression of the structural and functional cardiovascular damage. In cardiovascular disease related to ischemia/reperfusion injury, redox imbalance triggers the activity of a number of signaling pathways mediated by reactive oxygen species

(ROSs) and reactive nitrogen species (RNSs) [7]. Consequently, in cardiac surgery with extracorporeal circulation, electrical and structural myocardial remodeling due to the excessive production of these reactive species may lead to the development of arrhythmias such as atrial fibrillation [8]. Furthermore, percutaneous transluminal coronary angioplasty following acute myocardial infarction results in heart reperfusion damage, thus enhancing the infarct size [9]. These cardiac abnormalities are associated with major changes in oxidative stress-related biomarkers. However, to date there is no available data allowing to have an optimal biomarker approach with the ability to help identify patients at risk for cardiovascular disease, to monitor the efficacy of treatments, and to develop new pharmacological tools. Further studies are still lacking to achieve the validation of oxidative stress-related biomarkers to implement strategies suitable to improve the prevention and reduction of cardiovascular risk. The aim of this review was to present a novel focus on the bases supporting the applicability of reliable biomarkers accounting for the antioxidant potential, as well

as oxidative stress and nitrosative stress associated with clinical settings of cardiovascular diseases. Particularly, we analyze the biomarkers related to ischemia/reperfusion and essential hypertension, two examples of processes mediated by ROS and RNS. This novel strategy, together with other biomarkers, could contribute to the development of changes in health diagnostics and clinical research.

2. Oxidative Stress

The oxidation and reduction reactions in biological systems (redox reactions) represent the basis for numerous biochemical mechanisms of metabolic changes [10]. Oxidative stress constitutes a unifying mechanism of injury of many types of disease processes, and it occurs when there is an imbalance between the generation of reactive oxygen and nitrogen species and the antioxidant defense systems in the body, so that the latter become overwhelmed [11]. ROS are widely recognized as important mediators of cell growth, adhesion, differentiation, senescence, and apoptosis [12]; however, when these species are found in excess they might cause cell damage either directly or working as intermediates in diverse signaling pathways. RNS may have also a similar behavior.

In pathophysiological conditions, the main sources of ROS include the mitochondrial respiratory electron transport chain, xanthine oxidase (XO) activation through ischemia-reperfusion, the respiratory burst associated with neutrophil activation, and arachidonic acid metabolism. In the cellular metabolism, the oxygen molecule itself is reduced to water after forming, as successive intermediates, superoxide anion ($O_2^{\bullet -}$), hydrogen peroxide (H_2O_2), and hydroxyl radical. It is estimated that between 0.15 and 2% of cellular O_2 consumption results in superoxide formation [13].

Xanthine oxidoreductase functions as either xanthine dehydrogenase (XDH) form, which transfers electron to NAD^+ , and generates NADH or XO form, which transfers electron to O_2 and generates oxidative stress. Due to ischemia-induced cellular calcium overload converts XDH to XO, under reperfusion phase, enhanced XO can produce more ROS, such as superoxide, H_2O_2 , and hydroxyl radicals [14].

Oxidants generated by cellular NADPH oxidases participate in many biological processes, serving both as critical elements of signaling pathways and important effector molecules [15–20]. The enzyme NADPH oxidase catalyzes the one electron reduction of O_2 to generate $O_2^{\bullet -}$, using NADPH as the source of electrons. There appear to be at least three isoforms of NADPH oxidase expressed in the vascular wall. Furthermore, they are distributed in nearly all cell types and tissues, and many cells express more than one NADPH oxidase protein isoform [21].

Nitric oxide (NO) synthases (NOS) are a family of enzymes that convert the amino acid L-arginine to L-citrulline and NO. Three main isoenzymes exist in mammals that are regulated by distinct genes: a constitutive neuronal NOS (nNOS or NOS-1), an endotoxin- and cytokine-inducible NOS (iNOS or NOS-2), and a constitutive endothelial NOS (eNOS or NOS-3) [22]. NOS-1 and NOS-3

are constitutively expressed and Ca^{+2} /calmodulin dependent, while the inducible form (NOS-2) is induced by inflammatory stimuli and is Ca^{+2} /calmodulin independent [23]. eNOS plays a major role in the regulation of vascular function. For instance, eNOS synthesizes NO by a two-step oxidation of the amino acid L-arginine, thereby leading to activation of guanylyl cyclase. The resulting second messenger cGMP in turn activates the cGMP-dependent kinase, which leads to decrease in intracellular Ca^{+2} concentrations, thereby causing vasorelaxation. However, it has become clear from studies with the purified enzyme that eNOS may become uncoupled in the absence of the NOS substrate L-arginine or the cofactor tetrahydrobiopterin. Uncoupled eNOS state results in the production of $O_2^{\bullet -}$ instead of NO [24, 25].

Myeloperoxidase is a neutrophil and monocyte enzyme that amplifies the reactivity of H_2O_2 through generation of hypochlorous acid, free radicals, and RNS. The myeloperoxidase enzyme (MPO) produces hypochlorous acid from H_2O_2 and chloride anion during the neutrophil's respiratory burst. It requires heme as a cofactor. In addition, it oxidizes tyrosine to tyrosyl radical using H_2O_2 as oxidizing agent [26]. Both hypochlorous acid and tyrosyl radical are cytotoxic and used by the neutrophil to kill bacteria and other pathogens.

In physiological conditions, cells have an antioxidant defense system to prevent ROS and RNS damage. Antioxidant substances are small molecules that can scavenge free radicals by accepting or donating an electron to eliminate the unpaired condition. Typically, this means that the antioxidant molecule becomes a free radical in the process of scavenging a ROS to a more stable and less reactive molecule. In most cases the scavenger molecule provides hydrogen radical that combines with the free radical. Consequently, a new radical that has an enhanced lifetime compared with the starting one is generated, for instance, due to a conjugated system [27]. However there are many mechanisms whereby antioxidants may act such as: (1) scavenging reactive oxygen species or their precursors, (2) inhibiting the formation of ROS, (3) attenuating the catalysis of ROS generation via binding to metal ions, (4) enhancing endogenous antioxidant generation, and (5) reducing apoptotic cell death by upregulating the anti-death gene Bcl-2 [28].

Enzymatic antioxidant defenses include superoxide dismutase (SOD), glutathione peroxidase (GSH-Px), and catalase (CAT), among others. Nonenzymatic antioxidants include a variety of mostly exogenous biological molecules, such as ascorbic acid (vitamin C), α -tocopherol (vitamin E), reduced glutathione (GSH), carotenoids, flavonoids, polyphenols, and other exogenous antioxidants [29].

The first line of cellular defense against oxidative injury includes the antioxidant enzymes CAT, SOD, and GSH-Px [30]. The SOD converts superoxide anion to H_2O_2 , which is a substrate for CAT and GSH-Px. Catalase metabolizes H_2O_2 to water and oxygen and GSH-Px reduces both H_2O_2 and organic hydroperoxides when reacting with GSH [31]. Reduced glutathione is present at high concentrations in all mammalian cells, especially in the renal cells, hepatocytes, and erythrocytes [32]. GSH protects protein thiol groups from nonenzymatic oxidation or can act as a cosubstrate of GSH-Px [33].

Exogenous antioxidants, such as vitamins C and E, exist at a number of locations, namely, on the cell membrane, intracellularly, and extracellularly. They react with ROS to either remove or inhibit them. The hydrophobic lipid inside of membranes requires a different spectrum of antioxidants. Vitamin E is thought to be the most important antioxidant found within lipid membranes in the body [34].

Polyunsaturated fatty acids (PUFAs) are prone to be oxidized by free radicals and other ROSs [35], which may be alleviated by dietary supplementation of antioxidants such as vitamin C, vitamin E, and carotenoids [36]. Nevertheless, the administration of repeated high doses of PUFA may render the organism less susceptible to lipid peroxidation due to adaptive intracellular mechanisms involving enhancement of the antioxidant response via Nrf2 activation [37]. Increasing extracellular concentrations of fatty acids and low-density lipoproteins induce iNOS expression in many cell types including pancreatic β -cells, vascular smooth muscle cells, and macrophages [38]. Similarly, feeding a high saturated-fat diet to rats increases iNOS activity in liver and colon [39] and stimulates free radical production and oxidative DNA damage in skeletal muscle mitochondria and the whole body [40, 41].

Cellular effects of ROS will rely on its cellular exposure levels [42]. Cell exposure to low to moderate ROS levels triggers a survival response and reinforces ROS scavengers of the antioxidant defence system to elicit a cytoprotective effect. The molecular mechanism believed to be responsible for this adaptive change involves enhanced antioxidant activity achieved by upregulating several housekeeping genes partly under the control of nuclear factor erythroid 2-related factor-2 (Nrf2) [43]. Nrf2 is normally sequestered in the cytosol by Kelch-like ECH-associated protein 1 (Keap1). Upon oxidative stimulation, ROS could either oxidise Nrf2 or covalently modify Keap1 thiol groups, leading to Nrf2 dissociation from Keap1 and subsequent nuclear translocation. In the nucleus, Nrf2 binds to antioxidant response elements (AREs) in target gene promoters [44], which increase the expression of antioxidant enzymes such as catalase, glutathione peroxidase, glutathione reductase, glutathione S-transferase, NADPH:quinone oxidoreductase 1, and heme oxygenase-1 [37].

Cell exposure to high levels of ROS induces nuclear factor kappaB (NF- κ B) activation, which triggers a proinflammatory response characterized by increased levels of TNF- α , IL-1 β , IL-6, and IL-8, increased expression of adhesion molecules, such as E-selectin, VCAM-1, and ICAM-1, and promotion of oxidative stress. NF- κ B is one member of a ubiquitously expressed family of Rel-related transcription factors that serve as critical regulators of inflammatory related genes such as tumor necrosis factor (TNF) and IL-1 [45]. NF- κ B mainly exists in the cytosol as a preformed trimeric complex that consists of the inhibitory protein I- κ B and the P50/P65 protein dimer. ROS induces redox changes that result in phosphorylation of the I- κ B subunit, thereby activating its proteolytic digestion. When the inhibitor subunit is dislodged from the P60/P65 heterodimer, NF- κ B can translocate to the nucleus, bind DNA, and initiate transcription [42].

ROS can mediate cellular damage through direct interaction with cellular components. The redox reaction mediated by ROS determines an electron loss in a target molecule, which alters its structure and function. Virtually any cellular component can be a target of ROS interaction. Lipids, DNA, and proteins have been shown to interact with ROS and mediate cellular damage [46], given that it is not surprising that chronic and acute exposure to excessive levels of ROS can induce cell death processes (apoptosis, necrosis, and autophagy) [47–50].

Acute myocardial infarction (AMI) is a clinical model of oxidative stress by ischemia reperfusion. ROSs are major initiators of myocardial damage during ischemia/reperfusion. In pathophysiological conditions, AMI is initiated by myocardial ischemia due to coronary artery obstruction leading to an initial increase in ROS levels associated with a loss of antioxidant enzyme function, together with leakage of antioxidant enzymes into the extracellular fluid which in normal physiological conditions, kept oxygen free radical production under homeostatic control, and as a result, the unbalanced burst of free radicals on reperfusion easily overwhelms the available counteractive enzymes, so that control of ROS generation is lost [28]. ROSs including H₂O₂, O₂^{•-}, hydroxyl radical, and peroxynitrite have been shown to increase upon reperfusion of the heart following ischemia [51]. However, ROS seems to increase significantly after a few minutes of reperfusion, but its increase during ischemia alone is still controversial.

After an ischemic episode of the myocardium, left ventricle remodeling is known to occur; although its underlying mechanism is multifactorial, ROS and inflammatory cytokines may cause cardiodepressive reaction [52–54]. It is of interest to remark that ROSs also stimulate the production of inflammatory cytokines, and, in turn, inflammatory cytokines stimulate ROS formation. In chronic stage, ROS and inflammatory cytokines activate the matrix metalloproteinases [55, 56], thereby eliciting degradation of collagens which may cause a slippage in myofibrillar alignment causing left ventricular dilatation [57]. Oxidative stress may result in cellular defects including a depression in the sarcolemmal Ca²⁺-pump ATPase and (Na + K)-ATPase activities, changes leading to decreased Ca²⁺-efflux and increased Ca²⁺-influx, respectively [58]. The depression in Ca²⁺-regulatory mechanism by ROS ultimately results in intracellular Ca²⁺ ([Ca²⁺]_i) overload and cell death. On the other hand, an increase in [Ca²⁺]_i during ischemia induces the conversion of xanthine dehydrogenase to xanthine oxidase and subsequently results in generating superoxide radicals [59].

Oxidative stress can participate in the development of hypertension. Compelling experimental evidence indicates that ROSs play an important pathophysiological role in the development of hypertension. This is due, in large part, to O₂^{•-} excess and decreased NO bioavailability in the vasculature and kidneys, and to ROS-mediated cardiovascular remodeling [60–62]. In human hypertension, biomarkers of systemic oxidative stress are elevated [63]. Physiologically, ROSs are produced in a controlled manner at low concentrations and function as signaling molecules

[64] to maintain vascular integrity by regulating endothelial function and vascular contraction-relaxation balance. Under pathological conditions, increased ROS bioactivity leads to endothelial dysfunction, an essential pathophysiological mechanism in the development of hypertension. Reactive oxygen species can inhibit the three major endothelium-dependent vasodilator pathways, that is, NO, prostacyclin, and endothelial derived hyperpolarizing factor (EDHF) [65]. Superoxide anions not only reduce the bioavailability of NO but also directly inhibit its main target, soluble guanylyl cyclase [66–68]. Furthermore, superoxide anions can react with NO leading to the formation of peroxynitrite (ONOO^-) which uncouples NO synthase by oxidizing, on one hand, the Zn-thiolate complexes within the enzyme [69] and, on the other hand the essential cofactor tetrahydrobiopterin (BH_4) [70]. When endothelial nitric oxide synthase (eNOS) becomes unstable and uncoupled, it leads to subsequent less NO production and more superoxide generation [71]. Additionally, ONOO^- inhibits guanylyl cyclase, inactivates the prostacyclin synthase by tyrosine nitration, and further enhances oxidative stress by inhibiting superoxide dismutase [66–72].

Furthermore, increased ROS levels promote vascular smooth muscular cells growth, increased contractility, monocyte invasion, lipid peroxidation, inflammation, and increased deposition of extracellular matrix proteins, all of which are important factors in hypertensive vascular damage [73, 74]. A schema showing the cellular response to ROS is depicted in Figure 1.

3. Biomarkers

Epidemiologic literature that focuses on antioxidant status and chronic disease risk has in the past relied primarily upon biomarkers of exposure to antioxidant nutrients. However, this approach is in essence using exposure estimates to a select group of nutrients as a surrogate for estimating actual oxidative defense or oxidative stress status. Emphasis is now being placed on developing functional biomarkers of oxidative stress status, that is, biomarkers that integrate the effect of exposure to oxidants coupled with the full range of antioxidant protective mechanisms *in vivo* [75]. Many of such biomarkers are being studied including various measures of oxidation products of lipid, DNA, and protein (Table 1). Some of these biomarkers are now being applied in research of pathologies related to oxidative stress.

Lewandrowski et al. suggest the following features for an ideal cardiac biomarker, among others: (1) High concentration in the myocardium, (2) absence from nonmyocardial tissue, (3) high sensitivity and specificity in blood samples, (4) rapid release into the blood following myocardial injury, (5) favorable kinetics permitting detection of acute coronary syndromes and myocardial necrosis in days after the person seeks care, and (6) correlation between blood level of marker and extent of myocardial injury and prognosis [76].

3.1. Oxidative Stress-Related Biomarkers. Lipid peroxides have been estimated by measurement of thiobarbituric acid

reactive substances in plasma by the method of Ohkawa et al. [77]. In this method, in the peroxidation process, peroxides are decomposed to aldehydes like malondialdehyde (MDA), which can be detected by thiobarbituric acid that gives an easily measurable pink color. All of these are termed thiobarbituric reactive species (TBARS) [78]. This method is one of the most widely used assays to assess peroxidation in the whole organism. The reaction of peroxynitrite with lipids leads to peroxidation and MDA and conjugated diene formation [79]. It is noteworthy also that MDA products may behave as toxic bifunctional electrophiles, due to reactivity with proteins, phospholipids, and DNA, generating stable products at the end of a series of reactions to form propane adducts [80]. Consequently, a change in the properties of the molecule, for example, in its charge profile, could result in modified cell-matrix interactions [81]. Lipid peroxides, normally present in lipoproteins or membranes, are known to further induce lipid peroxidation, inhibit the mitochondrial electron transport system, and to oxidize sulfhydryl groups on proteins, hence altering its function or otherwise disrupting signal transduction pathways [82].

The nonenzymatic peroxidation of arachidonic acid results in the formation of F₂-isoprostanes, which are also products of lipid peroxidation. The levels of these compounds can serve as an index of lipid peroxidation in several diseases [83, 84]. Because of their mechanism of formation, specific structural features that distinguish them from other free radical-generated products and their chemical stability, they can provide a reliable index of lipid peroxidation *in vivo* in a variety of clinical settings associated with oxidative stress [85]. In addition, this determination provides a noninvasive method for dose selection to monitor the response to antioxidant treatment [86].

Another lipid peroxidation product is 4-hydroxynonenal (4-HNE), a highly cytotoxic and reactive α , β -aldehyde that is generated during various physiological and pathophysiological conditions based on the production of ROS [87, 88]. 4-HNE is also considered one of the most specific and sensitive measures of lipid autooxidation [89, 90]. HNE-4 can be measured by standard methods [91, 92].

Vitamin E can be measured by the method of HPLC method [93]. The determination of alpha-tocopherol in human serum is reported. UV-Detection is possible by using either one wavelength of 300 nm or after a lambda change mode with typical wavelengths for alpha-tocopherol (292 nm). According to short retention times (10 min) and rapid extraction the method is useful for clinical research and allows about 50 analyses per day and operator. Vitamin C (ascorbic acid) can be estimated by the method of Roe and Kuether [94]. This involves oxidation of ascorbic acid by copper followed by treatment with 2,4-dinitrophenylhydrazine that undergoes rearrangement to form a product with absorption maximum at 520 nm [95].

Total antioxidant activity is measured by the ferric reducing antioxidant power (FRAP) assay of Benzie and Strain [96]. FRAP is a test which measures the combined antioxidant effect of the nonenzymatic defenses in a biological fluid. At low pH, when a ferric-tripyridyltriazine (FeIII-TPTZ)

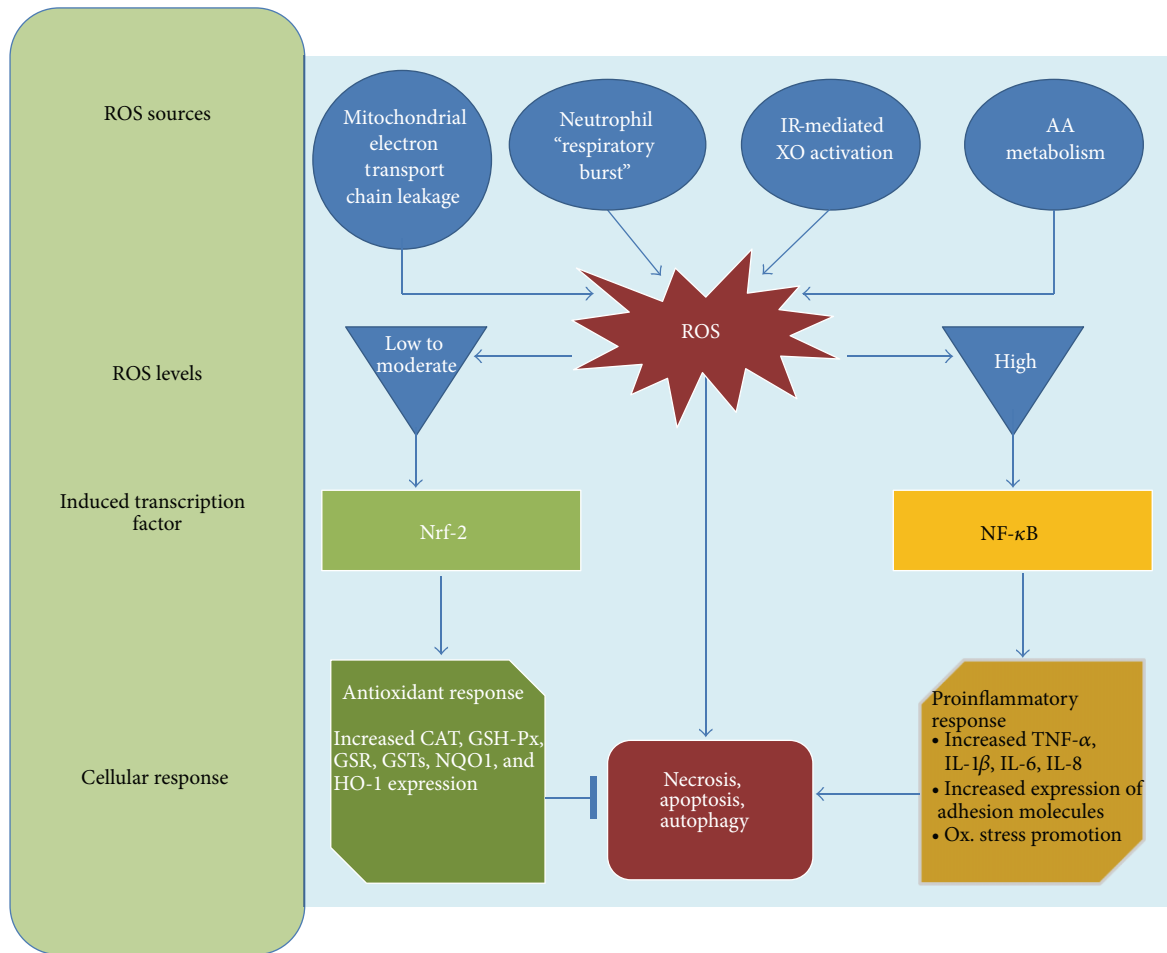


FIGURE 1: Cellular response to reactive oxygen species. Several pathophysiological conditions determine an increase on intracellular ROS levels. Transcription factor induction is dependent in the ROS levels. Low to moderate ROS levels induce Nrf-2 activation which increases the expression of antioxidant enzymes such as catalase, glutathione peroxidase, glutathione reductase, glutathione S-transferase, NADPH:quinone oxidoreductase 1, and heme oxygenase-1 while high levels of ROS induce NF-κB activation, which trigger a proinflammatory response characterized by increased levels of TNF-α, IL-1β, IL-6, and IL-8, increased expression of adhesion molecules, such as E-selectin, VCAM-1, and ICAM-1, and promotion of oxidative stress. AA: arachidonic acid; CAT: catalase; GSH-Px: glutathione peroxidase; GSR: glutathione reductase; glutathione S-transferase: GSTs; HO-1: heme oxygenase-1; NADPH:quinone oxidoreductase 1: NQO1; IL-1 β: Interleukin 1β; IL-6: Interleukin 6; IL-8: Interleukin 8; NF-κB: nuclear factor kappa-light-chain-enhancer of activated B cells; Nrf-2: nuclear factor- (erythroid-derived 2-) like 2.

complex reduces the ferrous (FeII) form, an intense blue color with an absorption maximum at 593 nm develops.

Another oxidative stress biomarker is the protein carbonylation. The direct damage of proteins during oxidative stress can give rise to the formation of protein carbonyls, which may serve as biomarkers for general oxidative stress, in addition to the data provided by lipid peroxidation. Most of the assays [97, 98] for the detection of protein CO groups involve derivatisation of the carbonyl group with 2,4-dinitrophenylhydrazine (DNPH), which leads to formation of a stable dinitrophenyl (DNP) hydrazone product. This then can be detected by various means, such as spectrophotometric assay, enzyme-linked immunosorbent assay (ELISA), and one-dimensional or two-dimensional electrophoresis followed by western blot immunoassay. At present, the measurement of protein CO groups after their derivatisation

with DNPH is the most widely utilized measure of protein oxidation [99].

3.2. Antioxidant Potential Biomarkers

Superoxide Dismutase (SOD). The term superoxide dismutase characterizes a family of proteins differing in their structure and cofactors, among them being Mn-SOD and Cu-Zn-SOD. SOD activity enhances the spontaneous dismutation of superoxide radicals to H₂O₂ [46]. SOD can be measured utilizing the technique of Misra and Fridovich [100] based on inhibition of the formation of nicotine amide adenine dinucleotide, phenazine methosulfate, and amino blue tetrazolium formazan.

Catalase (CAT). The end product of the dismutation reaction –H₂O₂– can be removed by the activity of the enzyme

TABLE 1: Biomarkers of oxidative stress.

Type of biomarker	Location	Reference
Ferric reducing ability of plasma	Extracellular	Benzie and Strain, 1996 [96]
Carbonyls	Extracellular	Mohanty et al., 2010 [98]
Lipid peroxidation		
Malondialdehyde	Extracellular	Ohkawa et al., 1979 [77]
F2-isoprostane	Extracellular	Collins et al., 1999 [202]
4-Hydroxynonenal	Extracellular	Halliwell and Gutteridge, 2007 [89]
Plasma vitamins		
Vitamin C	Extracellular	Roe and Kuether, 1943 [94]
Vitamin E	Extracellular	Teissier 1996 [93]
Antioxidant enzymes		
Superoxide dismutase	Intracellular	Misra and Fridovich, 1972 [100]
Catalase	Intracellular	Aebi and Bergmeyer, 1974 [101]
Glutathione peroxidase	Intracellular	Flohé and Günzler, 1984 [102]
GSH/GSSG ratio in erythrocyte	Intracellular	Hissin and Hilf, 1976 [104]
Prooxidant enzymes		
Xanthine oxidase	Intracellular	Haining and Legan 1967 [106]
NADPH oxidase	Intracellular	Nauseef, 2008 [17, 18]
Others		
Endothelial microparticles	Extracellular	Burger and Touyz, 2012 [109]
Endothelial progenitor cells	Extracellular	Touyz and Schiffrin, 2012 [111]
Ischemia modified albumin	Extracellular	Sinha et al., 2003 [194]

catalase. CAT is an enzyme with a very high K_M for its substrate and can remove H_2O_2 present in high concentrations [46]. The activity of CAT can be measured through the non-spontaneous decomposition of hydrogen peroxide, by the method described by Aebi [101].

Glutathione Peroxidase (GPx). The biochemical function of glutathione peroxidase is to reduce lipid hydroperoxides to their corresponding alcohols and to reduce free H_2O_2 to water. In contrast to catalase, peroxidase possesses high affinity for and can remove H_2O_2 even when it is present in low concentration [46]. Its activity can be estimated by the method described by Flohé and Günzler [102].

Thiol Index. GSH/GSSG ratio in erythrocyte. The GSH/GSSG index is a parameter of intracellular redox status. GSH is the major endogenous antioxidant produced by the cells, participating directly in the neutralization of free radicals and reactive oxygen compounds, as well as maintaining exogenous antioxidants such as vitamins C and E in their reduced forms [103]. GSH and GSSG levels can be measured in erythrocytes by the method described by Hissin and Hilf [104].

Xanthine Oxidase (XO). It is a key enzyme in reactive oxygen species formation, which plays a major role in cell oxidative stress. This enzyme catalyzes the oxidation of hypoxanthine to xanthine and can further catalyze the oxidation of xanthine to uric acid [105]. Xanthine oxidase can be estimated by the method of Haining and Legan [106].

NADPH Oxidase. The NADPH-oxidase complex utilizes electrons to produce superoxide radicals from the oxygen

molecule [46]. NADPH oxidase may be measured through chemiluminescence [107] or electrochemistry [108] methods, among others. Approaches to quantitate oxygen consumption, extracellular release of $O_2^{\bullet-}$ or H_2O_2 , and intracellular $O_2^{\bullet-}$ production provide reliable assessment of NADPH oxidase activity in a given population of cells [21].

3.3. Other Biomarkers of Cardiovascular Damage. Endothelial microparticles (EMPs) are nuclear fragments of cellular membrane shed from stressed or damaged cells. Typically defined as having a diameter of 0.1 to 1.0 μm , microparticles contain surface proteins and cytoplasmic material of the parental cells [109].

Flow cytometry has been largely used to quantify EMP levels in clinical samples, although this widely used methodology suffers from limitations according to MP probing, sizing, and counting in a standardized manner. For probing, EMP detection still presents the limitations of lacking fully specific antigens. CD144 was proposed as a marker for EMP detection. Others strategies have been developed as the combination of multicolor antibodies (CD31+/CD41-, CD31+/CD42b-, CD105+/CD45-) and monochrome composite markers (CD144+ CD105+, CD146+ CD105+) [110].

Endothelial progenitor cells have been a biomarker of cardiovascular damage. Of the various factors important in the protection of the endothelium are endothelial progenitor cells (EPCs). EPCs are bone marrow-derived cells capable of developing into mature endothelial cells [111]. EPCs are mobilized from the bone marrow into the peripheral blood in response to tissue ischemia or injury and through their differentiation improve blood flow and tissue repair [112].

The precise criteria for identifying EPCs by flow cytometry remain a contentious issue; however, the combined surface expression of CD133, CD34, and VEGFR2 is generally accepted as the most stringent criteria for EPCs [113, 114].

Ischemia modified albumin (IMA) has been another biomarker of cardiovascular damage. Under physiological conditions, transition metals can bind tightly to the exposed N-terminus of albumin [115]. In the presence of myocardial ischemia, structural changes take place in the N-terminus of the protein [116, 117] which reduce its binding capacity, possibly, in part, as a result of exposure to ROS. IMA rises in patients who develop myocardial ischemia during percutaneous coronary intervention (PCI) [84]. Serum IMA can be measured by the albumin cobalt binding (ACB) test. The first generation ACB test method has been validated and described in previous publications [118, 119].

4. Essential Hypertension

Hypertension has become an emerging epidemic and important worldwide public-health challenge, because it is one of the most common chronic conditions [120–123]. The importance of high blood pressure as a major cause of serious diseases has been recognized in most western countries for more than 50 years [124]. Essential or primary hypertension is a heterogeneous disorder in which secondary causes are not present. It accounts for 95% of all cases of hypertension [125], with different patients having different causal factors that lead to high blood pressure. The identification of population at risk is a challenge that requires laboratory blood tests; however, biomarkers to ensure regular monitoring of patients having essential hypertension are not available to date. Evidence has been provided that activation of the vascular endothelial cells in the presence of cardiovascular risk factors promotes oxidative stress and vascular inflammatory response, for either classic (smoking, aging, hypercholesterolemia, hypertension, hyperglycemia, and a family history of premature atherosclerotic disease) and novel (obesity, elevated C-reactive protein, and chronic systemic infection) risk factors, leading to acceleration of atherosclerotic vascular disease [126]. Therefore, appropriate biomarkers of these processes should give some clues for the management of early metabolic changes, otherwise leading to stages of permanent and progressive vascular lesions. Several studies have demonstrated excessive amounts of ROS in essential hypertensive patients and various animal models of hypertension [6, 46, 127–130].

4.1. Role of ROS in Vascular Homeostasis. The vasculature is a rich source of ROSs, which are produced in endothelial, adventitial, and smooth muscular cells. The regulation of vascular tone may be impaired by changes affecting the interaction between these vascular cells [131]. The endothelium senses mechanical stimuli, such as pressure and shear stress, as well as chemical stimuli induced by vasoactive substances. In response, it releases agents that regulate vasomotor function, trigger inflammatory processes, and affect hemostasis. Many crucial vasoactive endogenous compounds

are produced by the endothelial cells to control the functions of vascular smooth muscle cells and of circulating blood cells [132]. These complex systems determine a fine equilibrium which regulates the vascular tone. Impairments in endothelium-dependent vasodilation lead to the so-called endothelial dysfunction. The endothelium modulates several biological processes through the balance between opposing mechanisms such as vasoconstriction/vasodilation, pro-coagulation/antithrombosis, cell proliferation/apoptosis, among others. This dynamic tissue layer constitutes a source and/or target of multiple growth factors and vasoactive mediators involved in regulating the physical and biochemical properties of the systemic vessels, as well as vascular contractility and cell growth. Concerning the regulation of vasomotor tone, the endothelium plays a protective role by generating vasorelaxing substances, such as NO, PGI₂, and endothelium-derived hyperpolarizing factor (EDHF) [133], which oppose the vasoconstricting action. Vascular smooth muscle cells (VSMCs) are essential for good performance of the vasculature. They can alter the luminal diameter which enables blood vessels to maintain an appropriate blood pressure. VSMC are fit not only for short-term regulation of the vessel diameter, but also for long-term adaptation, via structural remodeling by changing cell number and connective tissue composition. The main function of vascular smooth muscle tonus is to regulate the caliber of the blood vessels in the body. Excessive vasoconstriction leads to hypertension, having physical effects on the vascular wall, such as tensile stress caused by pressure and shear stress caused by flow [134].

The vasoconstricting action of hormones, such as endothelin-1, angiotensin II, or urotensin II, among others, is mediated by ROS. Furthermore, ROS mediate many pathophysiological processes, such as growth, migration, apoptosis and secretion of inflammatory cytokines, as well as physiological processes, such as differentiation, by direct and indirect effects at multiple signaling levels. It was reported that H₂O₂ activates phospholipase C on VSMC through tyrosine phosphorylation and that this activation has a major role in rapid [Ca²⁺]_i mobilization in this type of cells [135]. Ultimately, the physiological or pathophysiological effect of ROSs is dependent on their steady state concentration. Oxidative stress can lead to unbalanced vasoconstricting and vasodilating forces in favor of the first. Indeed, this impairment in vasomotor tone regulation belongs to a more complex phenomenon named endothelial dysfunction, characterized by a shift of the actions of the endothelium toward reduced vasodilation, a proinflammatory state, and prothrombotic setting [6].

A variety of enzymatic and nonenzymatic sources of ROS exist in blood vessels. Enzymatic sources of ROS include NADPH oxidases located on the cell membrane of polymorphonuclear cells, macrophages, and endothelial cells and cytochrome P450-dependent oxygenases. The proteolytic conversion of xanthine dehydrogenase to xanthine oxidase provides another enzymatic source of both superoxide and H₂O₂ (and therefore constitutes a source of the highly reactive hydroxyl radicals) and has been proposed to mediate deleterious processes in vivo. In addition to NADPH

oxidase, the best characterized source of ROS, several other enzymes may contribute to ROS generation, including NO synthase, lipoxygenase, cyclooxygenases, xanthine oxidase, and cytochrome P450 enzymes. Also mitochondria could be considered a major source of ROS: in situations of metabolic perturbation, increased mitochondrial ROS generation might trigger endothelial dysfunction, possibly contributing to the development of hypertension [136].

4.2. Relationship between Oxidative Stress and Essential Hypertension. Systolic blood pressure and diastolic blood pressure correlated inversely with plasma antioxidant potential (e.g., FRAP) and directly with some oxidative stress-related parameters (e.g., F2-isoprostanes) [137]. On the other hand, other plasma blood pressure modulator levels, such as AT-II, ET-1, renin, aldosterone, and homocysteine, among others, showed no significant differences between normotensive and hypertensive subjects. These associations strongly suggest a role for oxidative stress as a modulator of blood pressure. These results are in agreement with previous studies reporting increased oxidative stress and decreased antioxidant capacity in patients with essential hypertension compared with normotensive subjects [138, 139]. These findings also suggest consideration of oxidative stress as a novel therapeutic target for the therapy of essential hypertension. F2-isoprostanes have been proposed as a new factor involved in the modulation of BP [140]. F2-isoprostanes have a direct correlation with blood pressure levels in normotensive and hypertensive subjects [137].

Long-term hypertension produces irreversible endothelial dysfunction due in part to self-sustaining ROS production [5, 141, 142]. In patients with already established endothelial damage, measurement of oxidative stress-related biomarkers will be of less use. In order to establish a functional interpretation of hypertension through oxidative stress-related biomarkers, the early phases of the disease should be studied. In the endothelial dysfunction phase, previous to the development of hypertension, the ROS fluctuations will appropriately reflect the balance between ROS production and their scavenging or depuration by the antioxidant defense system in the endothelium.

The role of the antioxidants vitamins C and E has emerged as a possible therapy for decreasing oxidative stress, thereby lowering BP. In fact, a concomitant decrease in BP and oxidative stress was shown following Vitamins C and E supplementation in patients with essential hypertension [143].

In contrast with these findings, the results of short-term trials of antioxidant supplements on blood pressure have been inconsistent [144, 145] and long-term clinical trials have failed to consistently support the antihypertensive effects of vitamins C and E in patients at high cardiovascular risk. This controversy may be explained by the patient inclusion and exclusion criteria, which did not consistently consider the influence of factors such as already established target organ damage, smoking status, physical activity, and intake of nonantihypertensive (e.g., statins) medication, which are potential confounders of oxidative stress status.

4.3. Antioxidant Therapy Effectiveness May Be Closely Dependent on Clinical Setting and Oxidative Status. Antioxidant therapy should be rather effective in the early stages of hypertension or atherosclerosis by preventing the oxidative-stress mediated “positive feedback loop” of progression from reversible endothelial dysfunction to atherosclerotic plaque formation [146]. Most patients participating in clinical trials are in the advanced stages of this progression (i.e., with atherosclerosis or cardiovascular disease) and the “positive feedback loop” will have generated irreversible structural changes such as vascular remodeling or atherosclerosis. Therefore the lack of benefit of antioxidant therapy in diminishing CV events may reflect incomplete information on the actual role of oxidative stress, which should be considered as a necessary component of the enrolled patients, as antioxidants may have no effect on the absence of marked oxidative stress and already established cardiovascular disease.

In all animal studies with positive results, the vitamin E administration started before the constitution of histological atherosclerotic changes in the structure of the arterial wall [147]. Furthermore, the protective effect shown by vitamins C and E supplementation in the prevention of immune atherosclerotic lesions occurring after cardiac transplant also supports this assumption. [148, 149]. Correction of hypertension or prehypertension and other cardiovascular risk factors leading to endothelial dysfunction in their early stages may help reduce the burden of disease and mortality attributable to high blood pressure [150, 151].

Considerable evidence suggests that oxidative stress, which results in an excessive generation of ROS, plays a key role in the pathogenesis of hypertension. Antioxidant therapies have been evaluated in order to decrease ROSs production or increase their scavenging. In this line, polyphenols, widespread antioxidants in fruits, vegetables, and wine, have demonstrated their beneficial role in prevention and therapy of hypertension, by acting as free-radical scavengers, metal chelators, and in enzyme modulation and expression. Numerous studies have demonstrated the importance of naturally occurring dietary polyphenols in promoting cardiovascular health and emphasized the significant role that these compounds play in limiting the effects of cellular aging. Polyphenols such as resveratrol, epigallocatechin gallate (EGCG), and curcumin have been acknowledged for having beneficial effects on cardiovascular health, while some have also been shown to be protective in aging [152]. Polyphenols activate and enhance eNOS expression by several signaling pathways, increase glutathione levels, and inhibit ROS-producing enzymes such as NADPH and xanthine oxidases. Accordingly, antioxidant activity of phenolics in leaves of three red pepper (*Capsicum annuum*), mainly putrescine and anthocyanine derivatives, demonstrated a great potential for radical scavenging activity against oxidative stress [153]. In addition, some polyphenols can exert beneficial effects other than those related to their antioxidant properties. Thus, luteolin can reduce VSMCs' proliferation and migration [154], and effects also reported for resveratrol [155].

Also, polyphenols have the capacity to improve the endothelial control of vascular tone not only in several experimental models of cardiovascular diseases such as

hypertension but also in healthy and diseased humans. Experiments with isolated arteries have shown that polyphenols cause nitric oxide (NO) mediated endothelium-dependent relaxations and increase the endothelial formation of NO [156], via the PI3-kinase/Akt pathway, leading to a rapid activation of endothelial NO synthase through its phosphorylation at Ser1177 [157].

It would then be concluded that the time window for an antioxidant therapy in essential hypertension is narrow, and as such should be focused on new onset and high-risk hypertension individuals having biomarkers consistent with a high ROS load.

As an example, in the Framingham Study, subjects who had isolated diastolic hypertension at baseline were generally male overweight or obese smokers and were at inordinately higher risk (23-fold) than those with optimal blood pressure to develop systo-diastolic hypertension [151]. As mentioned before, patients with these characteristics have shown favorable response to antihypertensive antioxidant therapy [139].

4.4. Cellular Biomarkers of Endothelial Health: EPCs and EMP. Endothelial progenitor cells (EPCs) are novel biomarkers for cardiovascular risk and show considerable potential as they reflect the body's capacity for endothelial repair [109]. EPCs have been validated as CV risk biomarkers for both low- and high-risk population [158–160]. Regarding essential hypertension, EPCs count in hypertension is decreased as is their overall function. Hypertension is associated with some specific EPCs function alterations, such as reduced in vitro migratory capacity [161] and increased senescence [162].

EPCs usefulness as a CV risk biomarker shows great promise but a consensus characterization of EPCs is still lacking [112], and further studies are needed to determine the usefulness of measuring EPC absolute number or functional capacity in different clinical settings [109]. Antioxidant agents, such as vitamin C, lower circulating EMP levels in patients with diabetes and dyslipidemia after myocardial infarction [110].

In turn, endothelial microparticles (EMPs) are considered as potential CV biomarkers, which stems from the fact that they directly reflect endothelial damage [109]. Microparticles circulate in healthy individuals and their levels increase in cardiovascular and atherothrombotic diseases [163]. In subjects with high risk of coronary heart disease, baseline levels of EMP predicted outcome, independent of Framingham score and of C-reactive protein (CRP) and brain natriuretic peptide levels [110].

In patients with hypertension, plasma levels of EMP correlated with systolic and diastolic blood pressures [111].

5. Ischemia-Reperfusion Myocardial Damage

The main clinical model of ischemia-reperfusion injury is acute myocardial infarction (AMI), the leading cause of mortality worldwide. According to the WHO, in 2008 ischemic heart disease accounted for 7.25 million deaths (12.8%). Systemic thrombolysis and percutaneous coronary angioplasty

(PCA) have been used to recover the myocardial perfusion, with the latter being the most successful, as it allows to re-establish the blood flow in the cardiac zones affected by the occlusion of a branch of the coronary artery. Nevertheless, as a consequence of this procedure, the ischemic zone is reperfused, giving rise to an ischemia-reperfusion event that generates increased production of ROS (oxidative stress) [164]. Therefore, this widely performed procedure leads to an enhancement of the previously established tissue damage (lethal reperfusion), as these reactive species attack biomolecules such as lipids, DNA, and proteins and trigger cell death pathways [165]. As ischemia-reperfusion injury remains a neglected therapeutic target [166], biomarkers of ischemia reperfusion could be used to measure the effectiveness of novel cardioprotective strategies to prevent lethal reperfusion.

A series of biochemical and metabolic changes in myocardial tissue occur due to deprivation of oxygen and nutrient supply during ischemia. Consequently, mitochondrial damage and ATP depletion impair myocardial contractile function [166]. Anaerobic glycolysis due to the absence of oxygen results in the accumulation of lactate and intracellular pH reduction (to <7.0). The latter activates the Na^+ - H^+ ion exchanger, thus extruding protons from the cell in exchange for Na^+ entry. Furthermore, the impaired function of ($\text{Na} + \text{K}$)-ATPase contributes to exacerbating the intracellular Na^+ and Ca^{2+} overload [167]. At the onset of reperfusion, the level of tissue oxygenation increases, leading to a burst of ROS generation, thereby accounting for the syndrome of reperfusion injury [164]. Neutrophils are the primary source of ROS during reperfusion, although endothelial cells and cardiomyocytes can also generate these reactive species. Increased ROS production is mainly due to activation of XO in endothelial cells, mitochondrial electron transport chain reactions in cardiomyocytes, and NADPH oxidase in inflammatory cells [168].

Several biomarkers could be used to demonstrate the major role of oxidative stress in the pathophysiology of ischemia-reperfusion damage in myocardial infarction. Extracellular biomarkers of ischemia-reperfusion damage encompass products of lipid peroxidation, plasma antioxidant vitamin levels, total antioxidant capacity of plasma, protein carbonylation, and uric acid level. In addition, intracellular biomarkers such as activity of antioxidant enzymes, thiol index (GSH/GSSG ratio), carbonyl levels, and F2-isoprostane, among others, are suitable to estimate oxidative stress injury.

5.1. Lipid Peroxidation. Increased lipid peroxidation is thought to be a consequence of oxidative stress. It occurs due to an impairment altering the dynamic balance between prooxidant and antioxidant mechanisms. In ischemia, the ATP is drastically reduced and is converted to hypoxanthine and then to uric acid by xanthine oxidase upon reperfusion. During this process, enormous amounts of superoxide radicals are formed and can simulate Haber-Weiss reaction for further generation of ROS, thus initiating lipid peroxidation [169]. It has been shown that in AMI

patients, the increase of oxidative stress markers, such as lipid peroxidation products, mirrors the decrease in low-density lipoprotein-cholesterol [170]. A reason for increased lipid peroxidation in plasma of AMI patients may be a poor enzymatic and nonenzymatic antioxidant defense system. Pasupathi et al. in a cohort of 100 patients with evidence of AMI, showed a significant rise in TBARS levels ($P < 0.001$), a lipid peroxidation biomarker indicative of elevated oxidative stress [171]. Furthermore, these authors showed an increased concentration of MDA in the circulation of total AMI patients, indicating increased lipid peroxidation. These results are in agreement with previous reports [172]. Nevertheless, lipid peroxidation might be increased due to the presence of systemic diseases including hyperlipidemia, diabetes mellitus, or renal disease, in addition to myocardial infarction [173], decreasing its specificity in this type of patients.

Also, F2-isoprostane, a representative isoprostane, has been reported to be a reliable marker for oxidant stress *in vivo*. In addition, it was reported that the level of immunoreactive 8-iso PGF(2 alpha) is increased in the great cardiac vein in patients with AMI [174]. The study showed transcatheter F2-isoprostane generation, being a reliable biomarker for the size of myocardium exposed to oxidant stress.

5.2. Plasma Antioxidant Vitamin Levels. In AMI patients, Pasupathi et al. showed lower levels of vitamins E and C compared with controls [171]. This is in accordance with studies of Singh et al. [175], who demonstrated that there was a significant drop in vitamins C, E, A, and β -carotene, whereas lipid peroxides were significantly higher in AMI patients, compared with controls. This indicates severe damage to antioxidant system, which is unable to combat oxidative stress and inflammation [176].

5.3. Total Antioxidant Capacity (TAC). This biomarker has been described as a critical tool for assessing redox status [177]. The TAC or related antioxidants may play an important role in protecting the organism from free-radicals-mediated damage [178]. The role that such compounds play in AMI development is important, since their presence may decrease the damage resulting from blood ROS during reperfusion.

5.4. Antioxidant Enzymes. Antioxidants constitute the foremost defense system that limit the toxicity associated with free radicals. It is known that plasma antioxidant capacity decreases and oxidative/antioxidative balance shift to the oxidative balance in patients with MI. SOD along with CAT and GPx, the preventive antioxidants, play a very important role in protection against lipid peroxidation [179]. Scott et al. showed that SOD, CAT, and GPx activities were significantly lower in MI and ischemic heart disease patients than in control subjects. Besides, decrease of SOD, CAT, and GPx activity was much more pronounced in smokers than in nonsmokers with AMI, thus making those individuals more vulnerable to oxidative stress. Free radical-scavenging enzymes such as SOD, CAT, and GPx are the first line of cellular defense against oxidative injury, decomposing $O_2^{\bullet-}$

and H_2O_2 , otherwise interacting to form the more reactive hydroxyl radical. These enzymes protect the red cells against $O_2^{\bullet-}$ and H_2O_2 -mediated lipid peroxidation [179]. Decreased activities of SOD and CAT in the erythrocyte in AMI patients have been reported [180, 181], likely due to inactivation by cross-linking or to exhaustion of the enzymes by increased peroxidation [182].

Glutathione peroxidase (GPx) catalyzes peroxide reduction utilizing GSH as the substrate and converting it to GSSG. Its activity has been reported decreased in erythrocytes of AMI patients [171]. Also, inactivation of GPx after endogenous exposure to products of lipid peroxidation [183] or of NO [184] has been reported. GSH is one of the most important endogenous intracellular antioxidants. It plays the role of a sulfhydryl (SH) group provider for direct scavenging reactions. GSH acts both as a substrate in the scavenging reaction catalyzed by GPx and as a scavenger of vitamins C and E radicals. A lowered serum concentration of reduced glutathione (GSH) has been considered the predictor of coronary restenosis after percutaneous coronary intervention, and its deficiency could lead to pronounced postreperfusion syndrome [185]. Decrease GSH concentration may also contribute to decreased GPx activity, because GSH is one of its substrates [186]. In accordance with this it has been shown that plasma and erythrocyte GSH concentration were significantly decreased in AMI patients. It may be due to an increased utilization of GSH [180]. In a single center prospective cohort trial with a 15-month follow-up [187], the primary endpoint was major adverse cardiac events (MACE), defined either as re-PCI or coronary artery bypass grafting, MI, cardiac death, or cardiac rehospitalization, or a combination thereof. Results showed that GSH levels are significantly decreased in the whole MACE group. Authors acknowledge that this was a preliminary study with a relatively low number of subjects. Due to this limitation, they observed a relatively low number of MACEs during follow-up. Also they used preprocedural GSH concentrations as a discriminative variable, and we did not monitor the fluctuations in the GSH concentration over time, to use it as a harmonic (time-dependent) variable.

High plasma MPO is reported to be a risk factor for early adverse cardiac events in patients with chest pain [188] or acute coronary syndromes [189] and to be associated with endothelial dysfunction [190, 191]. In myocardial infarction a retrospective study concluded that levels of MPO were independently predictive of 5-year survival and might provide additional prognostic information if used in combination with the established markers, left ventricular ejection fraction and NT-pro-BNP [192].

Ischemia modified albumin (IMA) is a recently developed biomarker of transient myocardial ischemia [193]. Circulating IMA is increased in patients with myocardial ischemia, after percutaneous coronary intervention [194] or in acute coronary syndromes [195]. The test has recently been licensed by the US Food and Drug Administration for diagnostic use in suspected myocardial ischemia. IMA is serum albumin in which the N-terminus has been chemically modified. It has been proposed that ROSSs such as superoxide ($O_2^{\bullet-}$)

and hydroxyl ($\cdot\text{OH}$) radicals generated during myocardial ischemia-reperfusion modify the N-terminus of serum albumin resulting in IMA formation. In vivo modifications of the albumin N-terminus are proposed to be related to the ROS production during myocardial ischemia reperfusion. For example, an increase in IMA was observed in patients minutes after transient occlusion and reperfusion during coronary angioplasty [195]. Roy et al. in 2006 showed for the first time that IMA formation is directly related to ROS in vitro, generating and providing novel information on the nature of the species contributing to IMA formation, namely $\cdot\text{OH}$ [196].

Polyphenols induce cardiovascular protective effects, thanks to their antioxidant, vasodilator, anti-inflammatory, anti-fibrotic, antiapoptotic and metabolic properties. Polyphenol-dependent cardioprotection occurs mainly through NO and via several powerful prosurvival cellular cascades that involve metabolic intermediates [197]. It has been reported that the polyphenol luteolin protects against myocardial injury and enhances cellular antioxidant defense capacity through the activation of Akt and ERK signal pathways that leads to Nrf2 activation and subsequently heme oxygenase-1 induction [198]. In addition, it preserves cardiac function, reduces infarct size and cardiomyocyte apoptotic rate after ischemia/reperfusion injury in diabetic rats [199]. These authors also demonstrated that luteolin has an anti-apoptotic role in an experimental model of ischemia/reperfusion in rats [199], being a potential candidate for preventing and treating cardiovascular diseases [200].

Nevertheless, common oxidative stress-related biomarkers for both pathologies could be the diminution of TAC of plasma and increased lipid peroxidation products, such as F2-isoprostanes and MDA in percutaneous coronary angioplasty following coronary infarction [139, 201].

6. Concluding Remarks and Perspectives

Increasing body of evidence accounts for the validity of using biomarkers to assess the factors involved in pathogenesis and clinical outcome of human diseases. Several biomarkers could be used to demonstrate the major role of oxidative stress in cardiovascular pathologies, such as those derived from essential hypertension and ischemia reperfusion damage. Indeed, the applicability of oxidative stress-related biomarkers arises from the role of ROS in the pathophysiology of these clinical settings. The role of ROS in the development of essential hypertension arises from both decreased vasodilation due to a diminished bioavailability of NO and prostacyclin together with an increased vasoconstriction by their cellular effect on VSMC, thus accounting for blood pressure elevation. The contribution of ROS to myocardial oxidative damage caused by infarction is due to direct and cellular mediated effects of ROS, such as necrosis and apoptosis, respectively.

Extracellular oxidative stress-related biomarkers include total antioxidant capacity of plasma, plasma antioxidant vitamin levels, lipid peroxidation product levels, protein carbonylation, and uric acid level. Intracellular oxidative

stress related biomarkers include the activity of antioxidant enzymes, prooxidant enzymes, and thiol index.

Cardiac biomarker testing in the clinical laboratory continues to evolve based on the availability of improved technologies and the results of clinical trials among patients with acute coronary syndromes. Although still at its infancy, the use of cellular biomarkers of endothelial dysfunction has the potential to yield powerful mechanistic insight into endothelial dysfunction. They provide a direct window to the active state of the endothelium or to endothelial repair. There is ample consensus that these biomarkers provide a direct window to the active state of the endothelium or to endothelial repair and for these reasons, they represent special value as biomarkers in these clinical settings. In the same fashion, ischemia modified albumin value lies in how it specifically denotes the presence of ischemia in the organism.

In summary, the applicability of oxidative stress-related biomarkers to study the pathogenesis and clinical outcome of essential hypertension and ischemia reperfusion cardiac pathologies arises from the role of ROS in cardiovascular diseases. Lipid peroxidation biomarkers such as plasma F2-isoprostane levels have been widely studied in the assessment of in vivo oxidative stress-related human disease. However, novel biomarkers are emerging to account for additional pathological processes. Thus, increased inflammatory and oxidative stress biomarkers are associated with endothelial progenitor cell dysfunction, which is reversed upon treatment with anti-inflammatory or antioxidant drugs. Therefore, inflammatory and redox signaling pathways that interfere with EPC bioactivity need to be further investigated.

Abbreviations

AMI:	Acute myocardial infarction
ARE:	Antioxidant response elements
BP:	Blood pressure
ACB:	Albumin cobalt binding
CAT:	Catalase
CABG:	Coronary artery bypass grafting
CO:	Carbon monoxide
CV:	Cardiovascular
CRP:	C-reactive protein
DNPH:	2,4-dinitrophenylhydrazine
DNP:	Stable dinitrophenyl
ELISA:	Enzyme-linked immunosorbent assay
EDHF:	Endothelium-derived hyperpolarizing factor
FRAP:	Ferric reducing antioxidant power
EMP:	Endothelial microparticles
EPCs:	Endothelial progenitor cells
FeIII-TPTZ:	Ferric-tripyridyltriazine
GSH:	Reduced glutathione
GSH-Px:	Glutathione peroxidase
H ₂ O ₂ :	Hydrogen peroxide
OH:	Hydroxyl
4-HNE:	4-hydroxynonenal
IMA:	Ischemia modified albumin
MACE:	Major adverse cardiac events
MPO:	Myeloperoxidase
MDA:	malondialdehyde

MI:	Myocardial infarction
NAD ⁺ :	Nicotinamide adenine dinucleotide
NADH:	Nicotinamide adenine dinucleotide dehydrogenase
NADPH:	Nicotinamide adenine dinucleotide phosphate
NO:	Nitric oxide
NOS:	Nitric oxide synthases
Nrf2:	Nuclear factor erythroid 2-related factor-2
nNOS:	A constitutive neuronal NOS
iNOS:	Cytokine-inducible NOS
eNOS:	Constitutive endothelial NOS
O ₂ ^{•-} :	Superoxide anion
[•] OH:	Hydroxyl radical
PCI:	Percutaneous coronary intervention
PUFA:	Polyunsaturated fatty acids
ROS:	Reactive oxygen species.
RNS:	Reactive nitrogen species
SOD:	Superoxide dismutase
SH:	Sulfhydryl
TBARS:	Thiobarbituric reactive species
TNF:	Tumor necrosis factor
VSMC:	Vascular smooth muscle cells
XO:	Xanthine oxidase
XDH:	Xanthine dehydrogenase.

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