Diabetic cardiomyopathy and its mechanisms: Role of oxidative stress and damage

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ABSTRACT
Diabetic cardiomyopathy as an important threat to health occurs with or without coexistence of vascular diseases. The exact mechanisms underlying the disease remain incompletely clear. Although several pathological mechanisms responsible for diabetic cardiomyopathy have been proposed, oxidative stress is widely considered as one of the major causes for the pathogenesis of the disease. Hyperglycemia, hyperlipidemia, hypertension- and inflammation-induced oxidative stress are major risk factors for the development of microvascular pathogenesis in the diabetic myocardium, which results in abnormal gene expression, altered signal transduction and the activation of pathways leading to programmed myocardial cell deaths. In the present article, we aim to provide an extensive review of the role of oxidative stress and anti-oxidants in diabetic cardiomyopathy based on our own works and literature information available.

INTRODUCTION
The occurrence of systemic hyperglycemia in diabetes can ultimately cause long-term damage to multiple organs, leading to severe complications. The microvasculature is a key target of hyperglycemic damage, and damage to small blood vessels can initiate and/or exacerbate systemic complications. Diabetic cardiovascular injury includes both cardiac and peripheral blood vessels. However, research has focused largely on peripheral vascular injury. Retinopathy, neuropathy and nephropathy are all caused, in part, by changes in blood vessels. However, research has focused largely on peripheral vascular injury. Retinopathy, neuropathy and nephropathy are all caused, in part, by changes in blood flow or abnormal vessel growth1–3. A disorder of the heart muscle in patients with diabetes is called diabetic cardiomyopathy (DCM), which can lead to inability of the heart to circulate blood through the body effectively, a state known as heart failure4. DCM is only said to exist if there is no coronary artery disease to explain the heart muscle disorder, although most heart failure in patients with diabetes results from coronary artery disease. Although recent studies have extended our understanding of the pathogenesis of the DCM, the mechanism by which hyperglycemia causes DCM remains incompletely understood1–3.

Oxidative stress has been linked both to the onset of diabetes and its complications5–9. Recent studies have shown that oxidative damage induced by reactive oxygen or nitrogen species (ROS and/or RNS) derived from hyperglycemia plays a critical role in diabetic injury in multiple organs. Thus, the relationship between oxidative stress and DCM is a major focus of current research. Although these pathogenic factors probably cause DCM through different mechanisms10–15, the major contribution of these pathogenic factors to DCM is oxidative stress16. Therefore, anti-oxidant prevention and therapy is critically important to prevent or postpone the development of diabetic cardiovascular complications. Here, we review the features of DCM in term of its physiological and pathogenic bases, the critical role of oxidative stresses in the pathogenesis of DCM and potential anti-oxidant therapy for DCM, with a focus on targets of the nuclear factor, erythroid-2-related factor 2 (Nrf2), and metallothionein (MT) based on the literature and our own studies.

PATHOPHYSIOLOGICAL BASE OF DCM
Emerging evidence from experimental, pathological, epidemiological and clinical studies has shown that diabetes mellitus causes cardiac functional and structural changes, independent of hypertension, coronary artery disease or any other known cardiac diseases, which supports the existence of DCM17–19. DCM is characterized by a disproportionate increase in left ventricular (LV) mass and myocardial fibrosis, and, as an early complication of diabetes, is manifested by diastolic dysfunction followed by abnormalities in systolic function20. One particularity of DCM is the long latent phase, during which the disease progresses, but is completely asymptomatic. A prominent ‘a’ wave can also be noted in the jugular venous pulse, and the
cardiac apical impulse might be overactive or sustained throughout systole. After the development of systolic dysfunction, LV dilation and symptomatic heart failure, the jugular venous pressure might become elevated, and the apical impulse would be displaced downward and to the left. These changes are accompanied by a variety of electrocardiographic changes that could be associated with DCM in 60% of patients without structural heart disease, although usually not in the early asymptomatic phase. Later in the progression, a prolonged QT interval might be indicative of fibrosis.

In most cases, DCM is detected with concomitant hypertension or coronary artery disease. When presenting with other cardiovascular complications (i.e., ischemic heart disease or hypertension), diabetic patients have a much worse prognosis than non-diabetic patients, and are more prone to progress to congestive heart failure. Diastolic dysfunction characterized by increased ventricular wall stiffness, and increased diastolic relaxation time is prevalent at early stages of the cardiomyopathy. Cardiomyocyte death is paralleled by fibroblast replacement, and leads to interstitial fibrosis mediated primarily by transforming growth factor-β (TGF-β), whereas cell death is considered to be a result of the induction of cardiac oxidative stress, although several other mechanisms have been also associated with DCM.

**GENERAL MECHANISMS UNDERLYING DCM**

Extensive evidence from experimental models of type 1 and type 2 diabetes implicates hyperglycemia as one of the main pathogenic mechanisms in DCM. An important consequence of high glucose-induced cellular injury is the formation of advanced glycation end-products (AGEs) resulting from the non-enzymatic glycation and oxidation of proteins and lipids. Recently, high levels of AGE products have been found in cardiac tissue of diabetic patients. Activation of the protein kinase C/diacylglycerol (PKC/DAG) signaling pathway is one of the mechanisms by which hyperglycemia exerts adverse cardiovascular effects. PKC activation by hyperglycemia contributes to cardiac fibrosis by stimulating connective tissue growth factor (CTGF) expression as shown in transgenic PKC-β2 mice. Consistently, the overexpression of PKC-β2 isoform in the heart of transgenic mice resulted in LV hypertrophy, fibrosis and decreased LV ejection fraction (LV EF), similar to that of DCM. PKC inhibition attenuated diastolic dysfunction, myocyte hypertrophy and collagen deposition, and preserved cardiac contractility in a rodent model of diabetic diastolic heart failure, suggesting that PKC-inhibition could represent a novel therapeutic strategy for the prevention of diabetes-associated cardiac dysfunction.

Intracellular Ca\(^{2+}\) is a major regulator of cardiac contractility. Diabetes is associated with the impairment in myocardial performance as a result of abnormalities in contractile and regulatory protein expression, and in cardiomyocyte Ca\(^{2+}\) sensitivity. In a study of human cardiomyocytes from type 2 diabetes patients, Ca\(^{2+}\) dysregulation was indicated by reduced myofilament Ca\(^{2+}\) sensitivity. Diabetes is associated with the activation of the renin–angiotensin system with consequent overproduction of angiotensin II (Ang II), its main physiological effector molecule. Ang II contributes to several alterations in diabetes including fibrosis by stimulating extracellular matrix component synthesis, apoptosis/proliferation, vascular inflammation and oxidative damage. The oxidative damage includes deoxyribonucleic acid (DNA) damage, which could be related to activation of poly-(ADP-ribose) polymerase (PARP) enzymes.

The endoplasmic reticulum (ER) is an organelle entrusted with lipid synthesis, calcium homeostasis, protein folding and maturation. Perturbation of ER-associated functions results in an evolutionarily conserved cell stress response, the unfolded protein response that is also called ER stress. ER stress is aimed initially at compensating for damage, but can eventually trigger cell death if ER stress is excessive or prolonged. Now ER stress has been associated with numerous diseases. For instance, our recent studies have shown the important role of ER stress in diabetes-induced cardiac cell death. It is known that apoptosis has been considered to play a critical role in diabetic cardiomyopathy. Inhibition of ER stress can result in the prevention of DCM.

Apoptosis has been considered as the outcome of cardiomyocytes in response to diabetic hyperglycemia, hyperlipidemia, inflammation and ER stress, and is also the key initiating factor to stimulating resident cardiomyocyte hypertrophy and fibroblast proliferation, both leading to cardiac remodeling and eventually dysfunction. In fact, autophagy, which is a process of self-cannibalization used by mammalian cells for the recycling of cellular contents, alters the heart of diabetic individuals. Like ER stress, autophagy can be both adaptive and maladaptive under specific settings. In OVE26 type 1 diabetes mice, the adenosine monophosphate (AMP)-activated protein kinase (AMPK) is reduced along with cardiac dysfunction and decreased cardiac autophagy. Genetic inhibition of AMPK in cardiomyocytes attenuates cardiac autophagy, exacerbates cardiac dysfunction and increases mortality in diabetic mice. More importantly, chronic AMPK activation with metformin, one of the most used antidiabetes drugs and a well-characterized AMPK activator, significantly enhances autophagic activity, preserves cardiac function and prevents most of the primary characteristics of DCM in OVE26 mice, but not in dominant negative-AMPK diabetic mice. Therefore, AMPK activation protects cardiac structure and function by increasing cardiac autophagy in the diabetic heart, whereas ceramide synthase-mediated lipid-induced autophagy and hypertrophy in cardiomyocytes is a cause of DCM. Therefore, an urgent need exists to clarify the mechanism of pathogenesis of DCM. Emerging evidence shows that diabetes induces cardiomyocyte apoptosis and suppresses or stimulates cardiac autophagy, indicating that the interplay between autophagy and apoptotic cell death pathways is important in the pathogenesis of DCM.
OXIDATIVE STRESS

Although several potential mechanisms underlying DCM have been addressed, all these pathogenic factors either induce or result from oxidative stress. Oxidative stress occurs when the production of ROS and/or RNS exceeds their degradation by anti-oxidant defenses. PARP proteins are overexpressed in diabetes\textsuperscript{46,47} as ROS- and/or RNS-induced oxidative DNA damage. Hyperglycemia in both type 1 diabetes and type 2 diabetes\textsuperscript{48,49} is associated with increased ROS and RNS production in mitochondria, as shown in Figure 1.

Mitochondria are a major source of ROS production. Cellular sources of ROS generation within the heart include cardiac

Figure 1 | Outline of the possible physiological roles of redox signaling in maintaining the balance between various reactive oxygen species and/or reactive nitrogen species, and anti-oxidants under (a) normal and (b) diabetic conditions. AGE, advanced glycation end-products; Ang II, angiotensin II; ASK1, apoptosis signal-regulating kinase 1; AT1, angiotensin II-1 receptor; eNOS, endothelial nitric oxide synthase; GPx, glutathione peroxidase; GSH, glutathione; GSSG, glutathione disulfide; H$_2$O$_2$, hydrogen peroxide; HB, tetrahydrobiopterin; iNOS, inducible nitric oxide synthase; IR, insulin receptor; IRS1, insulin receptor substrate-1; IRS2, insulin receptor substrate-2; JNK, c-Jun NH2-terminal kinase; NF-$\kappa$B, nuclear factor-$\kappa$B; NO, nitric oxide; NOX, nicotinamide adenine dinucleotide phosphate oxidase; ONOO$^-$, peroxynitrite; PI3K/AKT, phosphoinositide 3-kinase/Akt; PKC, protein kinase C; PTP-1B, protein tyrosine phosphatase -1B; SOD, super oxide dismutase; TR, thioredoxin reductase; Trx-1, thioredoxin-1.
myocytes, endothelial cells and neutrophils. The electron transport chain generates superoxide radicals as an inevitable byproduct at complex I and complex III in the respiratory chain. ROS leads to cellular damage through several mechanisms (oxidation, interference with nitric oxide [NO] and modulation of detrimental intracellular signaling pathways). Therefore, increased ROS causes cardiac dysfunction by direct damage to proteins and DNA, and by inducing apoptosis.

Elevated free fatty acid (FFA) concentrations in the heart as a result of a lack of insulin-mediated glucose metabolism can increase ROS and/or RNS production from both non-mitochondrial sources, such as activation of nicotinamide adenine dinucleotide phosphate (NADPH) oxidases (NOX), and modulating the mitochondrial electron chain to generate superoxide. Activated NOX produces superoxide which, in turn, can combine with NO, forming highly reactive and damaging peroxynitrite species. NOXs have emerged as a culprit in this condition. Cellular sources of ROS and/or RNS generation within the heart include cardiac myocytes, endothelial cells and neutrophils. ROS lead to cellular damage through several mechanisms (oxidation, interference with NO and modulation of detrimental intracellular signaling pathways). Therefore, increased ROS and/or RNS causes cardiac dysfunction by direct damage to proteins and DNA, as well as by inducing apoptosis. Oxidative damage and/or loss of anti-oxidant defense could also stimulate a number of responses associated with ventricular remodeling. These include activation of matrix metalloproteinase to alter the architecture of the extracellular matrix, modulation of signal transduction pathways that initiate cardiomyocyte hypertrophy and apoptosis, as summarized in Figure 1.

Some experiments suggest that cytosolic ROS and/or RNS defense mechanisms might be more significantly impacted by diabetes than mitochondrial defense mechanisms. These include activation of matrix metalloproteinase to alter the architecture of the extracellular matrix, modulation of signal transduction pathways that initiate cardiomyocyte hypertrophy and apoptosis.

**ANTI-OXIDANTS IN DCM**

Given that oxidative and/or nitrosative stress seems a critical cause for the development of DCM, several anti-oxidant approaches have been investigated for their potential prevention and amelioration of DCM. Although anti-oxidant therapy has been extensively studied for its potential prevention or treatment of various diabetic complications, direct investigations for the prevention by anti-oxidants of DCM remain relatively scarce. Here, we provide some examples first, and then will focus on Nrf2 and MT as the major targets of anti-oxidative therapy for DCM based on our own and other works.

**Thioredoxin 1**

In recent studies, intramyocardial thioredoxin 1 (Trx1) gene therapy in diabetic rats showed the prevention of angiogenic impairment and cardiac dysfunction severity after cardiac infarction. They found that overexpression of the Trx1 gene could reduce oxidative stress and apoptosis and, thereby, the extent of ventricular remodeling.

**Tempol**

Super oxide dismutase (SOD) has been found to protect cardiomyocytes from superoxide-induced damage by converting superoxide to hydrogen peroxide. Tempol, 4-hydroxy-2,6,6-tetramethyl piperidinoxyl, is a membrane-permeable SOD mimetic that has been shown to attenuate the effects of peroxynitrite in vascular systems. In the diabetic heart, tempol was also found to significantly reduce diabetes-activated GSK-3β, associated with increases in caspase-3 cleavage and apoptotic cell death at 3 days after streptozotocin (STZ) injection. The anti-apoptotic effect of tempol was accompanied with attenuation of STZ-induced cardiac hypertrophy and fibrosis, observed at 28 days after STZ injection. Prevention of diabetes-induced cardiac fibrosis was observed at 8 weeks in diabetes patients after 8-week tempol treatment, along with a significant reduction of oxidative stress as evidenced by reducing the expression of p22 phox, ROS production and increasing anti-oxidant enzyme capacity.

**Curcumin**

Curcumin is the natural polyphenolic compound that was originally used in traditional Indian medicine over 3,000 years ago, and has shown diverse pharmacological properties including anti-oxidant activity. Soetikno et al. have shown the prevention of DCM by curcumin in a STZ-induced diabetic rat model, and also found that the prevention of DCM by curcumin is associated with the suppression of NOX activation leading to the overgeneration of ROS and/or RNS, either through inhibition of PKC or activation of Akt pathways. We also reported the prevention of DCM by curcumin analog C66 in a STZ diabetic mouse model through suppression of c-Jun NH2-terminal kinase-related oxidative stress.

**Coenzyme Q10**

Coenzyme Q10, a lipophilic cofactor of the mammalian mitochondrial electron transport chain, is not only crucial for mitochondrial energy production (adenosine 5'-triphosphate [ATP]), but also has emerged as an effective anti-oxidant. Coenzyme Q10 has shown some clinical benefits in a number of clinical trials. Huynh et al. have treated type 1 diabetic mice at 4 weeks after hyperglycemia that was induced by multiple-low doses of STZ (55 mg/kg/day for 5 days) with coenzyme Q10 or angiotensin-converting-enzyme inhibitor (ACE-I) ramipril for 8 weeks. They found that diabetes upregulated LV Nox2 and superoxide production along with LV oxidative damage along with LV diastolic dysfunction and remodeling. All these were almost completely prevented by coenzyme Q10, which was also compared favorably with improvements observed with ramipril. Therefore, coenzyme Q10 supplementation could thus represent an effective alternative to ACE-Is for
the treatment of cardiac complications in type 1 diabetic patients, as both medicines have multiple beneficial effects not only through suppression of oxidative stress.

**Resveratrol**

Resveratrol (RSV; trans-3,5,4′-trihydroxystilbene), a polyphenolic compound and naturally-occurring phytoalexin present in red wine and vegetable foods, was also used for the potential prevention or therapy for DCM. In experimental models of diabetes, RSV treatment was found to reduce the generation of ROS and the incidence of cardiomyocyte death, along with improvement of cardiac function.70-74

In summary, the aforementioned summarized anti-oxidants have been extensively applied in several other diseases, such as cancer and cardiovascular diseases, but whether these anti-oxidants can also be used for diabetic patients for the prevention of their DCM remains the subject of further systemic investigation, although the preliminary studies discussed here are promising.

**Metallothionein**

Metallothionein is a cysteine-containing (1/3 of 61 amino acids) and metal-binding protein. Physiological functions of MT include the maintenance of the homeostasis of essential metals and detoxification of heavy metals. MT maintains the redox status by binding and releasing metals (mainly zinc [Zn] under physiological conditions). Although four isoforms of MTs have been characterized, MT-I and MT-II exist as the major isoforms in various human and animal organs including the heart.75-76 The physical, chemical and biological properties of MT have been summarized by other reviews.77-80 Cardiac protection of MT against various oxidative damage induced by various oxidative stresses has been reviewed by others75,80-83. To address whether MT can protect the heart from diabetes, we have extensively explored the potential protection of MT from DCM since 200184,85. Type 1 diabetes was induced by a single injection of STZ in both cardiac-specific MT transgenic mice (MT-TG) and wild-type (WT) mice. No differences in hyperglycemia, bodyweight gain, and serum lipid peroxide levels were found between the MT-TG and the WT diabetic mice. However, serum creatine phosphokinase activity as a biomarker of cardiac injury was markedly increased in the WT diabetic mice. Significantly, cardiac injury was also found in the WT diabetic mice as evidenced by morphological changes. Therefore, these results provided the first evidence showing the preventive role of MT in the heart against diabetes-induced injury.84,85 Following these early studies, several investigations from the authors’ own laboratory and other laboratories86-90 in STZ-induced and transgenic OVE26 type 1 diabetic mouse models, and a high-fat diet-induced type 2 diabetic mouse model, further supported the significant protection of cardiac MT overexpression against diabetes-induced morphological and functional changes. Detailed review of the prevention of DCM by MT from several aspects, including experimental evidence, potential mechanisms and clinical implication, has been provided.41,91,92 In the following section, we provide some new insights based on the most recent studies.

We have shown the important role of diabetes-induced nitrosative damage in the development of DCM45,87,92, and MT can efficiently prevent DCM development through suppression of nitrosative damage. Recently, we found that diabetes can specifically induce ATPase and succinyl-CoA:3-ketoacid CoA transferase-1 (SCOT; Figure 2) and ATP synthase (Figure 3).93,94 We have identified a nitrated protein of 58 kDa as SCOT by mass spectrometry in the heart of diabetic mice. We found that although total SCOT expression was not significantly different between MT-TG and WT diabetic mice, the diabetic WT hearts showed significantly increased nitration content and dramatically decreased catalyzing activity of SCOT (Figure 2). Although SCOT nitration sites were identified at Tyr76, Tyr117, Tyr135, Tyr226, Tyr368 and Trp374, only Tyr76 and Trp374 were found to be located in the active site by 3-D structure modeling. However, only Trp374 showed a significantly different nitration level between the WT and MT-TG diabetic hearts. These results suggest that MT prevention of diabetes-induced pathological changes in cardiac tissues is most likely mediated by suppression of SCOT nitration at Trp374.94 In another study, using LC/MS analysis we showed that diabetes increased tyrosine nitration of the ATP synthase α subunit at Tyr271, Tyr311 and Tyr476, and the β subunit at Tyr269 and Tyr508, and also significantly reduced ATP synthase activity by ~32% (Figure 3). These changes were not observed in MT-TG diabetic mice.95 These results suggest that MT can preserve SCOT and ATP synthase activity in STZ-induced diabetes, probably through the inhibition of SCOT and ATP synthase nitration.

Velic et al.96 have utilized light microscopy (LM) and transmission electron microscopy (TEM) to show cardiomyocyte morphology and myocardial capillary basement membrane (CBM) thickness in the LV. Although the molecular mechanisms regulating CBM width remain elusive, it seems possible that despite a significant hyperglycemic environment, MT antioxidative activity could mitigate local oxidative stress and reduce downstream excess microvascular extracellular matrix (ECM) formation. In addition, the reduction of intra- and perivascular ROS and/or RNS might protect against incipient endothelial damage and the CBM thickening that results from such injury.

In summary, MT has shown strong cardioprotection from diabetes under different conditions: type 1 diabetes (STZ-induced and OVE26 transgenic diabetic mouse models), obesity-related diabetes in mice and even in humans with type 2 diabetes.95,96 As Zn is an important trace element that is required for more than 300 enzymes and transcription factors, and can induce cardiac MT97,98, and MT plays an important role in intracellular Zn, Zn supplementation could be an alternative approach to intervention of DCM in clinics in the future.
Figure 2 | 3-Ketoacid CoA transferase-1 (SCOT) protein expression, nitration, and activity in the hearts of diabetic wild-type (WT) and cardiac-specific metallothionein-overexpressing transgenic (MT-TG) mice. Type 1 diabetes was induced with a single dose of streptozotocin (STZ) in Friend virus B-type (wild-type) and cardiac-specific overexpression of MT-TG mice used for the study. (a) Cardiac tissues collected at the indicated post-induction times were examined by western blot to show changes in SCOT expression. (b) Cardiac tissues from non-diabetic and diabetic WT and MT-TG mice were immunoprecipitated with 3-nitrotyrosine (3-NT) antibody and western blotted with antibodies against succinyl-CoA:3-ketoacid CoA transferase, γ-glutamylcysteine synthetase, and NADPH quinone oxidoreductase (NQO1), heme oxygenase-1 (HO1), and γ-glutamylcysteine synthetase, to play a role in the process of detoxification, anti-oxidative stress and anti-inflammation. As Nrf2 is ubiquitously expressed, the implication of its roles in a variety of pathologies has been described mainly through the use of Nrf2 knock-out mice (Nrf2-KO) in different disease models.

The potential therapeutic action of Nrf2 in the treatment of several diseases including DCM has been suggested. In order to answer whether Nrf2 can provide protection against high glucose-induced oxidative damage, we carried out the first study to compare oxidative stress, apoptosis and the contractile activities in both Nrf2 WT cardiomyocytes and Nrf2-KO cardiomyocytes in vitro. Treated with different concentrations of glucose and 3-nitropropionic acid (inhibition of the mitochondrial respiratory complex II) caused increases of these effects in both kinds of cells in a concentration-dependent manner. Furthermore, the level of oxidative damage was more severe, and contractile activities were worse in Nrf2-KO cells than WT cells. It means that Nrf2 is critical in protecting cardiomyocytes from high glucose- and mitochondrial damage-induced oxidative lesions in cardiomyocytes. Following this early study, we and others have extensively investigated the potential protection of Nrf2 against diabetic complications including DCM. An early review regarding this topic is available. The following section we will provide some new information regarding the expression of Nrf2 in the heart under diabetic conditions, and how to protect the heart from diabetes by manipulating Nrf2 expression.

Nrf2 as an Adaptive Response Protein is Upregulated at the Early Stage of Diabetes, but Downregulated at the Late Stage

It is known that Nrf2 expression and function in cells in vitro and tissues in vivo are increased in response to oxidative stress. Several studies have shown the induction of ROS and/or RNS by high levels of glucose (HG) in cultured cardiovascular cells, and whether HG could elevate the Nrf2 expression and activation and its downstream gene expression in these cells has been investigated. Treatment with glucose at 20 and 40 mmol/L for 24 h increased the expression of Nrf2 messenger ribonucleic acid in primary cardiomyocytes or the H9C2 cardiac cell line. Nqo1, a prototype of Nrf2-regulated detoxification gene, was induced to be overexpressed in cardiomyocytes by such HG exposure too. Immunofluorescent confocal microscopic examination of the cells showed that HG treatment significantly increased the nuclear and total cell staining of Nrf2 in comparison with the control cells, indicating that glucose indeed increased the protein level and nuclear accumulation of Nrf2.

Nrf2

Nrf2, a transcription factor belonging to the Cap’n’coTr/basic region leucine zipper (CNC-bZIP) family of transcription factors, plays the role as a chief controller of intracellular redox status and the detoxification process. Nrf2 is bound to Kelch-like ECH-associated protein 1 (Keap1) in the cytoplasm, which facilitates its proteasomal degradation. Nrf2 can promote its target genes, such as NADPH quinone oxidoreductase (NQO1), glutathione S-transferase, heme oxygenase-1 (HO1) and γ-glutamylcysteine synthetase, to play a role in the process of detoxification, anti-oxidative stress and anti-inflammatory responses. As Nrf2 is ubiquitously expressed, the implication of its roles in a variety of pathologies has been described mainly through the use of Nrf2 knock-out mice (Nrf2-KO) in different disease models.

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Upregulation of Nrf2 and/or its downstream anti-oxidant genes in response to hyperglycemia were found not only in the cultured cells, but also observed in the heart of diabetic mice. We have used C57BL/6 mice to induce type 1 diabetes with a single dose of STZ. At 2 weeks after hyperglycemia onset, we found a significant upregulation of Nrf2 downstream genes NQO1 and HO1 messenger ribonucleic acid expression. Tan et al. have shown a different finding in terms of Nrf2 expression in the cardiac tissue from diabetic patients and controls.

Figure 3 | Adenosine 5’-triphosphate (ATP) synthase α and β subunit expression and nitration. Type 1 diabetes was induced with a single dose of STZ in Friend virus B-type (wild-type) and cardiac-specific overexpression of metallothionein transgenic (MT-TG) mice used for the study. (a) Total expressions of ATP synthase α and β subunits in wild-type (WT) and MT-TG mice from control or diabetes at 2, 4 and 8 weeks after diabetes onset were unchanged (30 μg total proteins/lane). (b) To define the nitration of ATP synthase α and β subunits, cardiac tissues (2 mg total proteins/lane) from WT control and diabetic mice along with MT-TG diabetic mice were immunoprecipitated with 3-nitrotyrosine (3-NT) antibody, followed by western blot for ATP synthase α and β subunits with anti-ATP synthase α and anti-ATP synthase β antibody, respectively. Diabetes-induced increased 3-NT in the heart was prevented in MT-TG mice. *P < 0.05; **P < 0.01. (c) The summary of LC/MS identifications of the nitrated sites for ATP synthase α and β subunit peptides. Bold letters in the table are LC/MS identifications for the nitrated sites of ATP synthase α and β subunit peptides, which are the nitrated sites at Tyr271, Tyr311, Tyr476, Tyr269 and Tyr508.

Tissue sections of left ventricles were obtained from autopsy heart specimens of humans with or without diabetes (all diabetic males had histories of hypertension and cardiac dysfunction). Nrf2 expression in the nuclei was significantly downregulated compared with the control hearts.

In support of the aforementioned point, our recent finding showed that Nrf2 protein expression was slightly increased in the heart of mice with hyperglycemia for 3 months, but significantly decreased in the heart of mice with hyperglycemia.
for 6 months\textsuperscript{107}. Combined with our early study\textsuperscript{103} in which Nrf2 downstream genes were increased in the heart of diabetic mice at 2 weeks after STZ-induced hyperglycemia, we assume that Nrf2 is adaptively trying to remain functional to overcome diabetic damage at the early stage of diabetes. At the late stage of diabetes, however, cardiac anti-oxidant function is further impaired, leading to a decrease in cardiac Nrf2 expression. Therefore, these aforementioned studies imply the preventive function of Nrf2 against diabetes-induced oxidative damage.

**Upregulation of Nrf2 Protects the Heart from Diabetes**

MG132 is a specific, potent, reversible and cell-permeable proteasome inhibitor, and thus is able to reduce the degradation of ubiquitin-conjugated proteins in mammalian cells. MG-132 as a proteasome inhibitor can upregulate Nrf2-mediated anti-oxidative function and downregulate nuclear factor-kappa-light-chain-enhancer of activated B cells (NF-\(\kappa B\))-mediated inflammation; therefore, it has been used to prevent diabetic renal damage\textsuperscript{108,109}. We have investigated too whether through the aforementioned two mechanisms MG-132 could provide a therapeutic effect on DCM in the OVE26 type 1 diabetes mouse model. OVE26 mice develop hyperglycemia at 2–3 weeks after birth, and show albuminuria and cardiac dysfunction at 3 months-of-age. Therefore, 3-month-old OVE26 diabetic and age-matched control mice were intraperitoneally treated with MG-132 daily for 30 months. Diabetic mice showed significant cardiac dysfunction, including increased LV systolic diameter and wall thickness, and decreased LV EF with an increase of the heart weight-to-tibia length ratio. Diabetic hearts showed structural derangement and remodeling (fibrosis and hypertrophy). In diabetic mice, there was also increased systemic and cardiac oxidative damage and inflammation. All of these pathogenic changes were reversed by MG-132 treatment. MG-132 treatment significantly increased the cardiac expression of Nrf2 and its downstream anti-oxidant genes with a significant increase of total anti-oxidant capacity, and also significantly decreased the expression of NF-\(\kappa B\) inhibitor alpha and the nuclear accumulation and DNA-binding activity of NF-\(\kappa B\) in the heart. These results suggest that MG-132 has a therapeutic effect on diabetic cardiomyopathy in OVE26 diabetic mice, possibly through the upregulation of Nrf2-dependent anti-oxidative function and downregulation of NF-\(\kappa B\)-mediated inflammation\textsuperscript{110}.

Sulforaphane (SFN) is a molecule containing an isothiocyanate functional group, which is obtained from cruciferous vegetables, such as broccoli, Brussels sprouts or cabbages, and shows anticancer and antidiabetic properties in experimental models\textsuperscript{111}. We have investigated whether the chronic use of SFN can prevent the development of DCM in a type 1 diabetes mouse model that was induced with multiple low-dose STZ (MLD-STZ). Diabetic and control mice were treated with SFN for 3 months and then kept for another 3 months without SFN treatment. SFN could almost completely prevent the development of DCM along with an upregulation of Nrf2 expression and transcription function in the heart. The data showed that the exposure of the cultured cardiac H9c2 cells to HG increased the fibrotic effect, which could be completely prevented by pretreatment with SFN, which also significantly increased Nrf2 expression and transcription. Silencing Nrf2 expression completely abolished the prevention by SFN of a HG-induced fibrotic effect, suggesting the important role of Nrf2 in the cardiac protection by SFN against HG in vitro or diabetes in vivo\textsuperscript{107}.

Similarly, diallyl trisulfide (DATS) is a most powerful anti-oxidant among the sulfur-containing compounds in garlic oil. It has been reported that treatment of H9c2 cells with HG resulted in an increase in intracellular ROS level and caspase-3 activity, which were markedly reduced by the administration of DATS. DATS treatment significantly increased Nrf2 protein stability and nuclear translocation, upregulated downstream gene HO-1 and suppressed its repressor, Keap1. However, apoptosis was not inhibited by DATS in cells transfected with Nrf2-specific small interfering ribonucleic acid. Similar results were also observed in high glucose-exposed neonatal primary cardiomyocytes and streptozotocin-treated diabetic rats fed DATS at a dose of 40 mg/kg bodyweight. The findings show that DATS protects against hyperglycemia-induced ROS-mediated apoptosis by upregulating the Nrf2 pathway to further activate Nrf2-regulated anti-oxidant enzymes in cardiomyocytes exposed to HG\textsuperscript{112}.

Therefore, Nrf2 as a new target of anti-oxidant therapy could have a great potential for clinical application, as these Nrf2 activators have a less toxic effect. In terms of the mechanism by which SFN prevents the heart from diabetes, we assumed that it is mediated by upregulation of Nrf2 because of the direct role of Nrf2 in the prevention of the HG-induced fibrotic effect in cultured H9c2 cardiac cells; however, whether the protection by SFN from DCM in vivo is also completely mediated by the upregulation of cardiac Nrf2 expression and function remains to be further examine, as well as whether SFN treatment is able to protect the heart from diabetes in the Nrf2-KO mouse model.

**CONCLUSIONS**

Diabetes mellitus has been recognized as a major cause of morbidity and mortality for decades. There is a large body of evidence to suggest that diabetic patients are prone to significant perturbations at the cellular and molecular level causing structural and functional abnormalities in the myocardium and vasculature, leading to DCM. As already described, some extra research steps are required so as to pinpoint the molecular mechanisms underlying the DCM before proceeding to the proposition of relevant clinical trials. New insights into the mechanisms that increase oxidative stress in diabetes might lead to novel treatment strategies. Clinically relevant translational studies in this direction might help to further unravel the underlying mechanisms of DCM.
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