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Oral Administration of an Angiotensin-Converting Enzyme 2 Activator Ameliorates Diabetes-Induced Cardiac Dysfunction

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Abstract

We evaluated the hypothesis that activation of endogenous angiotensin-converting enzyme (ACE) 2 would improve cardiac dysfunction induced by diabetes. Ten days after diabetes induction (streptozotocin, 50mg/kg, i.v.), male Wistar rats were treated with the ACE2 activator 1-[[2-(dimethylamino)ethyl]amino]-4-(hydroxymethyl)-7-[[4-(4-methylphenyl)sulfonyl]oxy]-9H-xanthen-9-one (XNT, 1mg/kg/day, gavage) or saline (control) for 30 days. Echocardiography was performed to analyze the cardiac function and kinetic fluorogenic assays were used to determine cardiac ACE and ACE2 activities. Cardiac ACE2, ACE, Mas receptor, AT₁ receptor, AT₂ receptor and collagen type I and III mRNA and ACE2, ACE, Mas, AT₁ receptor, AT₂ receptor, ERK1/2, Akt, AMPK- α and AMPK- β_1 protein were measured by qRT-PCR and western blotting techniques, respectively. Histological sections of hearts were analyzed to evaluate the presence of hypertrophy and fibrosis. Diabetic animals presented hyperglycemia and diastolic dysfunction along with cardiac hypertrophy and fibrosis. XNT treatment prevented further increase in glycemia and improved the cardiac function, as well as the hypertrophy and fibrosis. These effects were associated with increases in cardiac ACE2/ACE ratios (activity: ~26%; mRNA: ~113%; and protein: ~188%) and with a decrease in AT₁ receptor expression. Additionally, XNT inhibited ERK1/2 phosphorylation and prevented changes in AMPK- α and AMPK- β_1 expression. XNT treatment did not induce any significant change in AT₂ receptor and Akt expression. These results indicate that activation of intrinsic cardiac ACE2 by oral XNT treatment protects the heart against diabetes-induced dysfunction through mechanisms involving ACE, ACE2, ERK1/2, AMPK- α and AMPK- β_1 modulation.

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Keywords

Cardiomyopathy; ACE; Angiotensin-(1-7); Angiotensin II; AMP-activated protein kinases; MAP kinases

1. Introduction

Diabetic patients might present a peculiar cardiovascular entity, the diabetic cardiomyopathy, characterized by diastolic and systolic dysfunction without coronariopathy and hypertension [1, 2]. Cardiovascular diseases (CVD) are the most important cause of death in the diabetic population and diabetes increases two- to fourfold the risk of CVD [3]. Indeed, hyperglycemia is one of the metabolic and homeostatic abnormalities that increase the cardiovascular mortality in patients with diabetes [4, 5].

It has been well-known that hyperactivity of the Angiotensin-converting enzyme (ACE)/Angiotensin (Ang) II/AT₁ receptor axis of the Renin-angiotensin System (RAS) is associated with the establishment and progression of CVD and diabetes [6, 7]. On the other hand, recent studies have reported that ACE2 holds beneficial cardiovascular actions, such as anti-hypertensive [8], anti-fibrotic [9, 10], anti-oxidant [11], anti-inflammatory and anti-atherosclerotic [12, 13] effects. This enzyme is an important member of the RAS since it catalyzes the hydrolysis of the C-terminal residue of Ang II to produce the cardioprotective peptide Ang-(1-7) [14–16]. Thus, the axis composed by ACE2, Ang-(1-7) and Mas, the receptor for this peptide, plays a critical protective role in balancing the deleterious effects of the ACE/Ang II/AT₁ receptor axis.

It is consensus that blockage of the ACE/Ang II/AT₁ receptor axis improves morbidity, mortality and cardiovascular events in patients with CVD or high-risk diabetes [17–21]. However, the need for new therapies emerges due to ACE inhibitor intolerance and no consistent proof of specific cardiovascular protection by blockage of AT₁ receptor that exceeds or efficiently synergizes the effect of ACE inhibition [22, 23]. Recently, activation of the ACE2/Ang-(1-7)/Mas axis has been considered a suitable approach for modulating the pathological effects of ACE/Ang II/AT₁ receptor axis hyperactivity in CVD. In fact, ACE2 is a key regulator of the heart function [24] and its deficiency mimics or enhances the cardiac dysfunction induced by ACE/Ang II/AT₁ receptor axis hyperactivity [9, 13]. Genetic and pharmacological manipulation of ACE2 has been demonstrated to be an important strategy to treat diabetes. However, these studies have focused on organs other than the heart such as kidney and pancreas [25–27]. For instance, Bindom et al. [25] reported an improvement in fasting glycemia, glucose tolerance and islet insulin content along with an enhancement of beta cells viability in type 2 diabetic mice overexpressing ACE2 [25].

To explore the therapeutic potential of ACE2, we have identified small-molecule ACE2 activators based on the virtual screening of its structure [28]. One of these compounds is the 1-[(2-dimethylamino)ethylamino]-4-(hydroxymethyl)-7-[(4-methylphenyl)sulfonyl oxy]-9Hxanthene-9-one (XNT). Acute administration of XNT induced a dose-dependent hypotensive response in spontaneously hypertensive rats (SHR) while chronic treatment with XNT improved the cardiac function and reversed the cardiac and renal fibrosis in SHR [28]. Furthermore, XNT prevented the increase in right ventricular systolic pressure and hypertrophy in a monocrotaline-induced pulmonary hypertension model [29] and attenuated the thrombus formation in SHR [12]. Thus, since ACE2 causes beneficial end-organ outcomes in diabetes and reverses hypertension-induced cardiac fibrosis, we hypothesized that activation of endogenous ACE2 using XNT would produce therapeutic outcomes in diabetes-induced cardiac dysfunction. To test this hypothesis, physiological, histological and

molecular analyses were conducted to evaluate the effects of chronic oral XNT treatment in diabetic cardiomyopathy.

2. Material and methods

2.1. Materials

The ACE2 activator, XNT, was synthesized by Alchem Laboratories Corp. (Alachua, FL, USA). Fluorogenic substrates for ACE2 (catalog ID: ES007) and ACE (catalog ID: ES005) were obtained from R&D Systems (Minneapolis, MN, USA). The following antibodies were used in the western blotting assays: ACE2 (Santa Cruz sc20998, Santa Cruz, CA, USA), ACE (Millipore CD143, Billerica, MA, USA), Mas (Alomone AAR-013, Jerusalem, Israel), AT₁ receptor (Santa Cruz sc1173, Santa Cruz, CA, USA), AT₂ receptor (Alomone AAR-012, Jerusalem, Israel), ERK1/2 (phosphorylated and total: Cell Signaling #9106, Danvers, MA, USA and Santa Cruz sc93/sc154, Santa Cruz, CA, USA, respectively), Akt (phosphorylated and total: Cell Signaling #9271 and #9272, Danvers, MA, USA, respectively), AMPK- α (phosphorylated and total: Cell Signaling #2535 and #2603, Danvers, MA, USA, respectively), AMPK- β ₁ (phosphorylated and total: Cell Signaling #4181 and #4150, Danvers, MA, USA, respectively) and GAPDH (Sigma G8795, St. Louis, MO, USA). The following TaqMan[®] probes (Roche Indianapolis, IN, USA) were utilized in the RT-PCR assays: ACE2, Rn01416923-m1; ACE, Rn00561094-m1; Mas, Rn00562673-s1; AT₁ receptor, Rn02132799-s1; AT₂ receptor, Rn00560677-s1; collagen type I, Rn00801649-g1; and collagen type III, Rn01437683-m1.

2.2. Methods

2.2.1. Animals, diabetes induction and XNT treatment—All experimental protocols were performed in accordance with the Federal University of Minas Gerais (Brazil) and the University of Florida Institutional Animal Care and Use Committees, which are in compliance with the NIH guidelines. The experiments were performed in male Wistar rats (180–200g) obtained from CEBIO-UFGM (Belo Horizonte, MG, Brazil) and from Charles River Laboratories (Wilmington, MA, USA). They were housed in a light/dark cycle (12h/12h) room with standard rat chow and water *ad libitum*. Briefly, the rats were fasted (~16h), anesthetized (ketamine:xylazine, 60:6mg/kg, i.p.) and injected with streptozotocin (STZ; 50mg/kg i.v., Sigma; St. Louis, MO, USA) to induce diabetes. Control non-diabetic (CTL) rats were injected with ~0.2mL of sodium citrate buffer (10mmol/L, pH 4.5). Ten days after diabetes induction, the rats were assessed for blood glucose levels (Accu-Chek[®]; Roche, IN, USA). The animals with fasting blood glucose concentration over 126mmol/L were considered diabetic. The treatment with XNT (1mg/kg/day, gavage) or vehicle (saline pH 2–2.5; equivalent volume, gavage) was initiated ten days after diabetes induction and conducted for 30 days. The dose of 1mg/kg/day of XNT for oral administration was based on pilot experiments performed to determine the lowest amount of XNT able to induce cardiovascular effects. After testing the doses of 0.6mg/kg/day and 1mg/kg/day, we chose the dose of 1mg/kg/day based on the effects observed [30]. We have demonstrated in previous studies that XNT is able to activate ACE2 both *in vitro* and *in vivo* [10, 28].

2.2.2. Echocardiographic analysis—Transthoracic echocardiographic examination was performed using an Acuson Cypress[™] machine equipped with an 8-MHz linear-array transducer (Siemens; Munich, Germany). The rats were anesthetized with a ketamine/xylazine mixture (60:6mg/kg, i.p.) (CTL: n=8; STZ: n=7; STZ+XNT: n=6). Left ventricular systolic and diastolic functions were assessed by ejection fraction (EF) and mitral inflow pulsed-wave Doppler, respectively. Three measurements were performed: 1) initial - at diabetes induction (day 0 - D0); 2) intermediate - ~28 days after diabetes induction (day 28 - D28); and 3) final - at the end of the experiments (day 40 - D40). Two-dimensional guided

M-mode imaging at the papillary muscle level was used to measure the left ventricular end-systolic (LVESD) and end-diastolic (LVEDD) diameters and posterior wall thickness (LVPWT) during diastole. The EF was calculated from the M-mode echocardiogram using the equation: $EF(\%) = [(LVEDD^3 - LVESD^3) / LVEDD^3] \times 100$. Mitral inflow pulsed-wave Doppler velocity was recorded from the apical four-chamber view. All analyses were performed in a blinded way by the same echocardiographer and included morphological and functional parameters. Furthermore, to evaluate if the anesthesia used in our protocol could interfere in the parameters, we adjusted the data to heart rate and no significant differences were observed (data not shown).

2.2.3. Histological analysis—Heart beat was stopped in diastole using 10% KCl (i.v.). The hearts were fixed in 4% Bouin and stained with Hematoxylin & Eosin for cell morphometry (CTL: n=4; STZ: n=5; STZ+XNT: n=4) or with Picrosirius Red for fibrosis (CTL: n=3; STZ: n=3; STZ+XNT: n=4). Three sections (5 μ m) from each animal were visualized in a light microscope (BX41[®]; Olympus, Center Valley, PA, USA), photographed (Q-Color3[™]; Olympus, Center Valley, PA, USA) under x400 magnification and analyzed using the ImageJ software (<http://rsbweb.nih.gov/ij/>). Cardiomyocyte diameter of the left ventricular wall (~100 cardiomyocytes for each animal) was measured across the region corresponding to the nucleus. Only cardiomyocytes cut longitudinally with nuclei and cellular limits visible were considered for analysis. Cardiac interstitial fibrosis of the left ventricle was measured by area percentage analysis. All analyses were performed in a blinded way by the same researcher.

2.2.4. Insulin sensitivity test—Insulin sensitivity test was performed in overnight fed rats two days before the end of the treatment (day 38 - D38). After intraperitoneal injection of insulin (0.75 U/kg body weight; Sigma, MO, USA), tail-blood samples were taken at 0, 15, 30, 60 and 90 minutes for measurement of blood glucose levels (Accu-Chek[®]; Roche, IN, USA).

2.2.5. ACE and ACE2 activities—Enzymatic activity was measured in a microplate reader (BioTekSynergy[™] 2; BioTek, Winooski, VT, USA), as previously described [28]. Briefly, left ventricle samples (ACE: 30 μ g and ACE2: 60 μ g; n=4) were homogenized in a buffer composed by 75mmol/L Tris-HCl pH 7.5, 1mol/L NaCl and 0.5mmol/L ZnCl₂. All assays were performed in duplicate at pH 7.4 in a reaction mixture containing: 50 μ mol/L substrate, 5mol/L NaCl, 75mmol/L Tris-HCl and 0.5 μ mol/L ZnCl₂. Samples were read every minute for 4h immediately after the addition of the fluorogenic peptide substrates at 37°C. Background fluorescence readings were obtained from reactions without tissue samples and the final enzymatic activity of the samples was corrected by the obtained background value.

2.2.6. Real-time RT-PCR—Quantitative real-time RT-PCR (qPCR) was used to evaluate ACE2 (n=4), ACE (n=4), Mas (n=4), AT₁ receptor (n=4), AT₂ receptor (n=3), collagen type I (n=3) and collagen type III (n=4) mRNA levels in left ventricles. Total RNA was obtained using RNeasy-4-PCR kit (Applied Biosystems/Ambion; Austin, TX, USA) and 150ng (ACE, ACE2, Mas, AT₁ receptor and collagens) or 600ng (AT₂ receptor) of total RNA were reverse transcribed using the iScript[™] cDNA synthesis kit (Bio-Rad; Hercules, CA, USA) in 20 μ L of reaction mixture. The cDNA samples were amplified by qPCR using an ABI Prism[®] 7000 Detection system (Applied Biosystems, Foster City, CS, USA). Gene expression was quantified using the comparative Ct (threshold cycle) method [31] with GAPDH as an endogenous control.

2.2.7. Western blotting analysis—After the end of the XNT treatment, left ventricles were harvested and Western blotting technique was performed to quantify the protein density of ACE2, ACE, Mas, AT₁ receptor, AT₂ receptor, ERK1/2, Akt, AMPK- α and AMPK- β ₁. Briefly, proteins of the left ventricle were isolated, separated by electrophoresis, transferred to a nitrocellulose membrane and probed with one of the following primary antibodies: ACE2 (n=3; 100 μ g; 1:1000), ACE (n=3; 100 μ g; 1:1000), Mas (CTL: n=3; STZ: n=3; STZ+XNT: n=4; 100 μ g; 1:1000), AT₁ receptor (n=3; 15 μ g; 1:400), AT₂ receptor (n=3; 50 μ g; 1:400), ERK1/2 (n=3; 100 μ g; phosphorylated 1:1000 and total 1:5000), Akt (n=3; 50 μ g; phosphorylated 1:1000 and total 1:1000), AMPK- α (n=3; 100 μ g; phosphorylated 1:250 and total 1:500) and AMPK- β ₁ (CTL: n=3; STZ: n=3; STZ+XNT: n=4; 100 μ g; phosphorylated 1:1000 and total 1:2000). To obtain an endogenous control for protein quantification, the membranes were stripped and re-probed with GAPDH antibody (1:20000 or 1:30000) for ACE2, ACE, Mas, AT₁ receptor and AT₂ receptor or with the total ERK1/2, Akt, AMPK- α and AMPK- β ₁ antibodies. Protein bands were detected with Amersham™ ECL Plus Western Blotting Detection Reagents (GE Healthcare, Waukesha, WI, USA), their densities were acquired using the Molecular Imager® GS-800™ calibrated densitometer and quantified using the Quantity One 1-D analysis software (Bio-Rad, Hercules, CA, USA).

2.2.8. Statistical analysis—The data are expressed as mean \pm S.E.M. and $P < 0.05$ was considered statistically significant. Statistical analyses were performed using Student's *t*-test, ANOVA or Kruskal-Wallis analysis followed by the Newman-Keuls, Bonferroni or Dunn's post tests, as indicated in the Figure legends.

3. Results

3.1. Efficacy of the oral XNT treatment to increase left ventricular ACE2 activity

Firstly, we evaluated the effect of the oral XNT treatment during 30 days on cardiac ACE2 activity since in previous studies we have shown that XNT causes an increase in ACE2 activity *in vitro* [28]. XNT administration increased the activity of ACE2 in hearts by ~65% in normal rats (CTL: 0.43 \pm 0.08 vs. XNT: 0.71 \pm 0.03 Δ RFU/ Δ min/ μ g; $P < 0.05$).

3.2. Effects of XNT on cardiac dysfunction induced by diabetes

Given the pronounced cardiac ACE2 activation reached by the oral XNT treatment, we subsequently tested the effects of XNT on cardiac dysfunction observed in diabetic animals. The echocardiographic analysis revealed that diabetic rats presented diastolic dysfunction evidenced by decreases in the peak early (E) to atrial/late peak (A) velocity of the transmitral inflow ratio (E/A ratio) (Figs. 1A, 1B and 1C). The XNT treatment reduced these changes in the pattern of the E/A ratio (Fig. 1C). Indeed, we observed mitral regurgitation only in control diabetic animals (Fig. 1B). Along with the diastolic dysfunction, diabetic animals also showed a slight, but significant, decrease in the EF. The XNT treatment efficiently prevented the alteration in the EF (Fig. 1D). However, no significant differences in LVEDD and LVESD were observed between the diabetic and diabetic+XNT groups (Figs. 1E and 1F). Importantly, these findings were not accompanied by changes in systemic arterial pressure (data not shown).

3.3. Effects of XNT on cardiac remodeling induced by diabetes

The histological analysis showed that the treatment with XNT prevented the cardiac hypertrophy observed in diabetic rats (Figs. 2A and 2B). A similar effect was viewed when the cardiac fibrosis was analyzed, i.e. the XNT administration decreased the fibrosis induced by diabetes (Figs. 2C and 2D). Also, diabetes caused a significant increase in the expression of collagen type III in the heart and this effect was reduced by the XNT treatment (Fig. 2E).

On the other hand, no significant changes were seen in the collagen type I expression among any of the groups (Fig. 2F).

3.4. Modulation of the cardiac RAS induced by XNT in diabetic animals

To investigate the mechanisms underlying the effects of XNT on diabetic cardiomyopathy, we analyzed the balance between ACE and ACE2. Oral treatment with XNT increased the ACE2 activity, as well as the mRNA expression in hearts of diabetic animals. No significant changes were observed in the ACE2 protein expression among any of the groups (Table 1). On the other hand, ACE2 activation decreased the ACE mRNA and protein expression in diabetic rats. No significant changes were observed in the ACE activity among any of the groups (Table 1). More important, we evaluated the ratios between the two ACEs. As expected, diabetic animals presented a decreased ACE2/ACE protein ratio (~76%) and XNT efficiently increased this ratio by ~188%. The ACE2/ACE activity ratio was reduced by ~14% in diabetic rats and XNT administration increased it by ~26% when compared with diabetic animals. In terms of mRNA expression, the ratio between ACE2 and ACE was increased in diabetic rats (~34%) and a further increase of ~113% was induced by XNT administration (Table 1).

Additionally, the expression of AT₁ receptor, AT₂ receptor and Mas was examined. We observed that diabetes caused a trend toward enhancing the AT₁ receptor mRNA expression along with a significant increase in AT₁ receptor protein expression. These effects were absent in diabetic animals treated with XNT (Table 2). A robust increase without any significant change in protein expression was observed in Mas mRNA expression in diabetic rats. This effect was reverted by XNT administration (Table 2). Furthermore, diabetic animals presented an augmentation in the AT₂ receptor protein expression when compared with control rats and XNT did not interfere in this effect (Table 2). Regarding the AT₂ receptor mRNA expression, we observed RT-PCR amplification only in diabetic animals treated or not with XNT and any significant difference was viewed between these two groups (data not shown). In keeping with the beneficial effects of XNT on the ACE2/ACE ratios, diabetic animals treated with saline presented a reduction of ~44% in the Mas/AT₁ receptor protein expression ratio while an increase of ~78% in the Mas/AT₁ receptor protein expression ratio was observed in diabetic animals treated with XNT. Regarding the mRNA expression, the ratio between Mas and AT₁ receptor increased in diabetic rats treated or not with XNT (~125%) (Table 2).

3.5. Effects of XNT on hyperglycemia, AMPK and MAPK expression in diabetic rats

Since AMPK play a role in glucose homeostasis and in cardiac performance, we explored the putative involvement of these kinases in the cardiac protection triggered by XNT in diabetes. Diabetic animals presented hyperglycemia and XNT significantly prevented a further increase in blood glucose levels in diabetic animals after 30 days of treatment (Fig. 3A). Interestingly, diabetic rats treated with XNT presented an increased sensitivity to insulin. This effect was evident 90 minutes after the administration of insulin (Fig. 3B). Also, phosphorylation of the catalytic AMPK isoform (AMPK- α) was reduced while the regulatory AMPK isoform (AMPK- β_1) was augmented in hearts of diabetic rats. The XNT treatment prevented these effects on both AMPK isoforms (Figs. 4A and 4B).

Finally, we investigated the left ventricular expression of mitogen activated protein kinases (ERK1/2 and Akt) in diabetic rats treated with XNT. ACE2 activation reduced the increased phosphorylation of ERK1/2 observed in diabetic animals (Fig. 5A). On the other hand, no significant changes were viewed in Akt expression among any of the groups (Fig. 5B).

4. Discussion

The ACE2 is an important regulator of the cardiac function [24] and several studies have shown that this enzyme induces beneficial outcomes in CVD and diabetic nephropathy [9–13, 26, 27]. Despite the relevance of the diabetic cardiomyopathy, the role of ACE2 in cardiac function and structure in diabetic animals has not been fully investigated. Thus, in the current study we addressed the effects of an ACE2 activator on the cardiac functional and structural changes induced by diabetes.

The most significant findings of this study are that oral administration of an ACE2 activator produced beneficial outcomes in diabetes-induced cardiomyopathy and these effects were independent of decreases in plasma glucose levels. The use of a chronic treatment with the ACE2 activator, XNT, prevented the diastolic dysfunction induced by diabetes. An important cause of diastolic dysfunction is the structural remodeling of the heart due to increased myocardial stiffness [32]. On this way, along with the improvement in the cardiac function, the XNT inhibited hypertrophy, fibrosis and the increase in collagen type III mRNA expression induced by diabetes. Also, the XNT treatment modulated the ACE2 and ACE expression and the phosphorylation of ERK1/2. Indeed, additionally to the protective role in the cardiac function, several studies have shown that ACE2 plays a role in the maintenance of the heart structure. ACE2 activation prevented cardiac fibrosis and its replacement in ACE2 knockout mice attenuated the Ang II-induced adverse myocardial remodeling through inhibition of ERK1/2 [9, 10, 28]. In hearts, ERK1/2 is increased after four weeks of diabetes induction and it induces ventricular hypertrophy along with pathological remodeling, chamber dilation and diastolic dysfunction [33–35]. In fact, we have demonstrated that the anti-fibrotic effect of chronic administration of XNT in spontaneous hypertensive rats is associated with a decrease in ERK1/2 phosphorylation [10]. Therefore, ACE2 activation may contribute to the maintenance of the heart function and its structural integrity by balancing the activity of the ACE/Ang II/AT₁ receptor and ACE2/Ang-(1-7)/Mas axes of the RAS, as well as by inhibiting the ERK1/2 activity. One limitation of our study is that we did not measure the plasma and cardiac Ang II and Ang-(1-7) levels. However, it is important to note that qualitative immunohistochemical analysis showed that Ang-(1-7) immunostaining is increased in hearts of rats treated with XNT [10].

It has been suggested that ACE2 holds a role in glucose homeostasis [25]. Hyperglycemia induces myocardium damage through RAS activation, oxidative stress, calcium homeostasis alteration and structural changes due to glycation end product formation, hypertrophy and fibrosis [36–38]. We found that XNT prevented further increase in glycemia in diabetic animals. This effect might be involved in the cardioprotection observed in XNT-treated rats. Thus, activation of pancreatic ACE2 could favorably influence the heart, probably by regulating the pancreatic beta cells function and improving the metabolism of the heart. This premise is supported by the recent data of Bindom et al. [25], who showed that overexpression of ACE2 in the pancreas of db/db mice increases islet insulin content and beta cell proliferation, reduces beta cell apoptosis and improves fasting glycemia. It is important to note that because XNT prevented further increase in blood glucose levels and improved the insulin sensitivity in diabetic animals, this certainly contributed to the limitation of the development of cardiovascular damages. However, this was not the main mechanism underlying the beneficial effects of XNT since the blood glucose levels were still very high as 300 mg/dL in XNT-treated diabetic rats. Unfortunately, we did not investigate the pancreatic mechanisms involved in the XNT effects. Thus, future studies are needed to clarify the effects of XNT on glucose metabolism and insulin synthesis and degradation.

As an important cellular energy sensor and regulator of the metabolic homeostasis, activation of AMP-activated protein kinases (AMPK) leads to shifting towards an energy-producing state from an energy-consuming state [39, 40]. AMPK is a heterotrimeric protein consisting of α (catalytic), β and γ (regulatory) subunits [41–43]. It has been shown that AMPK can mediate GLUT4-dependent and insulin-independent muscular glucose uptake and increase fatty acid oxidation. In addition, its activity is inhibited in presence of high glucose concentration [44–48]. In hearts, inhibition of AMPK is involved in diabetic cardiomyopathy development and its activation can prevent post-ischemic cardiac dysfunction [49–51]. We demonstrated that diabetic animals treated with saline showed a decrease in the AMPK- α phosphorylation along with an increased AMPK- β_1 phosphorylation. Administration of XNT prevented these changes. Thus, it is tempting to suggest that ACE2 modulates AMPK activity improving cardiac metabolic imbalance and resulting in beneficial outcomes in diabetes-induced cardiac dysfunction.

Nowadays, there are few studies evaluating the biological effects of small molecule ACE2 activators. We have consistently demonstrated that these molecules are potent activators of ACE2 both *in vitro*, as well as *in vivo* [10, 12, 28, 29]. However, it is pertinent to note that off-target effects of XNT on these beneficial outcomes cannot be ruled out at the present time. In fact, XNT treatment induced an increase in ACE2 activity and mRNA expression, but no changes in protein expression, indicating that changes in the post-transcriptional control of the ACE2 gene expression may be induced by XNT.

5. Conclusion

In this study, we demonstrated a functional and structural protection in hearts of diabetic animals promoted by XNT, an ACE2 activator. This cardioprotection involved the enhancement of the ACE2/ACE and Mas/AT₁ receptor ratios and it was associated with reduction of the ERK1/2 phosphorylation, prevention of changes in AMPK- α and AMPK- β_1 expression and prevention of cardiac hypertrophy and fibrosis. Thus, ACE2 activators could be supplemented with antihyperglycemic drugs as a combined therapy for the control of end-organ damages caused by diabetes. Our findings provide a promising therapeutic strategy to treat diabetic cardiomyopathy by oral XNT.

Supplementary Material

Refer to Web version on PubMed Central for supplementary material.

Acknowledgments

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Highlights

- We tested if XNT, an ACE2 activator, would improve diabetes-induced heart dysfunction
- Diabetic rats showed heart dysfunction/hypertrophy/fibrosis which were blunted by XNT
- ACE2 activation also increased cardiac ACE2/ACE ratio and decreased AT₁ expression
- Inhibition of ERK and modulation of AMPK- α and β_1 are also involved in these effects
- Activation of intrinsic ACE2 protects the heart against diabetes-induced dysfunction

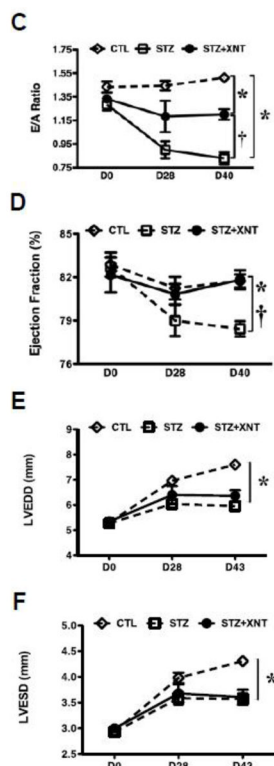
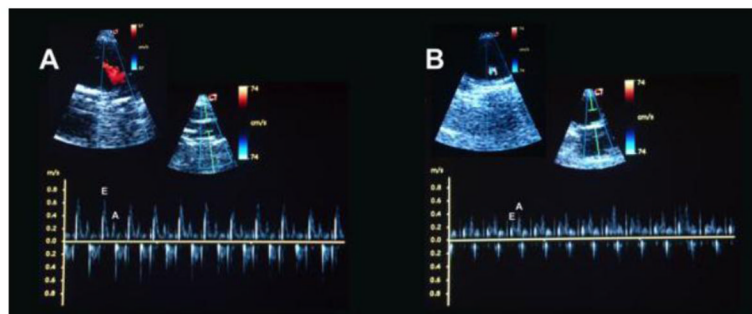
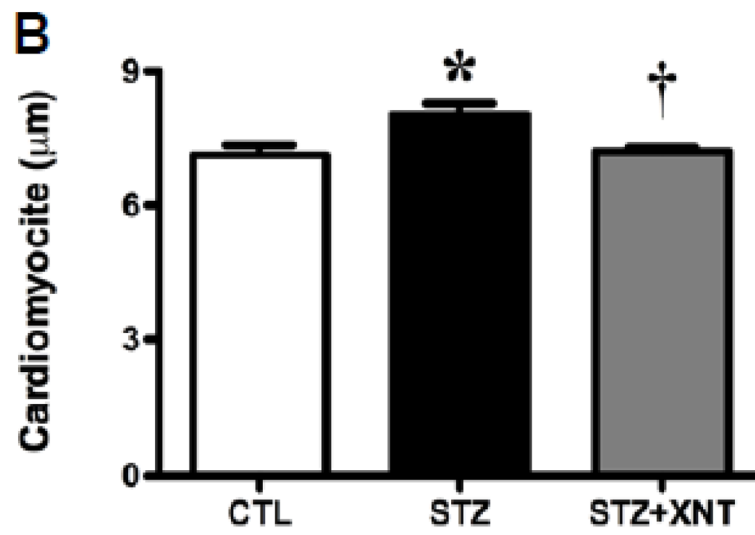
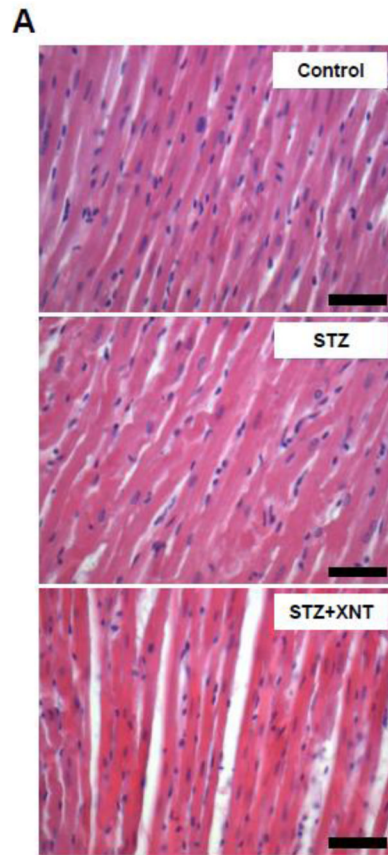
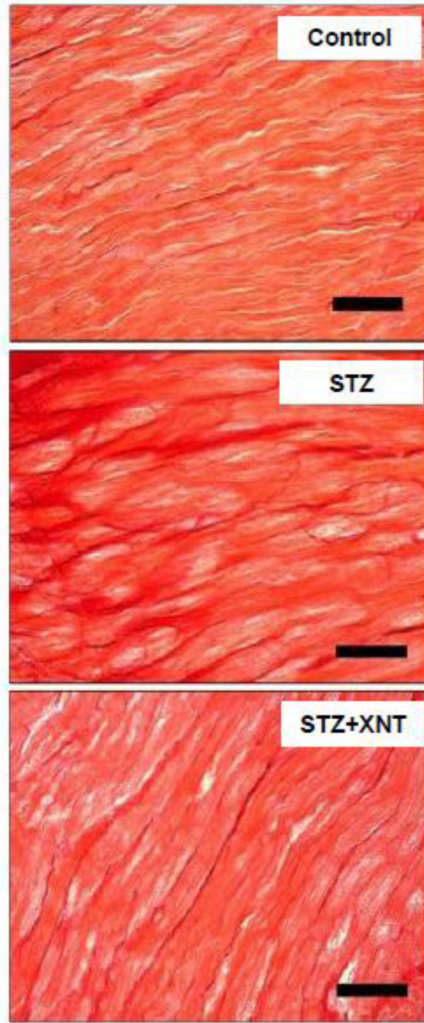


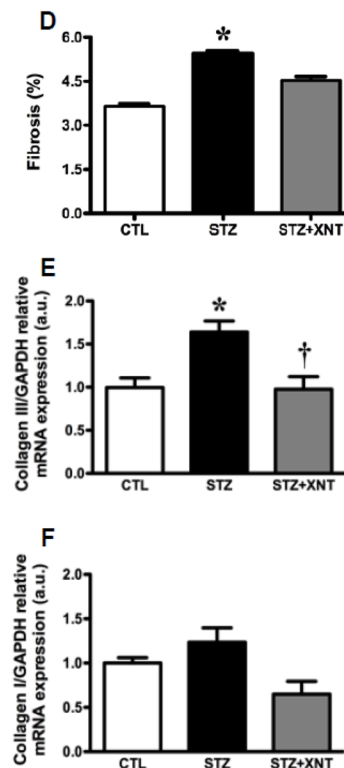
FIGURE 1.

Effects of ACE2 activation on cardiac function of diabetic animals. Representative transmitral pulsed-wave Doppler of a (A) control animal, displaying normal early (E) and late (A) mitral valve inflow waves and of a (B) diabetic animal treated with saline, showing inversion of the E and A waves. At the left upper side of the panel B it is possible to observe mitral regurgitation. (C) Peak velocity of E and A waves (E/A) ratio of the mitral valve inflow; (D) ejection fraction (%); (E) left ventricular end diastolic diameter (LVEDD, mm); and (F) left ventricular end systolic diameter (LVESD, mm) of control (CTL, n=8) and diabetic animals treated with saline (STZ, n=7) or XNT (STZ+XNT, n=6). The data are shown as mean \pm S.E.M. * P <0.05 vs. CTL and † P <0.05 vs. STZ. Two-way ANOVA followed by the Bonferroni post test.

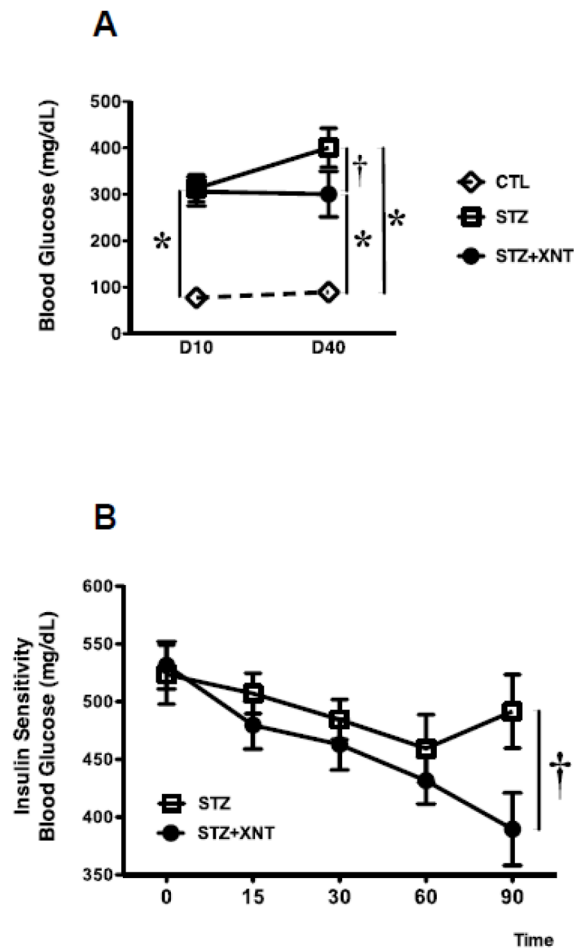


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**FIGURE 2.**

Effects of XNT on the cardiac remodeling induced by diabetes. (A) Representative photomicrographs of cardiomyocyte diameters of control animals, diabetic rats treated with saline (STZ) and diabetic animals treated with XNT (STZ+XNT). H&E (scale bar = 40 μm). (B) Cardiomyocyte diameter quantification (μm) (CTL: n=4; STZ: n=5; STZ+XNT: n=4). (C) Representative photomicrographs of left ventricular interstitial fibrosis of control animals, diabetic rats treated with saline (STZ) and diabetic animals treated with XNT (STZ+XNT). Intense red color indicates fibrosis. Picosirius-Red staining (scale bar = 40 μm). (D) Quantification of the interstitial fibrosis (%) (CTL: n=3; STZ: n=3; STZ+XNT: n=4). mRNA expression of collagen type (E) III and (F) I in control animals, diabetic rats treated with saline (STZ) and diabetic animals treated with XNT (STZ+XNT). Collagen type III (n=4) and collagen type I (n=3). The data are shown as mean \pm S.E.M. * P <0.05 vs. CTL and † P <0.05 vs. STZ. One-way ANOVA followed by the Newman-Keuls post test or Kruskal-Wallis analysis followed by the Dunn's post test.

**FIGURE 3.**

Effects of diabetes and XNT treatment on glucose regulation. **(A)** Fasting blood glucose levels (mg/dL) after 10 and 40 days (D10 and D40, respectively) of diabetes induction. CTL (n=15, white diamond), STZ (n=11, white rectangle) and STZ+XNT (n=16, black circle). **(B)** Insulin sensitivity in control and diabetic rats treated or not with XNT (STZ: n=11; STZ+XNT: n=8). The data are shown as mean \pm S.E.M. * P <0.05 vs. CTL and † P <0.05 vs. STZ. Two-way ANOVA followed by the Bonferroni post test.

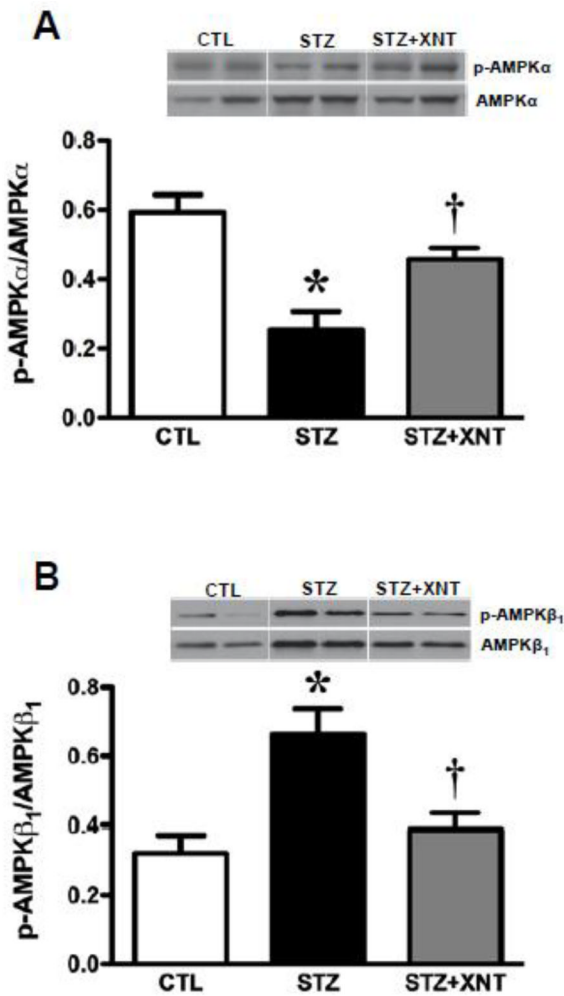


FIGURE 4.

Effects of XNT on cardiac AMPK expression in diabetic rats. (A) AMPK- α (total: 1:250, phosphorylated: 1:500; n=3) and (B) AMPK- β_1 (total: 1:1000, phosphorylated: 1:2000; CTL: n=3; STZ: n=3; STZ+XNT: n=4). The data are shown as mean \pm S.E.M. * P <0.05 vs. CTL and † P <0.05 vs. STZ. One-way ANOVA followed by the Newman-Keuls post test.



FIGURE 5. Effects of XNT on cardiac protein kinases expression in diabetic rats. (A) ERK1/2 (total: 1:1000, phosphorylated: 1:5000; n=3) and (B) Akt (total and phosphorylated 1:1000; n=3). The data are shown as mean \pm S.E.M. * P <0.05 vs. CTL and † P <0.05 vs. STZ. One-way ANOVA followed by the Newman-Keuls post test.

TABLE 1

Effects of XNT on the cardiac activity and expression of ACE2 and ACE.

	Activity (Δ RFU/min/ μ g)				mRNA (a.u.)				Protein (a.u.)				
	CTL	STZ	STZ+XNT	STZ	CTL	STZ	STZ+XNT	STZ	CTL	STZ+XNT	CTL	STZ	STZ+XNT
ACE2	0.49 \pm 0.07	0.54 \pm 0.06	0.76 \pm 0.06 ^{*†}	1.42 \pm 0.18	1.00 \pm 0.06	1.42 \pm 0.18	2.34 \pm 0.41 [*]	0.24 \pm 0.03	0.34 \pm 0.05	0.087 \pm 0.002	0.26 \pm 0.03 [*]	0.150 \pm 0.003 [†]	0.41 \pm 0.11
ACE	1.11 \pm 0.18	1.42 \pm 0.26	1.58 \pm 0.21	1.05 \pm 0.03	1.00 \pm 0.05	1.05 \pm 0.03	0.82 \pm 0.04 ^{*†}	0.087 \pm 0.002	0.087 \pm 0.002	0.087 \pm 0.002	0.26 \pm 0.03 [*]	0.150 \pm 0.003 [†]	0.150 \pm 0.003 [†]
ACE2/ACE	0.44 \pm 0.10	0.38 \pm 0.08	0.48 \pm 0.07	1.34 \pm 0.18	1.00 \pm 0.07	1.34 \pm 0.18	2.85 \pm 0.52 ^{*†}	0.95 \pm 0.17 [*]	3.93 \pm 0.55	3.93 \pm 0.55	0.95 \pm 0.17 [*]	2.73 \pm 0.71	2.73 \pm 0.71

The data are shown as mean \pm S.E.M. Activity: ACE2 (n=4, 60 μ g), ACE (n=4, 30 μ g), mRNA: ACE2 and ACE (n=4), Protein: ACE2 and ACE (n=3, 1:1000), GAPDH (1:30000). Please see the representative blots in the Supplementary data.

* $P < 0.05$ vs. Control (CTL) group and

[†] $P < 0.05$ vs. Diabetes (STZ) group. Kruskal-Wallis analysis followed by the Dunn's post test or One-way ANOVA followed by the Newman-Keuls post test (activity).

TABLE 2

Effects of XNT on the cardiac expression of Ang II and Ang-(1-7) receptors.

	mRNA (a.u.)				Protein (a.u.)			
	CTL	STZ	STZ+XNT	CTL	STZ	STZ+XNT	CTL	STZ+XNT
Mas	1.00±0.20	3.11±0.12*	1.84±0.19	0.12±0.01	0.12±0.01	0.11±0.01		
AT₁R	1.00±0.17	1.38±0.20	0.82±0.04 [†]	0.17±0.02	0.30±0.02*	0.16±0.02 [†]		
Mas/AT₁R	1.00±0.26	2.25±0.34*	2.24±0.33*	0.72±0.11	0.40±0.04	0.71±0.11		
AT₂R	-	-	-	0.04±0.01	0.15±0.01*	0.15±0.03*		

The data are shown as mean ± S.E.M. mRNA: Mas and AT₁ receptor (n=4), AT₂ receptor (n=3). Protein: Mas [1:1000, Control (CTL): n=3; Diabetes (STZ): n=3; Diabetes+XNT (STZ+XNT): n=4]; AT₁ receptor (1:400; n=4), AT₂ receptor (1:400; n=3). Please see the representative blots in the Supplementary data.

* $P < 0.05$ vs. Control (CTL) group and

[†] $P < 0.05$ vs. STZ group. One-way ANOVA followed by the Newman-Keuls post test.