


Endoplasmic Reticulum (ER) Stress in Part Mediates Effects of Angiotensin II in Pancreatic Beta Cells

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Introduction: The renin angiotensin aldosterone system (RAAS) is a hormone system known for its role in regulating blood pressure and fluid balance. Numerous RAAS inhibitors routinely prescribed for hypertension have also beneficial effects in type 2 diabetes (T2D) prevention. RAAS components are expressed locally in many tissues, including adipose tissue and pancreas, where they exert metabolic effects through RAAS bioactive hormone angiotensin II (Ang II). Pancreatic beta cells are specialized insulin-producing cells; they have also developed endoplasmic reticulum (ER), which contributes to beta cell dysfunction, when proteins are misfolded in disease states such as T2D. However, no studies have investigated the relationship between RAAS and ER stress in beta cells as a mechanism linking pancreatic RAAS to T2D. Hence, we hypothesized that Ang II treatment of beta cells increases ER stress and inflammation leading to reduced insulin secretion.

Methods: To test this hypothesis, we treated clonal INS-1E beta cells and human islets with Ang II and assessed changes in ER stress markers. INS-1E beta cells were also used for measuring insulin secretion and for assessing the effects of various RAAS and ER stress inhibitors.

Results: We demonstrated that Ang II significantly increased the expression of ER stress genes such as *Chop* and *Atf4* and reduced insulin secretion. Furthermore, inhibition of Ang II production with an angiotensin converting enzyme inhibitor (ACEi, captopril) significantly reduced ER stress. Moreover, the Ang II receptor blockade reduced ER stress significantly and rescued insulin secretion.

Discussion: This research provides new mechanistic insight into the role of RAAS activation via ER stress on beta cell dysfunction and provides additional evidence for protective effects of RAAS inhibition in T2D.

Keywords: beta cells, renin angiotensin aldosterone system, RAAS, type 2 diabetes, ER stress, inflammation

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Introduction

Type 2 diabetes mellitus (T2D) is an emerging epidemic worldwide, with the number of people affected rising significantly over the past several decades.¹ It is a complex metabolic condition that requires the simultaneous presence of two defects: insulin resistance and impaired pancreatic beta cell function.² Pancreatic beta cells in the early stages of insulin resistance, secrete more insulin to compensate for increasing insulin resistance; however, this leads to an extra burden on beta cells, which fail at some point resulting in T2D.³

Several mechanisms have been associated with pathogenesis and the development of T2D. One of these is the renin angiotensin aldosterone system (RAAS), which is classically known to regulate blood pressure and fluid balance. While numerous studies reported contributions of RAAS to T2D,^{4–6} and benefits of RAAS inhibitors in T2D prevention,⁷ the role of this system in beta cell function is relatively less explored. Angiotensinogen (Agt) is the primary precursor in RAAS and is catalyzed into smaller angiotensin peptides, including angiotensin II (Ang II), which is the main bioactive hormone product.⁸ Recently, components of RAAS have been discovered locally within various tissues, including adipose tissue and pancreas, with local functions.^{5,9–12} Mouse and human islets treated with Ang II demonstrated significantly reduced insulin secretion, which led to beta cell dysfunction.^{4,5}

Several clinical trials demonstrated that angiotensin receptor blockers (ARB) and angiotensin converting enzyme inhibitors (ACEi), primarily used as anti-hypertensive drugs also reduce T2D incidence by 25% in high-risk patients.^{7,13–15} This reduction is in part from protective actions on pancreatic islets through improved beta cell function.^{15–17} Furthermore, mice treated with ACEi have enhanced beta cell mass and improved beta cell proliferation markers.^{18,19} More importantly, pancreatic islets treated with agonists or blockers of the RAAS pathway points to a possible role of activated local RAAS in insulin secretion,^{5,20} but the precise mechanisms involved are not well established.

One of the downstream mechanisms through which RAAS contributes to obesity and insulin resistance is the endoplasmic reticulum (ER) stress.^{21,22} ER is a specialized secretory organelle, which is pivotal for protein folding. ER is especially critical in beta cells as around one million insulin molecules are produced per minute.²³ When ER is overburdened due to hyperinsulinemia, immature insulin and other unfolded proteins accumulate in ER lumen, leading to activation of unfolded protein responses (UPRs).²⁴ When UPR reaches beyond a certain capacity in cells, it results in ER stress, which then reduces insulin secretion leading to beta cell death and T2D.²⁵ Recently, microRNAs, which are post-transcriptional gene regulators, demonstrated regulation of cell survival or cell death through ER stress mediation in beta cells.²⁶ Specifically, miR-30 family and miR-708 were reduced or increased with ER stress, respectively.²¹ The detailed mechanisms by which RAAS regulates ER stress in beta cells are not completely understood. Hence, we hypothesized that activation of RAAS in beta cells led to increased ER stress and reduced insulin secretion; we further

hypothesized that both of which will improve with RAAS inhibition. The results of this study will increase understanding of the role of activated RAAS in beta cells on T2D progression and may help researchers develop potential new anti-diabetic therapeutic targets.

Materials and Methods

Cell Culture and Islet Procurement

Rat pancreatic INS-1E cell lines were a kind gift from Dr. Patrick Fueger (City of hope, CA) and approved by the Texas Tech University Research Ethics Committee. Cell lines were maintained in 5% CO₂ at 37°C in RPMI-1640 (Life Technologies, Carlsbad, CA) supplemented with 10% fetal bovine serum (FBS) (Thermo Fisher Scientific, Waltham, MA USA), 1 mM sodium pyruvate, 10 mM HEPES, 0.05 mM 2-mercaptoethanol, and 50 U/mL penicillin and 50 g/mL streptomycin (Thermo Fisher Scientific, Waltham, MA USA). Human islets were obtained through the Integrated Islet Distribution Program (<https://iidp.coh.org/>). We received shipments of human islets from de identified male subjects between 20–50 years with a BMI <30. Upon arrival, islets were hand-picked for further treatments.

INS-1E cells were treated with various doses (10–100 nM) of Ang II (Sigma-Aldrich, St. Louis, MO, USA) for 24 hours and 48 hours, respectively. Pancreatic human islets were treated with various doses of Ang II for 48 hours. For other experiments, various inhibitors such as 1 mg/mL captopril (Angiotensin-converting enzyme inhibitor; ACEi), 1000 nM P-186 (Angiotensin type 2 receptor inhibitor; P) and 1 mg/mL telmisartan (Angiotensin type 1 receptor inhibitor; T), 7.5 mM 4-phenyl butyric acid (4-PBA; ER stress inhibitor; Sigma-Aldrich, St. Louis, MO, USA) were used based on previous literature.^{21,27}

For glucose starvation treatments, cells were starved using HBSS buffer (1M NaHCO₃, 1M HEPES, 1M CaCl₂, 0.075M KCl, 2.5M MgSO₄, 5M NaCl, 1M KH₂PO₄ and 1g BSA) containing 3 mM glucose for 2 hours. Cells were then treated with HBSS buffer containing 15 mM glucose along with the respective treatments. After 2 hours, media collected was used for insulin secretion experiments, while cells were used for gene expression studies.

RNA Isolation and Gene Expression

RNeasy mini kit (Qiagen, Valencia, CA, USA) was used for RNA isolation, and cDNA was reverse transcribed using iScript reverse transcription supermix (BioRad, Hercules, CA, USA). Gene expression levels were

assessed by real-time quantitative polymerase chain reaction (RT-qPCR) using Sybr green master mix (BioRad, Hercules, CA, USA) and normalized to housekeeping genes (18S ribosomal RNA and GAPDH).

MicroRNA Expression

Total RNA was used for cDNA synthesis by reverse transcription using TaqMan™ Advanced microRNA cDNA Synthesis Kit (Thermo Fisher Scientific, Hereford, TX, USA). MicroRNA expression levels were assessed with real-time quantitative polymerase chain reaction (RT-qPCR) using the TaqMan™ Fast advanced master mix (Thermo Fisher Scientific, Hereford, TX, USA). The microRNA samples were normalized to housekeeping microRNAs (miR-191-5p and miR-186-5p).

Elisa

Insulin content was measured using enzyme-linked immunosorbent assay (ELISA) kits (EMD Millipore Corporation, St. Louis, Missouri, USA) according to the manufacturer's protocol.

Statistical Analyses

Results are presented as means \pm SEM where one-way ANOVA was performed with Tukey's posthoc test ($p < 0.05$) for experiments with more than two groups and analyzed using the Graph Pad Prism, version 8. On the contrary, Student's *t*-test was used for data with two groups. qRT-PCR assay results were analyzed using the CFX Manager software provided by Bio-Rad Laboratories, Inc., using the $2^{-\Delta\Delta CT}$ method. All in vitro experiments had at least three replicates.

Results

We tested whether Ang II activates ER stress in pancreatic beta cells using dose and time-dependent studies. We treated INS-1E cells with various doses of Ang II (10–100 nM) for 24 and 48 hours, respectively, to identify the optimal concentration of Ang II for inducing ER stress. We measured markers of ER stress, such as activating transcription factor 4 (*Atf4*) and CCAAT-enhancer-binding protein homologous protein (*Chop*). The expression of these genes was no different from untreated control cells at 24 hours (Figure 1A and B). However, *Atf4* mRNA levels were significantly increased by ~ 2.1 fold ($p < 0.05$) with 50 nM Ang II treatment at 48 hours compared to untreated control cells (Figure 1C). The *Chop* mRNA level increased significantly with all three doses of Ang II compared to control after 48

hours (Figure 1D). Ang II at 10 nM increased *Chop* levels by ~ 1.5 fold, while Ang II at 50 nM increased *Chop* mRNA by ~ 3 fold, respectively ($P < 0.05$) (Figure 1D). Consistent with these findings, glucose-stimulated insulin secretion was increased by ~ 8 fold ($p < 0.05$) as expected, with 15 mM glucose compared to 3 mM glucose at 24 hours (Figure 1E). Ang II at 50 nM in the presence of 15 mM glucose reduced insulin secretion significantly by ~ 4 fold ($p < 0.05$) compared to the 15 nM glucose at 24 hours (Figure 1E). At 48 hours, insulin secretion with 50 nM Ang II was significantly decreased by 3.7 fold compared to 15 mM glucose control, with no differences for 10 and 100 nM Ang II compared to control (Figure 1F). Based on the above findings, we chose 50 nM of Ang II to induce ER stress for subsequent experiments.

We validated the findings from INS-1E cells in human pancreatic islets treated with various doses of Ang II for 48 hours. As seen in rodent cells, *Atf4* gene level was significantly increased with 50 nM Ang II by ~ 2 fold (Figure 2A) ($p < 0.05$) while the *Chop* levels were increased by ~ 1.7 fold ($p < 0.05$) with Ang II at both 50 and 100 nM in human pancreatic islets (Figure 2B). Due to the cost and limited availability of human islets, other experiments were performed in INS-1E cells.

To further confirm the role of RAAS in ER stress, we treated INS-1E cells with captopril, an ACEi that inhibits the production of Ang II. Pancreatic beta cells treated with captopril for 48 hours significantly lowered gene levels of *Atf4* and *Chop* compared to control by ~ 1.5 fold ($p < 0.05$), as seen in Figure 3A and B. Furthermore, captopril treatment increased insulin secretion by ~ 1.6 fold ($P < 0.06$), which showed trending towards significance compared to control (Figure 3C).

Ang II effects are mediated downstream through two receptors of the RAAS, namely angiotensin type 1 receptor (AT1) and angiotensin type 2 receptor (AT2). As the RAAS pathway induces ER stress, we sought to determine the Ang II receptor type mediating these effects. We treated cells with Ang II in the presence of AT1 inhibitor (telmisartan) and AT2 inhibitor (P-186), either individually or in combination. While Ang II treatment alone induced ER stress as shown by significantly increased levels of *Atf4* and *Chop* mRNA levels by ~ 2 and 1.5 fold ($p < 0.05$) compared to control respectively (Figure 4A and B), treatment with either P-186 or telmisartan reduced both *Chop* and *Atf4* gene expression significantly, compared to Ang II-treated cells. Additionally, treatment with both inhibitors (telmisartan and P-186) also significantly lowered *Atf4* and *Chop* expression levels compared to control,

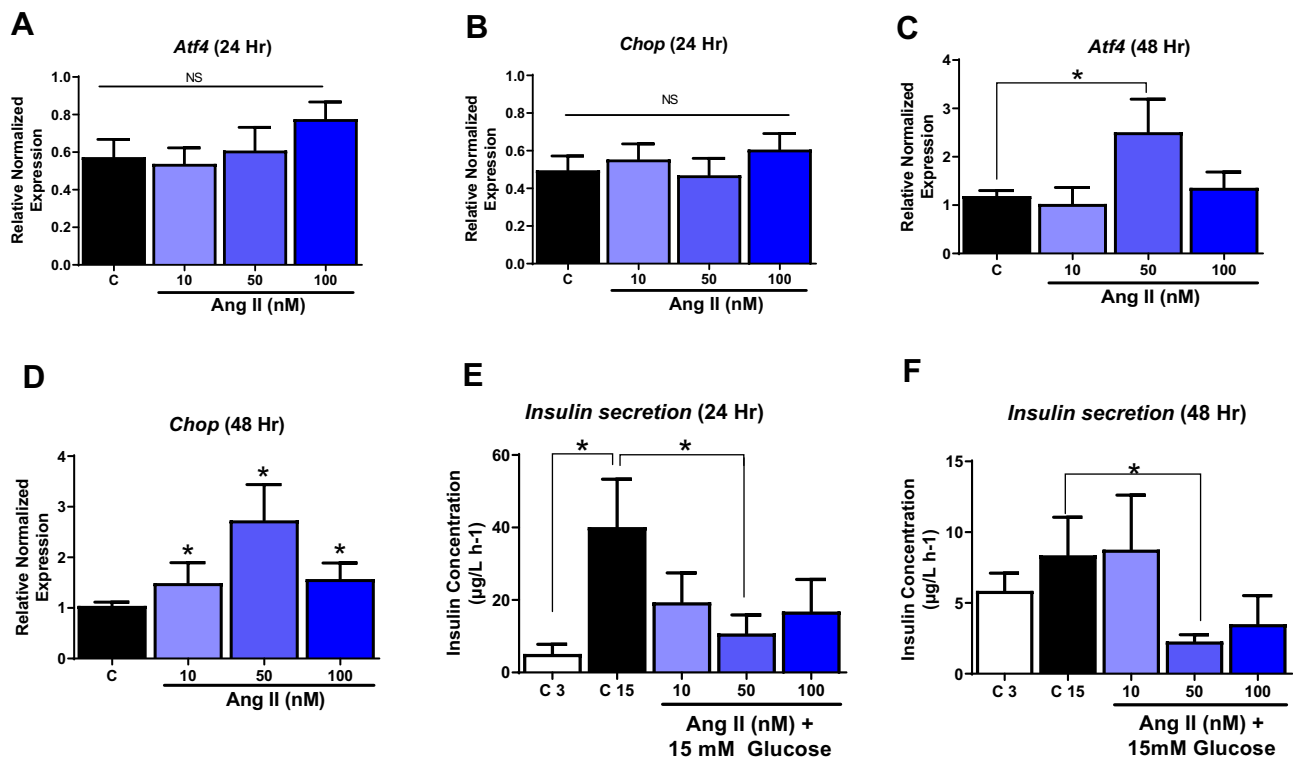


Figure 1 Dose and time-dependent effects of Angiotensin II. INS-1E cells were treated with different doses of Ang II to determine the optimal dose of Ang II to induce ER stress. (A and B) Ang II at all doses tested did not alter activated transcription factor 4 (*Atf4*) and C/EBP homologous protein (*Chop*) gene levels at 24 hours. (C) Ang II at 50 nM increased *Atf4* gene levels, and (D) Ang II at all doses tested increased *Chop* gene levels at 48 hours. (E) Ang II only at 50 nm reduced insulin secretion at 24 hours. (F) Ang II at 50 nM reduced insulin secretion at 48 hours. Data are presented as mean ± SEM (n=4). *p< 0.05 compared to control.

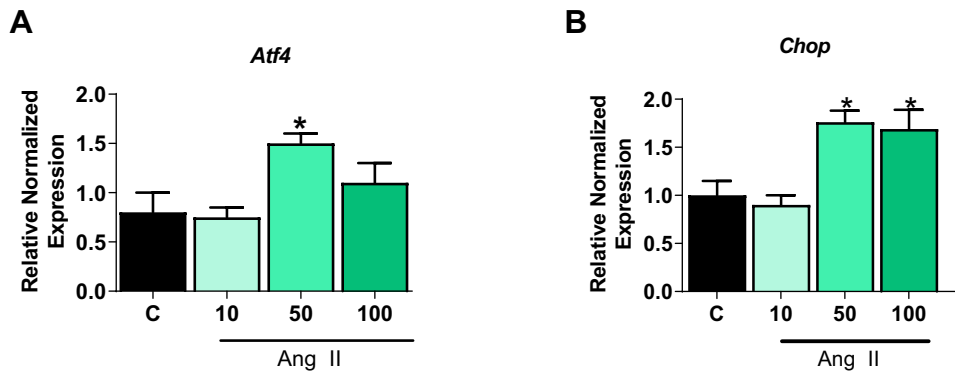


Figure 2 Dose and time-dependent effects of Angiotensin II on Human Pancreatic islets. Human Pancreatic islets were treated with different doses of Ang II for 48 hours to determine the optimal dose of Ang II to induce ER stress. (A) Ang II at 50 nM increased gene levels of activated transcription factor 4 (*Atf4*). (B) Ang II at both 50 and 100 nM increased C/EBP homologous protein (*Chop*) gene levels. Data are presented as mean ± SEM (n=4). *p< 0.05 compared to control.

suggesting ER stress is equally transmitted via both the AT1 receptor and AT2 receptor as shown in Figure 4A and B (p<0.05). Lastly, insulin secretion was significantly reduced by 50 nM of Ang II by ~ 2 fold but was rescued by telmisartan, P-186 individually as well as combined receptor treatments (Figure 4C).

To further validate Ang II effects on beta cells via ER stress, we tested whether the effects of Ang II could be reduced by ER stress inhibitor 4-PBA. INS-1E cells were

treated with Ang II (50 nM) alone or in the presence of 4-PBA for 48 hours, and ER stress markers were measured. As expected, Ang II induced ER stress by 1.5 fold; however, ER stress was significantly reduced with 4-PBA along with Ang II by ~ 3 and 5 fold (p<0.05), as shown by expression levels of both *Atf4* and *Chop* respectively in Figure 5A and B. Lastly, insulin secretion was significantly increased by ~ 1.9 fold (p<0.05) with Ang II along with 4-PBA compared to Ang II, as shown in Figure 5C.

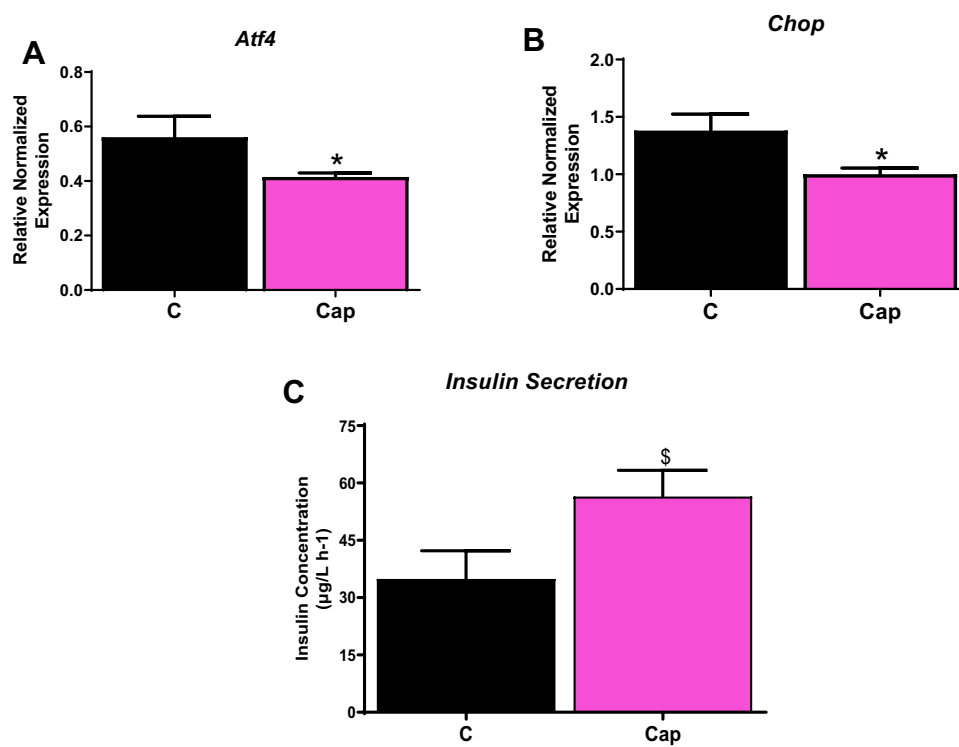


Figure 3 Captopril reduces ER stress. INS-IE cells treated with 100 μM captopril reduces ER stress as shown by (A) activated transcription factor 4 (*Atf4*) gene levels, (B) C/EBP homologous protein (*Chop*) expression gene levels and (C) improved insulin secretion. Data are presented as mean ± SEM (n=5 each group). *p< 0.05 compared to control. [§]p< 0.1.

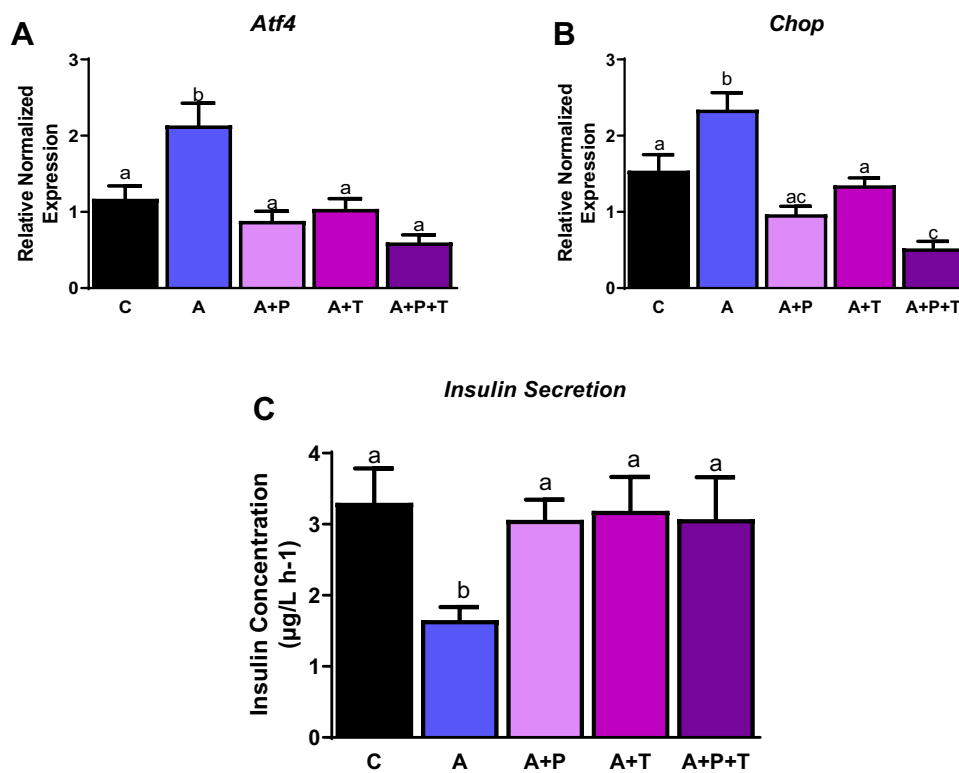


Figure 4 Angiotensin receptor inhibitors reduce ER stress. INS-IE cells treated with telmisartan (T; AT1 inhibitor), PI86 (P; AT2 inhibitor) reduce ER stress as shown by (A) activated transcription factor 4 (*Atf4*) gene levels, (B) C/EBP homologous protein (*Chop*) gene levels and (C) improved insulin secretion. Data are presented as mean ± SEM (n=4-5 each group). Means without the same letter are different.

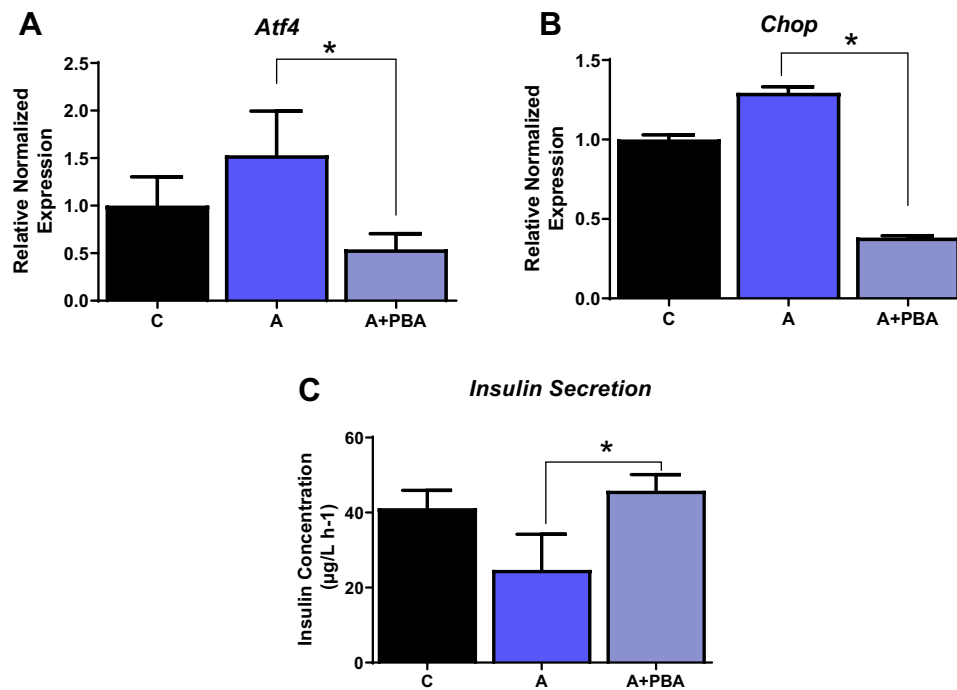


Figure 5 ER stress inhibitor reduces ER stress even in the presence of Angiotensin II. INS-1E cells treated with Ang II and ER stress inhibitor, 4-phenyl butyric acid reduces ER stress as shown (A) activated transcription factor 4 (*Atf4*), (B) C/EBP homologous protein (*Chop*) and (C) improved insulin secretion. Data are presented as mean \pm SEM (n=5 each group). *p< 0.05 compared to Ang II.

Since RAAS exerts proinflammatory effects in adipose tissue and adipocytes,¹⁰ we determined if activation of RAAS also led to inflammation in beta cells and, if so, whether this was mediated by Ang II receptors. We measured various inflammation markers, such as inducible nitric oxide synthase (*iNos*), nuclear factor-kappa B (*Nfkb*), and tumor necrosis factor-alpha (*Tnf- α*) in Ang-II treated INS-1E cells. Ang II significantly increased *iNos* mRNA levels by ~ 2 fold, while inhibition of AT1 and AT2 receptors individually or in combination reduced *iNos* mRNA levels, compared to Ang II (Figure 6A). Surprisingly, there were no differences with the above treatments for expression of *Nfkb* and *Tnf- α* genes (Figure 6B and C). In addition, we measured interleukin (IL)-6, interleukin-1 beta (*Il-1 β*), monocyte chemoattractant protein -1 (*Mcp-1*), and *Il-10* mRNA levels, but their expression levels were not detectable in INS-1E cells.

Lastly, as miRNAs have been previously reported to regulate ER stress and inflammation, and mediate Ang II effects in part in adipocytes,²¹ we specifically tested changes in miR-30 and miR-708-5p, negatively and positively associated with ER stress, respectively.^{21,28} As shown in Figure 7, Ang II treatment significantly reduced levels of miR-30c-5p by ~ 2 fold but not miR-30a-3p and miR-708-5p levels compared to control untreated cells.

Discussion

RAAS is a critical physiological regulator of blood pressure and fluid balance. However, additional local functions of RAAS have also been identified in several metabolic tissues, including adipose tissue and the pancreas.^{11,29} Components of RAAS (AT1, AT2, Agt, Ang II and ACE) are expressed in murine and human pancreatic islets with a local role within the pancreas.^{17,30-32} RAAS regulates islet blood flow;³³ however, specific non-hemodynamic functions in pancreatic beta cells are not well established.³⁴

In our current study, Ang II treatment, which mimics RAAS overactivation, induces ER stress in pancreatic beta cells and human islets. Furthermore, Ang II reduces insulin secretion in INS-1E cells. Additionally, inhibiting the RAAS pathway reduced ER stress and improved insulin secretion in beta cells. Furthermore, increased ER stress by Ang II was mediated via both AT1 and AT2 receptors. Moreover, we have shown that one of the common miRNA regulators of ER stress, miR-30, was also reduced in our studies. These studies find in part that ER stress induced by the RAAS pathway could be a mechanistic pathway that could contribute to T2D in beta cells.

The Ang II receptor, AT1, is primarily known to mediate vasoconstriction, oxidative stress, and inflammation,³⁵ while AT2 overall exerts beneficial properties such as vasodilation,

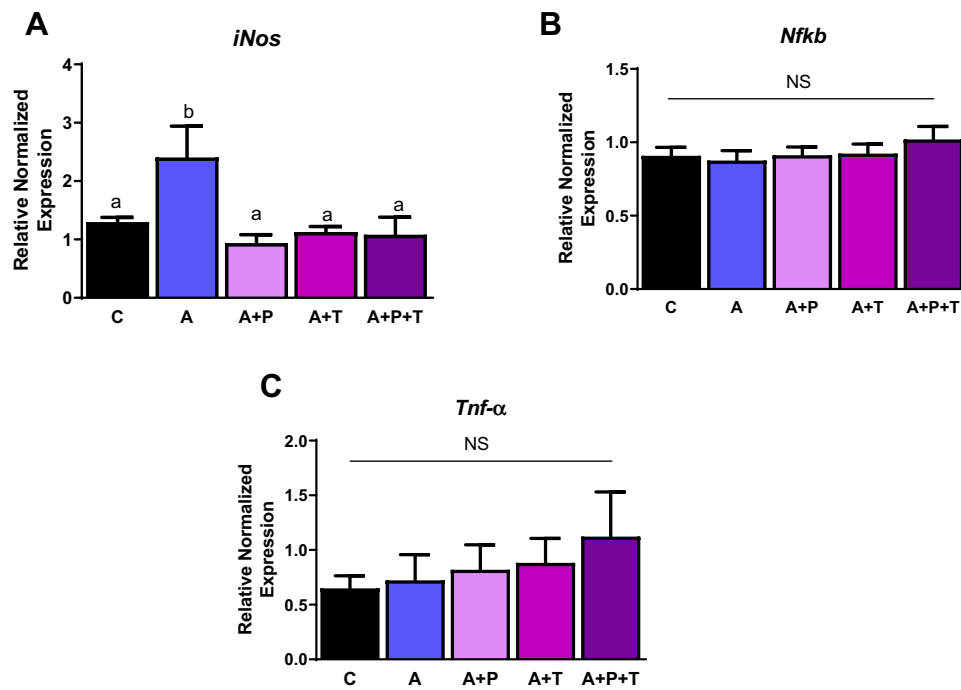


Figure 6 Angiotensin induces inflammation through *iNos*. INS-IE cells treated with telmisartan (T; AT1 inhibitor), P186 (P; AT2 inhibitor) reduce inflammation as shown by (A) inducible nitric oxide synthase (*iNos*) gene levels, (B) nuclear factor kappa-light-chain-enhancer of activated B cells (*Nfkb*) gene levels and (C) tumor necrosis factor alpha (*Tnf-α*). Data are presented as mean \pm SEM (n=5 each group). Means without the same letter are different.

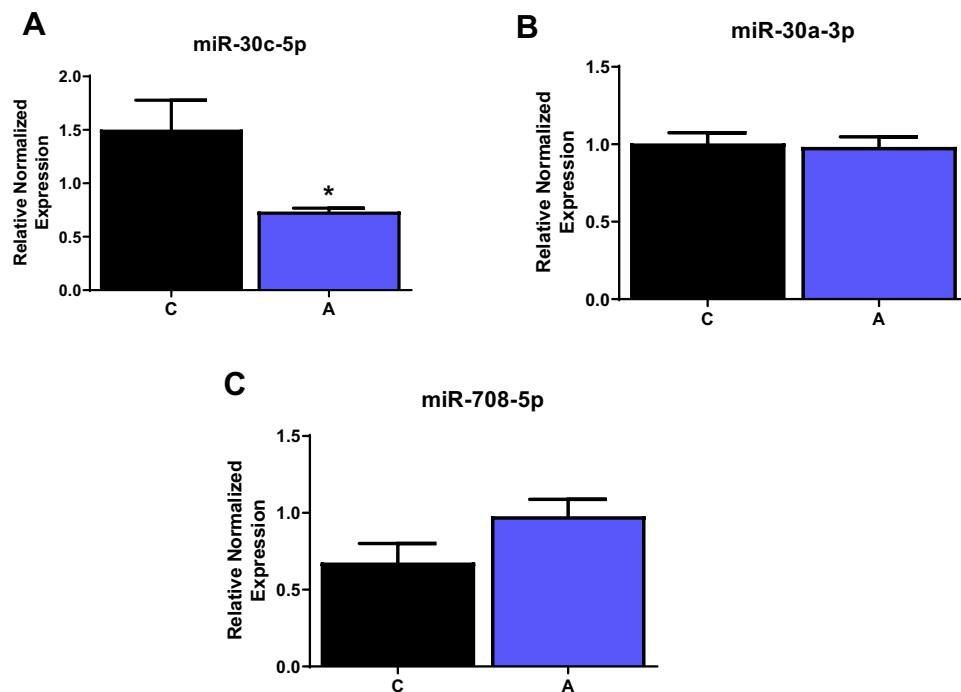


Figure 7 Alterations in microRNAs associated with ER stress. INS-IE cells were treated with Ang II, and levels of different miRNAs were measured. (A) *miR-30c* was reduced while (B) *miR-30a* and (C) *miR-708* was unaltered in the presence of Ang II. Data are presented as mean \pm SEM (n=3). *p<0.05.

anti-inflammatory, and anti-fibrotic effects.³⁶ In human islets, inhibition of AT1 receptor reduced ER stress and improved its antioxidant potential and insulin secretion.³⁷ These

findings agree with our results demonstrating that ER stress is reduced when interaction of Ang II with AT1 was blocked.³⁸ Additionally, we showed that ER stress was

also, in part, mediated by the AT2 receptor, typically considered a protective receptor in other cell types.^{36,39} However, in rodent clonal beta cells, overexpression of AT2 receptor increased apoptosis and reduced insulin secretion.⁴⁰ This suggests that AT2 receptor signaling may upregulate Chop activity, which overburdens ER, leading to increased apoptosis. This is in part consistent with our data showing that inhibition of AT2 reduced ER stress; however, whether this will trigger apoptosis needs to be further validated. Interestingly, in ob/ob mice, a model of T2D, AT1 receptor gene levels in islets were lower compared to healthy Wt mice where, AT2 receptor levels were on the other hand higher.⁴¹ The importance of such changes is specifically related to reduced ER stress and insulin secretion with AT2 blockers. In future studies, we will be using silencing approaches to specifically test the role of each RAAS component to understand detailed mechanisms linking Ang II receptor signaling to critical biological outcomes such as insulin secretion, apoptosis and other beta cell functions.

Several clinical trials with ACEi have shown reduced incidence of T2D in hypertensive patients.^{16,42-44} A couple of these studies have demonstrated that these beneficial effects were partly through improved insulin release.¹⁵ Additionally, increased ER stress reduces insulin biosynthesis.⁴⁵ Corroborating these findings, our data reported for the first time that ACEi inhibitors reduced ER stress and improved insulin secretion. The beneficial effects of ACEi could be potentially mediated through Mas receptor,⁴⁶ and worthy of future investigation.

We showed that Ang II treatment reduces insulin secretion, in agreement with few published studies.^{5,30,38} However, other studies have shown that activation of the RAAS pathway induced insulin secretion. This difference might be related to treatment conditions, such as culture media. Some insulin secretion experiments were conducted directly in media and did not use a starvation buffer (HBSS), which is typically used to measure glucose-stimulated insulin secretion.^{33,47} Also, the duration of Ang II stimulation to measure insulin secretion varied among studies. Ang II treatments could induce insulin secretion for shorter periods but could be deleterious at longer intervals. Similar effects were observed with IL-1 β . Clonal beta cells treated for a short time with IL-1 β potentiated insulin secretion, while longer treatment times inhibited insulin secretion.^{48,49} Consistent with these effects, we have previously shown that chronic RAAS overexpression in adipose tissue causes hyperinsulinemia and insulin resistance with increased adipocyte inflammation,¹⁰ while acute treatment of adipocytes with Ang II stimulated insulin signaling.⁵⁰

ER stress is also known to activate inflammation, and both of these pathological factors could be induced by Ang II.^{4,51,52} However, the precise inflammatory pathways which are induced by Ang II are not well established yet in beta cells as well in other tissues.^{53,54} Our study and others have demonstrated that activation of RAAS could contribute to inflammation.²¹ Furthermore, increased *Chop* expression, an ER stress marker, induced inflammation through the NF κ B pathway.⁵⁵ Additionally, proinflammatory mediators under pathological conditions also trigger ER stress, further suggesting these processes are intertwined.⁵⁶ Higher ER stress and inflammation, individually and in combination, could, in turn, cause insulin resistance, which could further activate RAAS leading to a vicious cycle.

Another well-established mechanism through which cellular processes are regulated is through miRNAs, also known to play a critical role in ER stress response.⁵⁷ In our study, miR-30a-3p was reduced by Ang II, consistent with studies in cardiac cells and adipocytes^{21,28} where Ang II reduced these miRNAs. However, whether Chop is a direct target of miR-30a-3p needs further evaluation using mimic/inhibitors and luciferase reporter assays.

Current study used 10–100 nM concentration to test Ang II effects in beta cells. This is much higher than 5–50 pM circulating levels found in healthy individuals.^{58,59} However, the levels could rise to ~200 pM in individuals suffering from obesity.⁶⁰ For example, subcutaneous adipose tissue Ang II levels are significantly higher than circulatory levels,⁶⁰ Ang II levels found locally in the canine pancreas is much higher than systemic levels.⁶¹ Ang II has a dramatic effect on insulin secretion, and such effects likely stem from local Ang II production. It is important to note that it is challenging to measure the amount of Ang II accurately due to its short life and low concentrations. However, studies have demonstrated that local RAAS generates high concentrations of Ang II in isolated human pancreatic islets.³⁰ It is also difficult to measure Ang II concentration directly in the human pancreas; hence it needs to be done in vitro using human or mouse islets.

In this original research, we primarily focused on the ATF4-CHOP pathway given its importance in beta cells.⁶² We also tested BiP, a key regulator of UPRs, but Ang II did not induce BiP at the doses tested (data not shown). It is possible that different doses of Ang II alter different ER stress biomarkers. Also, consistent with our findings in beta cells, Ang II induced ER stress in adipocytes by activating the ATF4-CHOP axis.²¹ However, studies have

also demonstrated that other players like ATF3, ATF6, and IRE1 α may be important as well in regulating beta cell proliferation.²⁷ Whether Ang II regulates other branches of ER stress in beta cells or related pathways needs to be tested in future studies. Lastly, we tested ER stress markers only at gene levels, but measuring protein levels or post translational modifications will provide a more comprehensive understanding, to better explain the role of Ang II overactivation in beta cell functions.

Conclusion

In conclusion, activation of the RAAS upregulates ER stress in clonal beta cells, which is detrimental in T2D especially, and could lead to reduced insulin secretion. Our studies demonstrated that inhibition of RAAS improves insulin secretion; thus, currently used antihypertensive drugs targeting RAAS are also promising targets for insulin resistance and T2D. Our studies add significantly to currently known benefits of RAAS inhibitors in T2D prevention, by unraveling a new mechanism in the pancreas, by which these inhibitors may improve T2D.

Abbreviations

Atf4, activating transcription factor; ACEi, angiotensin converting enzyme inhibitor; Ang II, angiotensin II; ARB, angiotensin receptor blockers; AT1, angiotensin type 1 receptor; AT2, angiotensin type 2 receptor; Agt, angiotensinogen; Chop, CCAAT-enhancer-binding protein homologous protein; ER, endoplasmic reticulum; ELISA enzyme-linked immunosorbent assay; iNos, inducible nitric oxide synthase; IL, Interleukin; Mcp-1, monocyte chemoattractant protein-1; Nf-kb, nuclear factor-kappa B; 4-PBA, 4-phenyl butyric acid; RT-qPCR, real-time quantitative polymerase chain reaction; RAAS, renin angiotensin aldosterone system; Tnf- α , tumor necrosis factor-alpha; T2D, type 2 diabetes mellitus; UPR, unfolded protein response.

Disclosure

The authors report no conflicts of interest in this work.

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