We previously reported that oxidative stress induced by long-term tacrolimus treatment impairs mitochondrial function in pancreatic beta cells. In this study, we aimed to investigate the therapeutic potential of coenzyme Q$_{10}$, which is known to be a powerful antioxidant, in mitochondrial dysfunction in tacrolimus-induced diabetic rats. In a rat model of tacrolimus-induced diabetes mellitus, coenzyme Q$_{10}$ treatment improved pancreatic beta cell function. The administration of coenzyme Q$_{10}$ improved insulin immunoreactivity within islets, which was accompanied by reductions in oxidative stress and apoptosis. Assessment of the mitochondrial ultrastructure by electron microscopy revealed that coenzyme Q$_{10}$ treatment increased the size, number, and volume of mitochondria, as well as the number of insulin granules compared with that induced by tacrolimus treatment alone. An in vitro study using a pancreatic beta cell line showed that tacrolimus treatment increased apoptosis and the production of mitochondrial reactive oxygen species, while cotreatment with coenzyme Q$_{10}$ effectively attenuated these alterations. At the subcellular level, tacrolimus-induced impairment of mitochondrial respiration was significantly improved by coenzyme Q$_{10}$, as evidenced by the increased mitochondrial oxygen consumption and ATP production. Our data indicate that coenzyme Q$_{10}$ plays an important role in reducing tacrolimus-induced oxidative stress and protects the mitochondria in pancreatic beta cells. These findings suggest that supplementation with coenzyme Q$_{10}$ has beneficial effects in tacrolimus-induced diabetes mellitus.

Tacrolimus (Tac), a calcineurin inhibitor (CNI), is the most commonly used immunosuppressant. However, chronic treatment with Tac following kidney transplantation is associated with an increased incidence of new-onset diabetes mellitus (DM) after transplantation (NODAT). A 5-year follow-up study of patients being treated with Tac after transplantation showed a frequency of DM of 41%.$^{2}$ The diabetogenic action of Tac is not understood, but pharmacologic calcineurin inhibition, important in preventing rejection during organ transplantation, is related to post-transplantation beta cell failure.$^{3-10}$ Various studies have revealed that this condition is partly related to the toxicity of Tac on beta cells in the pancreas, with oxidative injury playing a central role in Tac-induced beta cell dysfunction.$^{11-13}$

Coenzyme Q$_{10}$ (CoQ$_{10}$) acts as a diffusible electron carrier in the mitochondrial respiratory chain.$^{14}$ CoQ$_{10}$ also acts as a powerful antioxidant that removes free radicals, inhibits the initiation and propagation of lipid peroxidation in cellular biomembranes, and aids in the regeneration of $\alpha$-tocopherol.$^{15,16}$ Earlier papers have revealed the beneficial effects of CoQ$_{10}$ in diverse experimental diseases of oxidative tissues.$^{17-23}$ Therefore, CoQ$_{10}$ has the potential to preserve cells against Tac-induced oxidative stress and pancreatic beta cell injury.

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Received: 12 September 2018
Accepted: 17 May 2019
Published online: 29 May 2019
Recent studies have shown that Tac has a detrimental effect on mitochondrial function in beta cells. Impaired mitochondrial Ca\(^{2+}\) uptake, which is mechanistically linked to mitochondrial dysfunction, has been shown to occur following a reduction in the enzymatic activity of cytochrome c oxidase, the incidence of which is increased in DM following Tac treatment\(^{24}\). Based on this knowledge, the current study was conducted to investigate whether the concurrent treatment of CoQ\(_{10}\) can ameliorate Tac-induced DM by reducing oxidative stress via its anti-oxidative properties. We also evaluated whether the protective effect of CoQ\(_{10}\) is associated with the preservation of mitochondrial function, such as preventing impairments in oxygen consumption of mitochondria and ATP production and an increase in reactive oxygen species (ROS) production during Tac treatment.

**Materials and Methods**

**Ethics statement.** All procedures were performed in strict accordance with the recommendations of the ethical guidelines for animal studies. All experimental animal care protocols were approved by the Animal Care and Use Committee of the Catholic University of Korea (CUMC-2015-0028-02). Animals were sacrificed under xylazine/rompun anesthesia, and every effort was made to minimize animal suffering.

**Experimental rat model.** Male Sprague-Dawley (SD) rats (Orient Bio, Seongnamsi, Korea) were provided a 0.05% sodium diet (Research Diets, New Brunswick, NJ, USA), and water ad libitum. After 7 days, the mice were randomly divided into 4 groups (\(n = 8\) per group) and administered 1.5 mg/(kg\(\cdot\)day) Tac (Astellas, Ibaraki, Japan) or 1 mL/(kg\(\cdot\)day, subcutaneous) vehicle (Vh: olive oil; Sigma-Aldrich, St. Louis, MO, USA) with/without coenzyme Q\(_{10}\) (CoQ\(_{10}\): 20 mg/kg\(\cdot\)day, diluted in olive oil, daily oral gavage; Chong Kun Dang Pharm., Seoul, Korea) for 4 weeks. Route of administration and doses were selected as per those established in earlier reports\(^{25,26}\).

**Basic treatment protocol.** Animals were pair-feeding and their body weight was monitored. Rats were housed separately in metabolic cages (Tecniplast, Gazzada, Italy) for assessment of water intake and urine volume for 1 day after the 4-week treatment. On the next day, rats were anesthetized, and their blood and tissue were harvested for the following experiments. The Tac concentration in whole-blood was measured using liquid chromatography-tandem mass spectrometry (Abbott Diagnostics, Abbott Park, IL, USA)\(^{27}\).

**Pancreatic function and beta cell area in islets.** After the 4-week treatment, an intraperitoneal glucose tolerance test (IPGTT) was conducted for measurement of the glucose level. The data obtained in the IPGTT was calculated by trapezoidal estimation for the area under the curve of glucose (AUCg). The fasting insulin level in serum was examined using an ELISA Kit (Millipore Corp., St. Charles, MO, USA). HbA1c was measured in whole blood using Cobas Integra 800 (Roche Instrument Center AG, Rotkreuz, Switzerland). For quantification of the beta cell area, the numbers of beta cells were assessed in each sample using captured images following immunohistochemical staining for insulin (TDI Scope Eye Version 3.6 for Windows; Seoul, Korea). Insulin-positive cells were randomly selected and counted by randomly selecting 20 islets from the eight animals in each group.

**Immunohistochemistry.** After processing with the retrieval solution (pH 6.0), methanolic H\(_2\)O\(_2\), 0.5% of Triton X-100, and 10% of normal serum (Jackson ImmunoResearch, West Grove, PA, USA), the tissue sections were incubated for 1 day at 4°C with primary antibodies against insulin (18-0067, Invitrogen, Carlsbad, CA, USA), 8-hydroxy-2’-deoxyguanosine (8-OHdG: MOG-100P; JaICA, Shizuoka, Japan), 4-hydroxy hexenal (4-HHE: MHH-030n; JaICA), or Ki67 (ab15580, Abcam, Cambridge, UK) and then with peroxidase-conjugated antibodies (Molecular Probes, Carlsbad, CA, USA). The activity of peroxidase was evaluated using (4-HHE: MHH-030n; JaICA), or Ki67 (ab15580, Abcam, Cambridge, UK) and then with peroxidase-conjugated antibodies (Molecular Probes, Carlsbad, CA, USA). The number of 8-OHdG- or 4-HHE-labeled cells were determined by measuring approximately 20 randomly chosen non-overlapping islets from the eight animals in each group. Using histogram equalization, the percent positive area was calculated by quantitative analysis under the same intensity (TDI Scope Eye).

**Measurement of beta cell mass.** The relative volumes of beta cells were estimated by the point-counting method\(^{28}\). The relative beta cell volume was measured by classifying the number of points matching to the insulin-positive region by the number of points matching to the remaining pancreatic area. Cell masses were measured by multiplying the ratios of beta cells by the total pancreatic weight\(^{29}\).

**Glucose-stimulated insulin secretion (GSIS) assay.** As described earlier, islets were isolated from male SD rats (250–300 g) by digestion using collagenase\(^{30,31}\). The islets were pre-incubated in conditioned RPMI 1640 medium at 37°C for 24 h. The isolated islets were then treated with Tac (1 μg/mL) and CoQ\(_{10}\) (10–1000 ng/mL) for 12 h. This was followed by analysis of insulin secretion. The harvested islets were divided into groups of 30 islets and washed with Krebs-Ringer Modified Buffer (KRB), followed by addition of 2.8 mM glucose (basal). After washing with KRB, the islets were incubated with KRB containing 16.7 mM glucose for 1 h. The insulin level in the solution was examined using an ELISA kit (Millipore Corp., St. Charles, MO, USA).

**Measurement of 8-OHdG in serum.** The end product of oxidative DNA damage was measured by determining the concentration of 8-OHdG in serum. The 8-OHdG level was evaluated using an ELISA kit (Cell Biolabs, San Diego, CA, USA).

**Terminal deoxynucleotidyl transferase dUTP nick end labeling (TUNEL) assay.** The TUNEL assay was performed for the tissue sections using the Apoptosis Detection kit (Millipore Corp.) as per the
manufacturer's instructions. For double labeling with insulin, tissue sections were incubated with an insulin antibody (Invitrogen) followed by incubation with a Cy3-labeled antibody (Jackson ImmunoResearch). The double-labeled cells were counted in approximately 20 randomly selected non-overlapping islets per animal of each group.

Transmission electron microscopy. Electron microscopic inspection was processed as previously described32. The area and number of mitochondria per cell were evaluated in 40 random beta cells using an image analyzer (TDI Scope Eye).

Three-dimensional (3D) reconstruction of mitochondria. Long ribbons up to about 20 sections of isolated islets treated with drugs were cut at a thickness of 70–90 nm on an ultramicrotome (Leica Microsystems Ltd.). The area of interest was selected, and consecutive serial sections were imaged under an electron microscope (JEM 1010; JEOL, Tokyo, Japan). Outlines of individual mitochondria were manually traced with different colors through image stacks using the Photoshop software (Adobe Systems, San Jose, CA, USA). Each image was aligned, and the marked structures were masked and exported to the 3D modeling program Mimics v.19.0 (Materialise, Leuven, Belgium), along with information regarding slice thickness, actual pixel size, and image orientation. The mitochondrial volume and surface area in a stack consisting of equal numbers of sections were also calculated using Mimics v.19.0.

Cell culture. INS-1 (rat insulinoma cell line) was grown in conditioned RPMI-1640 medium in a humidified atmosphere containing 5% CO2. Cells were plated in culture dishes and treated with Tac (50 μg/ml, diluted in DMSO) and CoQ10 (1 pg/ml–10 μg/ml, diluted in DMSO) for 12 h.

Cell viability and apoptosis. For assessment of cell viability, INS-1 cells were plated in culture dishes at 90% confluence. On the following day, the cells were treated with Tac (50 μg/ml) and 1 pg/ml–10 μg/ml CoQ10 for 12 h. CCK-8 solution (Dojindo, Rockville, MD, USA) was added to each well to evaluate cell viability. For assessment of apoptosis, the cells detached by trypsin were treated with annexin V (BD Biosciences, San Jose,
CA, USA) in binding buffer (BD Biosciences). The cells were analyzed using a FACSCalibur flow cytometer (BD Biosciences). Data was reported as the percentage of fluorescent cells compared to the total cell number.

Measurement of mitochondrial ROS. The cells were treated with MitoSOX Red (Invitrogen) for 30 min at 37 °C to detect mitochondrial ROS (superoxide anion) levels using a FACSCalibur flow cytometer (BD Biosciences). Side and forward scatter data were obtained (10,000 events per sample).

Live cell time-lapse imaging. INS-1 cells were cultured in a 12-well plate and then incubated with 1μM of MitoSOX Red (Invitrogen) and drugs. Automated live-cell imaging was performed on a Lionheart FX Automated Microscope (BioTek Instruments Inc., Winooski, VT, USA) with attached plate handler and live-cell incubator using a 10× objective. Fluorescence images were captured every 1 h for 12 h, with optical emission and excitation wavelengths of 580 nm and 510 nm, respectively.
Oxygen consumption rate (OCR) experiments. Real-time OCR in the cells was evaluated using an XF24 Extracellular Flux Analyzer (Seahorse, Billerica, MA, USA). In a non-CO₂ incubator, the medium of the cells was changed to a running medium following incubation with drugs or Vh and treated for 1 h at 37 °C. The mitochondrial inhibitors used were 1 µM of oligomycin (ATP synthase inhibitor), 0.5 µM of carbonyl cyanide-4 (trifluoromethoxy) phenylhydrazone (FCCP), and 0.5 µM of rotenone/antimycin A (complex I and III inhibitor). Immediately following completion of the assay, each well was washed with phosphate-buffered saline, and lysis buffer (Pierce Biotechnology, Rockford, IL, USA) was added to lyse the cells in the well. Total protein was measured using the BCA Protein Assay Kit (Pierce Biotechnology). Then, the OCR data were normalized to the total protein content of each sample. Parameters for mitochondrial function were evaluated by measuring these mitochondrial inhibitor compounds as modulators to determine ATP production, basal respiration, spare respiratory capacity, and maximal respiration. A total of 3–4 wells were used for each group.

Statistical analysis. The values are expressed as the mean ± standard error of at least three independent tests. Using PRISM software, multiple comparisons were carried out by one-way analysis of variance with Bonferroni’s post-hoc test (Version 7.03 for Windows, GraphPad Software, La Jolla, CA, USA). Results with P values < 0.05 were considered statistically significant.

Results
Effect of CoQ₁₀ administration in an experimental model of Tac-induced DM. Table 1 displays alterations in the functional parameters of the experimental groups following treatment of Tac and CoQ₁₀ for 28 days. The Tac + CoQ₁₀ group showed larger reductions in intake-water and urine volume than that in the Tac group. Therefore, CoQ₁₀ did not affect the whole-blood trough level of Tac, implying that a drug-interaction did not occur at these doses.

Rats treated with Tac showed features of DM, as indicated by higher AUC₉ and HbA₁c values and lower levels of plasma insulin compared with those in animals administered with Vh. Addition of CoQ₁₀ reversed these changes (Fig. 1). Tac-treated islets observed lower insulin immunoreactivity and reduced numbers of beta cells than the Vh group, and these effects were reduced by treatment with CoQ₁₀ (Fig. 2). We also quantified the beta cell mass in the groups and found that a reduction in beta cell mass induced by Tac treatment was reversed by cotreatment with CoQ₁₀ as shown in Fig. 2c.

To assess the exact effects of the insulin secretion function of CoQ₁₀ during Tac-induced islet injury, we treated primary rat isolated islets with Tac and CoQ₁₀ in a culture setting and evaluated GSIS. As expected, Tac significantly decreased GSIS. The addition of CoQ₁₀ resulted in significantly higher insulin secretion levels compared with that of Tac alone, as shown in Fig. 3.
Effect of CoQ10 administration on Tac-induced oxidative stress and apoptosis in pancreatic beta cells.

The proliferation of beta cells was confirmed by tissue expression of Ki67 from the experimental groups (Fig. 4). Treatment of Tac reduced the number of Ki67-positive cells in the insulin-positive area compared with those in the Vh and CoQ10 groups, while concomitant CoQ10 treatment restored Ki67 staining. Figure 5 shows the co-immunohistochemical staining results for 8-OHdG (a–d,m) and 4-HHE (e–h,o) and TUNEL assays (i–p) using tissue sections from rats. (n) Serum 8-OHdG excretion.

**Figure 5.** Effects of CoQ10 on oxidative stress and apoptosis in pancreatic islets and sera during Tac-induced DM in rats. Representative images and quantification of immunohistochemistry for 8-OHdG (a–d,m) and 4-HHE (e–h,o) and TUNEL assays (i–p) using tissue sections from rats. (n) Serum 8-OHdG excretion. Arrowheads indicate 8-OHdG-, 4-HHE-, and TUNEL-positive cells. Scale bar = 50 µm (all panels). Data are presented as the mean ± SE (n = 8). One-way ANOVA was used to analyze the data. *P < 0.05 versus the Vh group; $P < 0.05 versus the CoQ10 groups; #P < 0.05 versus the Tac group.
restored by CoQ₁₀ treatment. We also examined whether CoQ₁₀ protects against apoptosis by Tac treatment. The number of TUNEL-positive cells were markedly higher in the Tac group than in the Vh group and were decreased with the cotreatment of CoQ₁₀ (Fig. 5i–l,p).

Effect of CoQ₁₀ administration on Tac-induced mitochondrial ultrastructural changes in pancreatic beta cells. To evaluate mitochondrial function, electron microscopy was used to assess the mitochondrial ultrastructure and quantify the mitochondrial number. For the islet cells obtained from the current experimental rats, Tac treatment decreased both the average mitochondrial area and number (Fig. 6a). By contrast, the administration of CoQ₁₀ recovered the average mitochondrial area and number. Electron microscopy also demonstrated that the number of insulin granules were markedly decreased by Tac treatment. CoQ₁₀ was effective in reducing this attenuation in number of granules, confirming that treatment with CoQ₁₀ protected islet function after Tac treatment (Fig. 6b–d). The effect of CoQ₁₀ on mitochondrial volume and surface area was evaluated by performing 3D reconstruction of consecutive sections of individual mitochondria in beta cells from isolated rat islets. Consistent with the previous results, reductions in mitochondrial volume and surface area induced by Tac treatment were significantly reversed by cotreatment with CoQ₁₀ (Fig. 6e,f).
CoQ10 reduces Tac-induced apoptosis and mitochondrial ROS production in INS-1 cells. Using the INS-1 pancreatic beta cell line, we further evaluated the protective effect of CoQ10 against mitochondrial injury during Tac treatment. INS-1 cells were subjected to different concentrations (1 pg/ml–10 µg/ml) of CoQ10 for 12 h during Tac-induced toxicity. CoQ10 improved INS-1 cell viability at all doses compared with that following Tac-only treatment (Fig. 7a). To examine whether CoQ10 inhibits mitochondrial pathways related to apoptosis, we carried out flow cytometric analysis with annexin V staining. The Tac-induced increase in the percentage of annexin V-positive cells was significantly attenuated by CoQ10 treatment (Fig. 7b,c).

Next, we evaluated whether treatment with CoQ10 reduced mitochondrial ROS accumulation during Tac treatment. Use of MitoSOX Red for mitochondrial superoxide (O2−) detection accompanied by flow cytometric analysis in INS-1 cells revealed that CoQ10 significantly attenuated Tac-induced MitoSOX Red fluorescence (Fig. 8a,b). To visualize the protective effect of CoQ10 against O2− accumulation in mitochondria, we performed time-lapse monitoring of MitoSOX Red intensity in the presence or absence of Tac and CoQ10 in the same field. As shown in Fig. 8c, MitoSOX Red fluorescence intensity was dramatically elevated in the Tac group, but this effect was attenuated by CoQ10 treatment.

CoQ10 increased mitochondrial respiration in Tac-treated INS-1 cells. The mitochondrial bioenergetics of whole cells was determined by calculating oxygen consumption over time after the sequential addition of inhibitors of mitochondrial function. This analysis revealed marked differences between the Tac and Tac + CoQ10 groups (Fig. 9a). Compared to Tac-only treatment, Tac + CoQ10 treatment resulted in higher rates of basal mitochondrial respiration, which is usually indicative of either a higher number of mitochondria or increased mitochondrial activity (Fig. 7b). Moreover, compared with Tac-only treatment, Tac + CoQ10 treatment resulted in significantly higher ATP-linked respiration and maximal respiration.

Discussion
The current study was performed to investigate whether cotreatment with CoQ10 is effective in ameliorating pancreatic beta cell dysfunction by Tac. The results of our study showed that CoQ10 attenuated hyperglycemia and restored the insulin secretion ability by reducing Tac-induced oxidative stress. These findings suggest that CoQ10 produces beneficial effects for reducing mitochondrial injury via its antioxidative properties during Tac-induced beta cell injury (Fig. 10). The results of our study thus provide a rationale for use of CoQ10 as supplemental therapy in Tac-induced DM in clinical practice.
In this study, we first evaluated whether the administration of CoQ10 was effective in controlling hyperglycemia in an experimental Tac-induced DM model. Recently, several clinical studies have shown that CoQ10 treatment reduces DM34–36. In this study, we found that the AUCg based on IPGTT was increased and the plasma insulin level and insulin-immunoreactivity in islets were decreased by chronic Tac treatment. However, cotreatment with CoQ10 significantly decreased the AUCg and increased levels of plasma insulin and GSIS compared with the levels observed following Tac treatment alone. Furthermore, immunostaining of insulin in islets revealed that concomitant treatment with CoQ10 resulted in increases in islet size and immunoreactivity for insulin, as well as preservation of islet morphology as demonstrated by less irregular islet boundaries and reduced vacuolization. These findings suggest that CoQ10 has beneficial effects on the preservation of beta cells during Tac treatment.

To elucidate the possible mechanism of CoQ10 in the control of hyperglycemia, we evaluated the expression of markers of oxidative stress in the experimental groups, as oxidative stress has been suggested to be an important pathway in the pathogenesis of pancreatic beta cell injury. High levels of oxidative stress induced by Tac administration lead to islet cell death and dysfunction, with short- and long-term Tac treatment inducing the

**Figure 8.** Effects of CoQ10 during Tac-induced mitochondrial ROS production in INS-1 cells. (a,b) MitoSOX Red staining to detect mitochondrial superoxide anion (O$_2^-$) levels by flow cytometric analysis. (c) Representative time-lapse images of cells treated for 12 h with MitoSOX Red in the presence or absence of Tac and CoQ10 were taken from the same field. Scale bar = 1000 µm (all panels). Data are presented as the mean ± SE (n = 5). One-way ANOVA was used to analyze the data. *P < 0.05 versus the control group; ^P < 0.05 versus the CoQ10 groups; #P < 0.05 versus the Tac group.
production of ROS, thereby causing apoptotic cell death. Therefore, it has been proposed that a reduction in oxidative stress could protect from pancreatic beta cell injury. Based on this hypothesis, we focused on the antioxidative effect of CoQ₁₀ on Tac-induced oxidative injury in pancreatic beta cells. For this, we measured the expression of 8-OHdG and 4-HHE, markers of oxidative DNA and lipid damage, respectively, and found that while their expression levels were increased in beta cells in Tac-treated mice, administration of CoQ₁₀ attenuated these increases in expression levels. Indeed, there is evidence to show that the development of diabetes is

Figure 9. Effect of CoQ₁₀ on mitochondrial function during Tac-induced injury in INS-1 cells. (a) The areas under the curve for basal respiration, (b) ATP production, maximal respiration, proton leak, and non-mitochondrial respiration were calculated from the OCR. Data are presented as the mean ± SE (n = 5). One-way ANOVA was used to analyze the data. &P < 0.05 versus the control group; $P < 0.05 versus the CoQ₁₀ groups; #P < 0.05 versus the Tac group.

Figure 10. Proposed mechanism of the protective effect of CoQ₁₀ during Tac-induced pancreatic beta cell injury.
associated with increased oxidative stress and that CoQ₁₀ can scavenge ROS, resulting in an anti-hyperglycemic effect[22]. Overall, our data indicate that oral administration of CoQ₁₀ is effective in reducing Tac-induced oxidative stress in pancreatic beta cells.

We further evaluated whether the antioxidative effect of CoQ₁₀ ameliorated mitochondrial injury during Tac treatment. The mitochondria play important roles in pancreatic beta cells from glucose metabolism to insulin exocytosis, thereby ensuring tight regulation of glucose-stimulated insulin secretion[23]. Impairment in the mitochondrial function affects this metabolic coupling and ultimately leads to apoptosis and beta cell death. Furthermore, mitochondria are susceptible to oxidative damage and could be a major source of superoxide under disease conditions[24]. Based on this knowledge, we first examined rates of apoptosis, a mitochondrial pathway of cell death in vivo and in vitro, and found that CoQ₁₀ reduced the numbers of TUNEL- and annexin V-positive cells compared with those in the Tac group. Ultrastructural analysis revealed that Tac treatment caused reductions in the number, area, size, and volume of mitochondria, as well as the number of insulin granules, while CoQ₁₀ administration attenuated these changes. CoQ₁₀ also reduced accumulation of mitochondrial ROS and superoxide anion and restored basal respiration, ATP-linked respiration, and maximal respiration rates according to oxygen consumption over time. These data suggest that administration of CoQ₁₀ helps to maintain mitochondrial function during Tac-induced oxidative stress, resulting in a subsequent decrease in apoptotic cell death.

In addition to Tac, rapamycin analogs, sirolimus (SRL) and everolimus (EVR), have been widely used as immunosuppressants in transplantation, but they are also associated with an increased risk of DM. Many studies have demonstrated that both of these drugs cause mitochondrial dysfunction in pancreatic beta cells, eventually leading to decreased insulin release[25–27]. We have also confirmed that SRL or EVR treatment alone results in the development of hyperglycemia, accompanied by a reduction in the number of insulin granules and an increase in the expression of oxidative stress markers, in animal models[28]. However, addition of CoQ₁₀ attenuated SRL-induced hyperglycemia, as well as oxidative stress, in a rat model. At the subcellular level, CoQ₁₀ also improved not only the morphology of the mitochondria but also mitochondrial respiration[29]. Based on these findings, CoQ₁₀ supplementation would also be beneficial in the treatment of rapamycin-induced DM.

In conclusion, CoQ₁₀ plays an important role in reducing Tac-induced oxidative stress and protecting mitochondria in pancreatic beta cells. These findings suggest that CoQ₁₀ may be useful in the management of Tac-induced DM in the future.

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**Acknowledgements**
This work was supported by the Korean Health Technology R&D Project, Ministry for Health & Welfare, Republic of Korea (HI14C3417, HI16C1641), and Basic Science Research Program through the National Research Foundation of Korea (NRF) funded by the Ministry of Science, ICT & Future Planning (NRF-2018R1D1A1A02043014).

**Author Contributions**
S.W.L. and C.W.Y. designed the research and wrote the report; K.L., Y.J.S. and Y.Q. conducted the animal experiments; K.E.L. and H.L.K. conducted the electron microscopic analysis; S.W.L., J.H.Y. and E.J.K. performed experiments; K.E.L. and H.L.K. conducted the electron microscopic analysis; S.W.L. and C.W.Y. designed the research and wrote the report; K.L., Y.J.S. and Y.Q. conducted the animal experiments; K.E.L. and H.L.K. performed the cell culture studies; K.L., S.W.L., B.H.C. and C.W.Y. analyzed the data and edited the manuscript.

**Additional Information**

**Competing Interests:** The authors declare no competing interests.

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