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Beta-Amyloid Instigates Dysfunction of Mitochondria in Cardiac Cells

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Abstract: Alzheimer's disease (AD) includes the formation of extracellular deposits comprising aggregated β -amyloid ($A\beta$) fibers associated with oxidative stress, inflammation, mitochondrial abnormalities, and neuronal loss. There is an associative link between AD and cardiac diseases; however, the mechanisms underlying the potential role of AD, particularly $A\beta$ in cardiac cells, remain unknown. Here, we investigated the role of mitochondria in mediating the effects of $A\beta_{1-40}$ and $A\beta_{1-42}$ in cultured cardiomyocytes and primary coronary endothelial cells. Our results demonstrated that $A\beta_{1-40}$ and $A\beta_{1-42}$ are differently accumulated in cardiomyocytes and coronary endothelial cells. $A\beta_{1-42}$ had more adverse effects than $A\beta_{1-40}$ on cell viability and mitochondrial function in both types of cells. Mitochondrial and cellular ROS were significantly increased, whereas mitochondrial membrane potential and calcium retention capacity decreased in both types of cells in response to $A\beta_{1-42}$. Mitochondrial dysfunction induced by $A\beta$ was associated with apoptosis of the cells. The effects of $A\beta_{1-42}$ on mitochondria and cell death were more evident in coronary endothelial cells. In addition, $A\beta_{1-40}$ and $A\beta_{1-42}$ significantly increased Ca^{2+} -induced swelling in mitochondria isolated from the intact rat hearts. In conclusion, this study demonstrates the toxic effects of $A\beta$ on cell survival and mitochondria function in cardiac cells.

Keywords: Alzheimer's disease; beta-amyloid; cardiomyocytes; coronary artery endothelial cells; mitochondria



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1. Introduction

Alzheimer's disease (AD) is a progressive neurodegenerative disorder that is expected to affect 13 million individuals, mostly older adults, in the U.S. by 2050 [1]. The main pathological characteristics of AD are the formation of extracellular deposits comprising aggregated β -amyloid ($A\beta$) fibers and intracellular neurofibrillary tangles formed by hyperphosphorylated tau protein. These alterations are associated with the loss of synapses, mitochondrial structural and functional abnormalities, oxidative stress, inflammation, and neuronal loss. A growing body of experimental and clinical studies demonstrate an associative link between AD and cardiac diseases such as heart failure, ischemic heart disease, and atrial fibrillation [2]. It has been proposed that cardiac dysfunction leads to cerebral hypoperfusion (hypoxia) associated with brain oxidative stress and acidosis that, in combination with $A\beta$ aggregation, provoke neuronal degradation and progression of AD [3]. Oxidative stress can induce further production of $A\beta$ and tau protein, another critical component involved in the pathogenesis of AD [4]. These studies suggest a feedback loop embedded in the crosstalk between oxidative stress and $A\beta$ aggregation that stimulates the development and progression of AD. Although a causal role of cardiac dysfunction in AD progression can be explained, at least partially, by oxidative stress induced by decreased

cerebral blood flow (hypoperfusion), it remains unclear whether AD *per se* can increase the risk for developing cardiac abnormalities.

Protein misfolding plays a central role in the pathogenesis of AD, and recent studies identified A β aggregates in the heart of patients with heart failure and dilated cardiomyopathy [5–7]. It has been suggested that A β oligomers may compromise cardiac function and lead to cardiovascular disease, the second leading cause of death (after pneumonia) in patients with AD [8]. A β_{1-40} and A β_{1-42} , structurally similar to those found in the brain and similar amounts in the heart of AD patients, are associated with diastolic cardiac dysfunction [6]. Moreover, AD shares similar risk factors and common genetic profiles with cardiac diseases [5,9]. Mutations in *PSEN1* and *PSEN2*, associated with familial AD, and the apolipoprotein E $\epsilon 4$ allele, a risk factor for AD, were found in patients with dilated cardiomyopathy [5,10]. AD patients lacking symptomatic cardiovascular diseases exhibit cardiac dysfunction, including increased diastolic dysfunction and left ventricular wall thickness [6,11–13], as well as electrocardiographic abnormalities [12]. In patients with AD, A β_{1-40} and A β_{1-42} aggregates were found in the cardiomyocytes and interstitial spaces associated with cardiac dysfunction [6]. Plaque-like amyloid deposits observed in cardiomyopathy [5] could induce proteotoxicity and cell death in the heart [14]. However, despite a broad range of studies, the causal role of AD and amyloidosis in the development of cardiac abnormalities remains unknown. In addition, the source of the A β protein aggregates in the hearts of patients with AD is not entirely understood. However, high levels of A β_{1-40} oligomers have been detected in the cerebrospinal fluid of AD patients [15], and an increase in A β oligomers in peripheral plasma is now starting to be considered as a potential biomarker for AD [16]. Therefore, there is a potential for these toxic aggregation species to reach the heart. Although it is not yet clear how AD affects cardiac function, patients with AD are at increased risk for developing cardiac abnormalities.

The mechanisms of cell damage induced by A β are still subject to investigation; several lines of evidence demonstrate toxic intracellular effects of A β and suggest a role of the mitochondria in mediating these effects (reviewed in [17–19]). A β was found in the mitochondria of patients with AD [4,20] and increased A β aggregation was associated with mitochondrial dysfunction [21]. Together with oxidative stress and altered Ca²⁺ homeostasis, mitochondrial abnormalities have been shown to mediate neuronal dysfunction and cell death in AD patients and animal models of A β and tau pathology [22,23]. Mitochondrial dysfunction also mediates cerebral microvascular endothelial cell death induced by A β oligomers [21,24,25]. Mitochondria-mediated apoptosis can also be initiated by the effects of A β oligomeric species on membrane death receptors, such as DR4 and DR5, which trigger the extrinsic, and consequently the intrinsic, apoptotic pathways [26]. However, whether mitochondrial alterations act as a causal event and contribute to the etiology of AD or result from the disease pathology remains unknown. Notably, most of the studies elucidating the effect of A β on mitochondria have been conducted on brain tissue and neuronal or other brain cells, and thus, the mechanisms of A β production and its possible effects on cardiac cells remain unknown.

In this study, we elucidated the role of mitochondria in mediating the effects of A β_{1-40} and A β_{1-42} on cultured cardiomyocytes, coronary endothelial cells, and isolated cardiac mitochondria to clarify the potential capability of A β to impact mitochondrial bioenergetics and induce cardiac cell death. Our results demonstrated that A β_{1-40} and A β_{1-42} differentially affected cell viability and mitochondrial function; A β , mostly A β_{1-42} was found aggregated and instigated mitochondrial dysfunction associated with increased mitochondrial ROS (mtROS) and swelling, and reduced mitochondrial membrane potential ($\Delta\Psi_m$), and calcium retention capacity (CRC).

2. Materials and Methods

2.1. Animals

Adult Sprague Dawley male rats (275–325 g) were purchased from Taconic (Hillside, NJ, USA). All experiments were performed according to protocols approved by the UPR

Medical Sciences Campus Institutional Animal Care and Use Committee and conformed to the National Research Council Guide for the Care and Use of Laboratory Animals published by the US National Institutes of Health (2011, eighth edition).

2.2. Isolation of Mitochondria from Rat Hearts

Mitochondria were isolated from rat hearts by the method described previously [27]. Briefly, heart ventricles were cut and homogenized using a Polytron homogenizer in ice-cold sucrose buffer containing 300 mM sucrose, 20 mM Tris-HCl, and 2 mM EGTA, pH 7.2, and supplemented with 0.05% BSA. The heart homogenate was centrifuged at $2000 \times g$ for 3 min to remove cell debris. The supernatant was centrifuged at $10,000 \times g$ for 6 min to precipitate mitochondria and then washed again under the same conditions in the BSA-free sucrose buffer. The final pellet containing mitochondria was resuspended in the sucrose buffer.

2.3. Mitochondrial Permeability Transition Pore (mPTP) Opening

The swelling of mitochondria as an indicator of mPTP opening was determined in freshly isolated mitochondria (50 μ g) by monitoring the decrease in light scattering at 525 nm in the presence or absence of Ca^{2+} [28]. The swelling buffer contained 125 mM KCl, 20 mM Tris-base, 2 mM KH_2PO_4 , 1 mM MgCl_2 , 1 μ M EGTA, 5 mM α -ketoglutarate, 5 mM L-malate, pH 7.1. Mitochondrial swelling was measured by adding Ca^{2+} to a total (accumulative) concentration of 100, 200, 300 μ M with 5-min intervals at 37 °C using the Clariostar (BMG Labtech, Cary, NC, USA). The rates of swelling were calculated as decrements of absorbance values per minute ($\Delta A_{525} \cdot \text{min}^{-1} \cdot \text{mg}^{-1}$) and presented as a percentage of control.

2.4. Primary Primary Human Coronary Artery Endothelial Cells (HCAEC)

Primary HCAEC were cultured according to the manufacturer's recommendations (ATCC). The cells were cultured in Vascular Cell Basal Media supplemented with VEGF-based Endothelial Cell Growth Kit (5 ng/mL rh VEGF, 5 ng/mL rh EGF, 5 ng/mL rh FGF, 15 ng/mL rh IGF-1, 10 mM L-glutamine, 0.75 units/mL heparin sulfate, 1 μ g/mL hydrocortisone, 50 μ g/mL ascorbic acid, 2% fetal bovine serum, ATCC) and 1% antibiotic solution (Sigma-Aldrich, Burlington, MA, USA). Cells were maintained in 95% air and 5% CO_2 at 37 °C. Cells from passage 3–8 and less than 14 divisions were used, and less than 80% confluence was maintained during propagation.

2.5. H9c2 Cardiomyoblasts

H9c2 cardiomyoblasts were cultured according to the manufacturer's recommendations (ATCC). Briefly, the cells were cultured in DMEM based modified media (4 mM L-glutamine, 4.5 g/L glucose, 1 mM sodium pyruvate, and 1.5 g/L sodium bicarbonate) supplemented with 10% fetal bovine serum and 1% antibiotic solution (Sigma-Aldrich) and maintained in 95% air and 5% CO_2 at 37 °C. Cells maintained within 80–90% confluence from passages 3–10 were used in experiments. Mitochondrial bioenergetics, metabolism, and morphology of H9c2 cells are similar to primary cardiomyocytes [29].

2.6. Permeabilization of Cells

Cells were freshly harvested using trypsin-EDTA and then permeabilized for 10 min on ice in the buffer containing 300 mM sucrose, 10 mM Tris-HCl, 2 mM EGTA, pH 7.4, and 50 μ g/mL saponin. After the permeabilization, cells were washed with equilibration buffer (100 mM sucrose, 10 mM Tris-HCl, 10 μ M EGTA, pH 7.4), then dissolved in the incubation buffer (200 mM sucrose, 10 mM Tris-MOPS, 5 mM α -ketoglutarate, 2 mM malate, 1 mM P_i , 10 μ M EGTA-Tris, pH 7.4).

2.7. Mitochondrial CRC Assay

Freshly harvested and permeabilized cells were incubated at 37 °C in the 0.1 mL of the incubation buffer (200 mM sucrose, 10 mM Tris-MOPS, 5 mM α -ketoglutarate, 2 mM malate, 1 mM Pi, 10 μ M EGTA-Tris, pH 7.4) containing 100 nM Calcium Green-5N. Calcium was added to increase matrix Ca^{2+} load, and the fluorescence intensity was recorded by the CLARIOStar microplate reader (BMG Labtech).

2.8. Cellular ROS Assay

Freshly harvested and permeabilized cells were incubated at 37 °C in the 0.1 mL of the incubation buffer (200 mM sucrose, 10 mM Tris-MOPS, 5 mM α -ketoglutarate, 2 mM malate, 1 mM Pi, 10 μ M EGTA-Tris, pH 7.4) containing 50 mM sodium phosphate, pH 7.4, 50 μ M AmplifluTM Red, 0.1 U/mL HRP (Sigma-Aldrich) and fluorescence signals were monitored using CLARIOStar microplate reader (BMG Labtech).

2.9. Fluorescence Immunocytochemistry

Cells were fixed and permeabilized with ice-cold methanol for 5 min. Fixed cells were washed with PBS 3 times, then blocked with 2% BSA in PBS for 30 min. Antibodies against A β (ab11132), ATP5A (ab176569), and calreticulin (ab92516) were used as per the manufacturer's recommendation (Abcam, Waltham, MA, USA). The cells were incubated with the primary antibodies overnight at 4 °C, and then washed 3 times with PBS and incubated with Alexa Fluor 488 anti-mouse and/or Alexa Fluor 594 anti-rabbit secondary antibodies (Thermo Fisher, Waltham, MA, USA) for 1 h at room temperature. After incubation with secondary antibodies, the cells were washed 3 times with PBS, and 100 nM DAPI was added to visualize the nucleus. Images were captured by Olympus IX73 microscope with LUCPLFLN40X objective using Cellsense Dimension (Olympus, Center Valley, PA, USA) software. Image compositions were made using ImageJ.

2.10. Oligomerization and the Treatments of A β Peptides

The A β peptides A β ₁₋₄₀ and A β ₁₋₄₂ purchased from Sigma-Aldrich and Bon Opus Biosciences (Millburn, NJ, USA) were prepared as described previously [30]. First, lyophilized amyloid peptides were dissolved in 1,1,1,3,3,3-hexafluoro-2-propanol (HFIP) to 1 mM. After the peptides were completely monomerized, they were lyophilized and re-dissolved in DMSO to 10 mM, then were diluted to 1 mM by adding deionized water. The peptides were diluted in culture media to 0.2 mM and incubated at 4 °C for 48 h to form oligomers, which were then applied to cultured cells to a final concentration of 10 μ M [31]. Low-binding surface plastic wares were used to prepare the amyloid peptides. To treat H9c2 cells, media was changed to serum-free media, A β peptides were added, and incubated 48–96 h. For primary HCAEC, Vascular Cell Basal Media supplemented with VEGF-based Endothelial Cell Growth Kit was used without modification. The HCAEC growth media was refreshed every 2 days in the presence or absence of A β peptides for 2–20 days.

2.11. Analysis of Cell Viability

Cell viability was determined by the AlamarBlueTM Cell Viability Assay Reagent (Thermo Fisher) as previously described [32].

2.12. Analysis of Cellular ATP, $\Delta\Psi_m$, and mtROS

Cells were live stained with 5 μ M ATP-Red (Sigma-Aldrich), 10 μ M JC-1 (Thermo Fisher), and 2 μ M MitoSOX (Thermo Fisher) for quantification of cellular ATP levels, $\Delta\Psi_m$, and mtROS production, respectively, following the manufacturers' recommendation (Thermo Fisher). The fluorescence intensity of the dyes was measured using the CLARIOStar microplate reader (BMG Labtech). Fluorescence signals of ATP-Red and MitoSOX were normalized to the fluorescence intensity (blue signal) of the nucleus (Hoechst). JC-1 signals were presented as the red to green fluorescence intensity ratio of the dye.

2.13. Apoptosis Assay

For analysis of activated caspases 3 and 7, live cells were stained with 2 μ M CellEvent Caspase-3/7 Green Detection Reagent (Thermo Fisher) in the presence of 50 nM Hoechst 33342 (Thermo Fisher) as per the manufacturer's recommendation. Fluorescence signals were measured using the CLARIOstar microplate reader (BMG Labtech). The caspase fluorescence intensity was normalized to the fluorescence intensity (blue signal) of the nucleus (Hoechst).

2.14. Statistical Analysis

Data were analyzed using Student's *t*-test. Results are presented as mean \pm SEM. $p < 0.05$ was considered statistically significant. The number of biological samples but not technical replicates were used as a sample size.

3. Results

3.1. A β Decreased Cell Viability and Impaired Cell Morphology

The cell viability of HCAEC grown in the culture media containing 10 μ M A β_{1-40} or A β_{1-42} for 20 days was significantly decreased in comparison with control cells. The cell viability of A β_{1-40} or A β_{1-42} groups were 8.6% and 24% lower ($p < 0.05$) than the control, respectively (Figure 1A). Morphological analysis using phase-contrast microscopy in A β_{1-42} -treated cells revealed irregular-shaped lumps appearing bright compared to the control and A β_{1-40} -treated cells (Figure 1B). Incubation of H9c2 cardiomyocytes with 10 μ M A β_{1-42} for 96 h resulted in a 39% ($p < 0.05$) decrease of cell viability, whereas A β_{1-40} reduced the cell viability only by 8% (Figure 1C). H9c2 cells challenged with A β_{1-42} showed abnormally shrinking cell morphology, in addition to abnormal bright aggregates (Figure 1D).

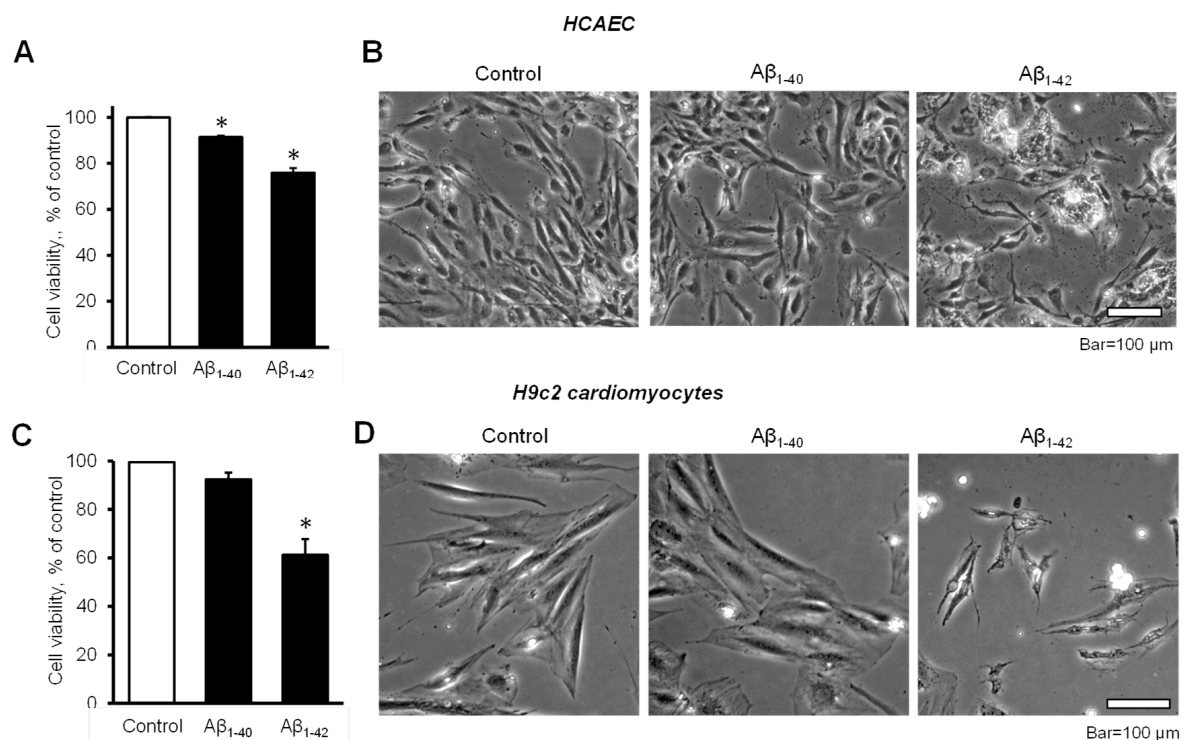


Figure 1. A β decreased cell viability and impaired cell morphology (A,B): Primary HCAEC grown for 20 days in the presence and absence of 10 μ M A β_{1-40} or A β_{1-42} . (A): Cell viability. $n = 4$ per group. * $p < 0.05$ vs. control. (B): Representative phase-contrast images of control (vehicle: DMSO) cells and cells grown with 10 μ M A β_{1-40} or A β_{1-42} . Bar = 100 μ m (C,D): H9c2 cardiomyocytes grown with A β_{1-40} or A β_{1-42} for 96 h. (C): Cell viability. $n = 4$ per group. * $p < 0.05$ vs. control. (D): Representative images of control (vehicle: DMSO) cells and cells grown with 10 μ M A β_{1-40} or A β_{1-42} . Bar = 100 μ m.

3.2. Aggregated A β Accumulated Inside of Cells

To investigate the possible inclusion of A β aggregates in subcellular compartments, primary HCAEC and H9c2 cardiomyocytes incubated with A β_{1-40} or A β_{1-42} were visualized using immunocytochemistry. Control primary HCAEC showed only low perinuclear signal, which could indicate a low level of A β and/or background staining, whereas A β_{1-40} -treated cells demonstrated the higher intensity of the cytoplasmic signal. Primary HCAEC challenged with A β_{1-42} showed a strong signal of irregular-shaped aggregates with a size of 1–50 μm (Figure 2A). Similar patterns of intracellular aggregation of A β were observed in H9c2 cells. The control group showed low perinuclear signal, whereas cells grown with A β_{1-40} showed more cytoplasmic signal (in addition to the nuclei) compared to the control. Cells grown with A β_{1-42} showed irregular-shaped aggregates in the cytoplasm (Figure 2B) that were absent in the control.

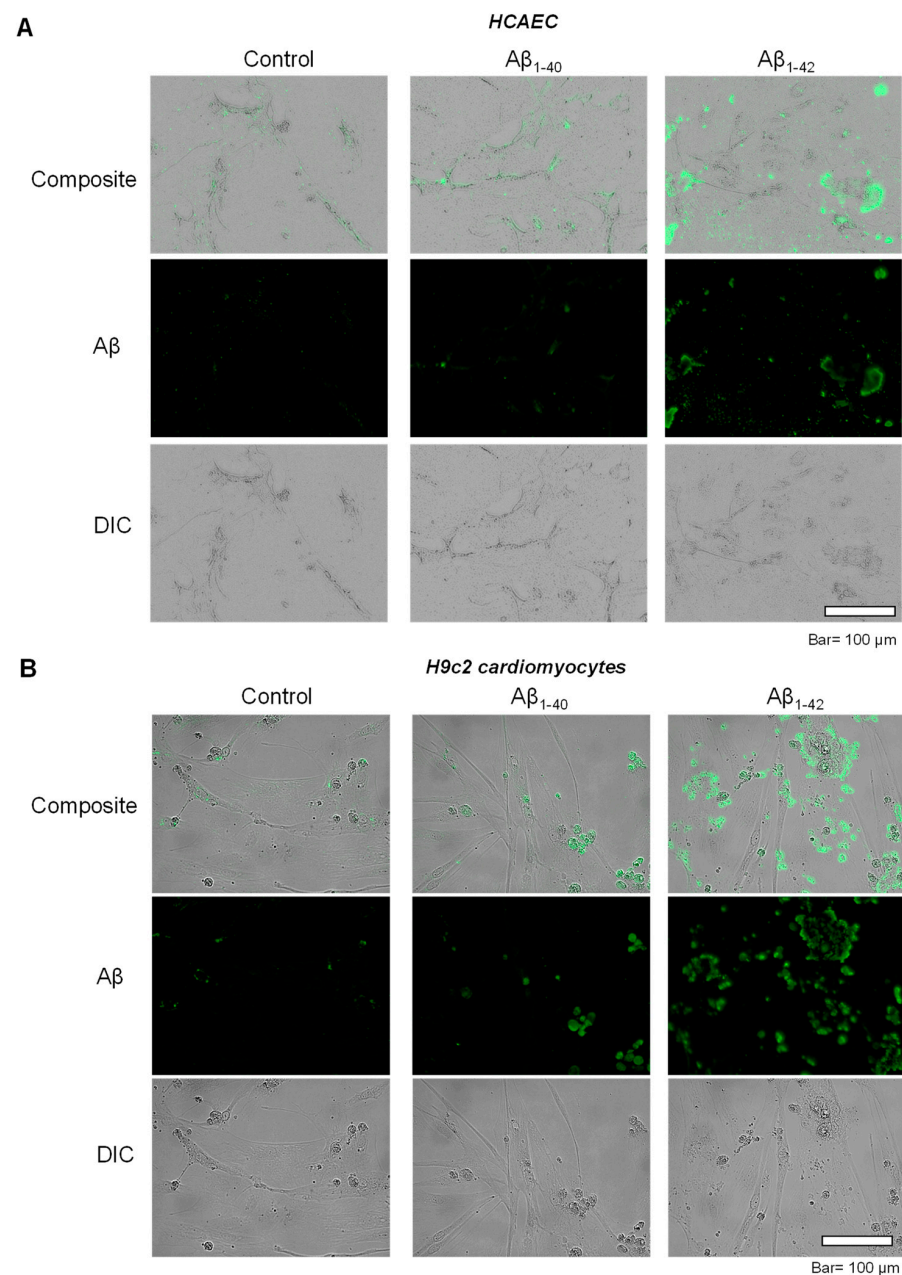


Figure 2. Accumulation of aggregated A β . (A): Immunostaining of A β in primary HCAEC grown with 10 μM A β_{1-40} or A β_{1-42} for 20 days. Bar= 100 μm (B): Immunostaining of A β in H9c2 cardiomyocytes grown with 10 μM A β_{1-40} or A β_{1-42} for 96 h. Bar = 100 μm .

3.3. A β Induced Mitochondrial and Cellular Dysfunction

Next, we investigated the role of mitochondria in the cells exposed to A β . The primary HCAEC incubated with 10 μ M A β_{1-42} for 20 days showed a 37% ($p < 0.05$) increase of mtROS, and a 179% ($p < 0.01$) increase of cellular (total) ROS, compared to the control cells (Figure 3A,B). A β_{1-40} and A β_{1-42} induced depolarization of the mitochondrial membrane as evidenced by decreased $\Delta\Psi_m$. The primary HCAEC treated with A β_{1-40} and A β_{1-42} showed a 20% ($p < 0.05$) and 41% ($p < 0.01$) decrease of $\Delta\Psi_m$, respectively, in comparison with the control (Figure 3C). A β_{1-42} induced a 2.4-fold ($p < 0.01$) increase of activated caspase 3/7, indicating increased apoptosis (Figure 3D). Interestingly, A β_{1-40} and A β_{1-42} increased ATP levels by 13% and 29% ($p < 0.01$), respectively (Figure 3E). Similar trends in cellular and mitochondrial parameters were observed in H9c2 cells. The cells incubated with 10 μ M A β_{1-42} for 20 days showed a 22% ($p < 0.01$) increase of mtROS, and a 24% ($p < 0.01$) increase of cellular ROS, compared to the control cells (Figure 3F,G). A β_{1-42} decreased $\Delta\Psi_m$ by 33% ($p < 0.05$) in H9c2 cells (Figure 3H), which showed 25% ($p < 0.01$) more activated caspase 3/7 than the control cells (Figure 3I). In H9c2 cells, only A β_{1-42} increased the ATP levels by 13% ($p < 0.05$), whereas A β_{1-40} had no significant effects (Figure 3J).

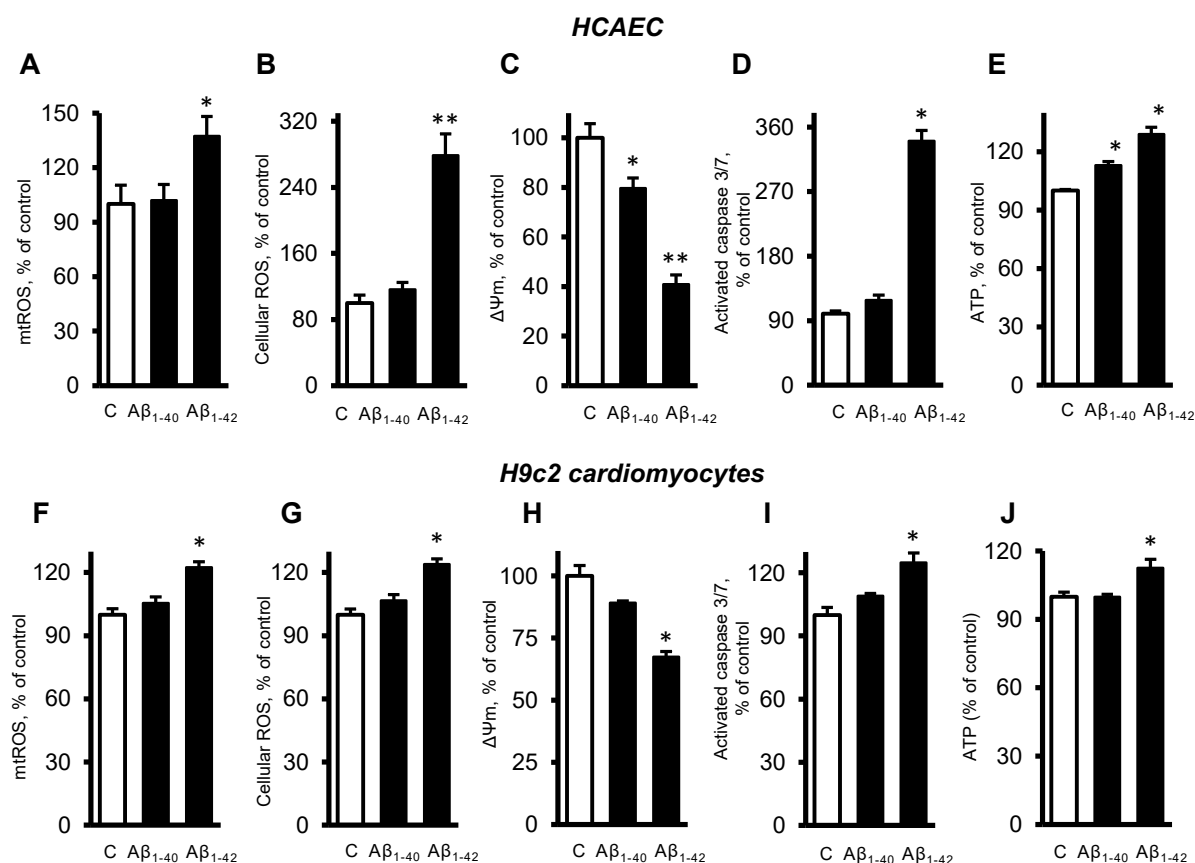


Figure 3. A β -induced mitochondrial and cellular dysfunction. (A–E): Primary HCAEC cultured for 20 days in the presence and absence of 10 μ M A β_{1-40} or A β_{1-42} . (A): mtROS levels measured by MitoSOX Red (Thermo Fisher). (B): Cellular ROS levels measured using Ampiflu Red (Sigma) in permeabilized cells. (C): $\Delta\Psi_m$ measured by JC-1 (Thermo Fisher). (D): Activated caspase 3/7 levels measured using the CellEvent™ Caspase-3/7 Green Detection Reagent (Thermo Fisher). (E): ATP levels measured by ATP-Red (Sigma). (F–J): H9c2 cardiomyocytes cultured for 96 h in the presence or absence of 10 μ M A β_{1-40} /A β_{1-42} . All parameters were measured by the same methods used for primary HCAEC (A–E). (F): mtROS levels. (G): Cellular ROS levels. (H): $\Delta\Psi_m$. (I): Activated caspase 3/7 levels. (J): ATP levels. $n = 4$ per group for all parameters of primary HCAEC and H9c2 cardiomyocytes. * $p < 0.05$, ** $p < 0.01$ vs. control (C).

Figure 4. Cells and mitochondria demonstrated early response to A β . (A–G): Primary HCAEC grown for 48 h in the presence and absence of 10 μ M A β ₁₋₄₀ or A β ₁₋₄₂. (A): Cell viability after 48 h of treatment. $n = 4$ per group. (B): Representative phase-contrast images of control (vehicle: DMSO) cells and cells treated with 10 μ M A β ₁₋₄₀ or A β ₁₋₄₂ for 48 h. Bar = 100 μ m (C): mtROS levels (D): Cellular ROS levels. (E): $\Delta\Psi_m$. (F): Activated caspase 3/7 levels. (G): ATP levels. (H–N): H9c2 cardiomyocytes grown for 48 h in the presence and absence of 10 μ M A β ₁₋₄₀ or A β ₁₋₄₂. (H): Cell viability after 48 h of treatment. (I): Representative phase-contrast images of cells. Bar = 100 μ m (J): mtROS levels (K): Cellular ROS levels. (L): $\Delta\Psi_m$. (M): Activated caspase 3/7 levels. (N): ATP levels. $n = 4$ per group for all parameters of primary HCAEC and H9c2 cardiomyocytes. * $p < 0.05$ vs. control (C).

Cardiomyocytes also showed dark, grainy aggregates in the cytoplasm of incubated with A β_{1-42} for 48 h. The control and A β_{1-40} -treated groups did not show remarkable changes in morphology (Figure 4I). A β_{1-42} induced a 17% ($p < 0.05$) increase of mtROS and a 19% ($p > 0.05$) increase of cellular ROS after 48 h of incubation (Figure 4J,K). A β_{1-42} induced a 15% ($p < 0.05$) decrease of the $\Delta\Psi_m$ (Figure 4L) and increased caspase 3/7 activation by 15% ($p < 0.05$) compared with the control (Figure 4M). Both A β_{1-40} and A β_{1-42} had no significant effect on the ATP levels in H9c2 cells (Figure 4N). Analysis of mitochondrial CRC demonstrated high sensitivity of the cells to A β_{1-42} . Incubation of primary HCAEC with 10 μ M A β_{1-42} for 48 h decreased the mitochondrial CRC by 15% ($p < 0.05$) compared to control cells (Figure 5A,B). Likewise, 10 μ M A β_{1-42} induced a 20% ($p < 0.05$) decrease of the mitochondrial CRC in H9c2 cardioblasts (Figure 5C,D). A β_{1-40} did not affect the mitochondrial CRC in both types of cells.

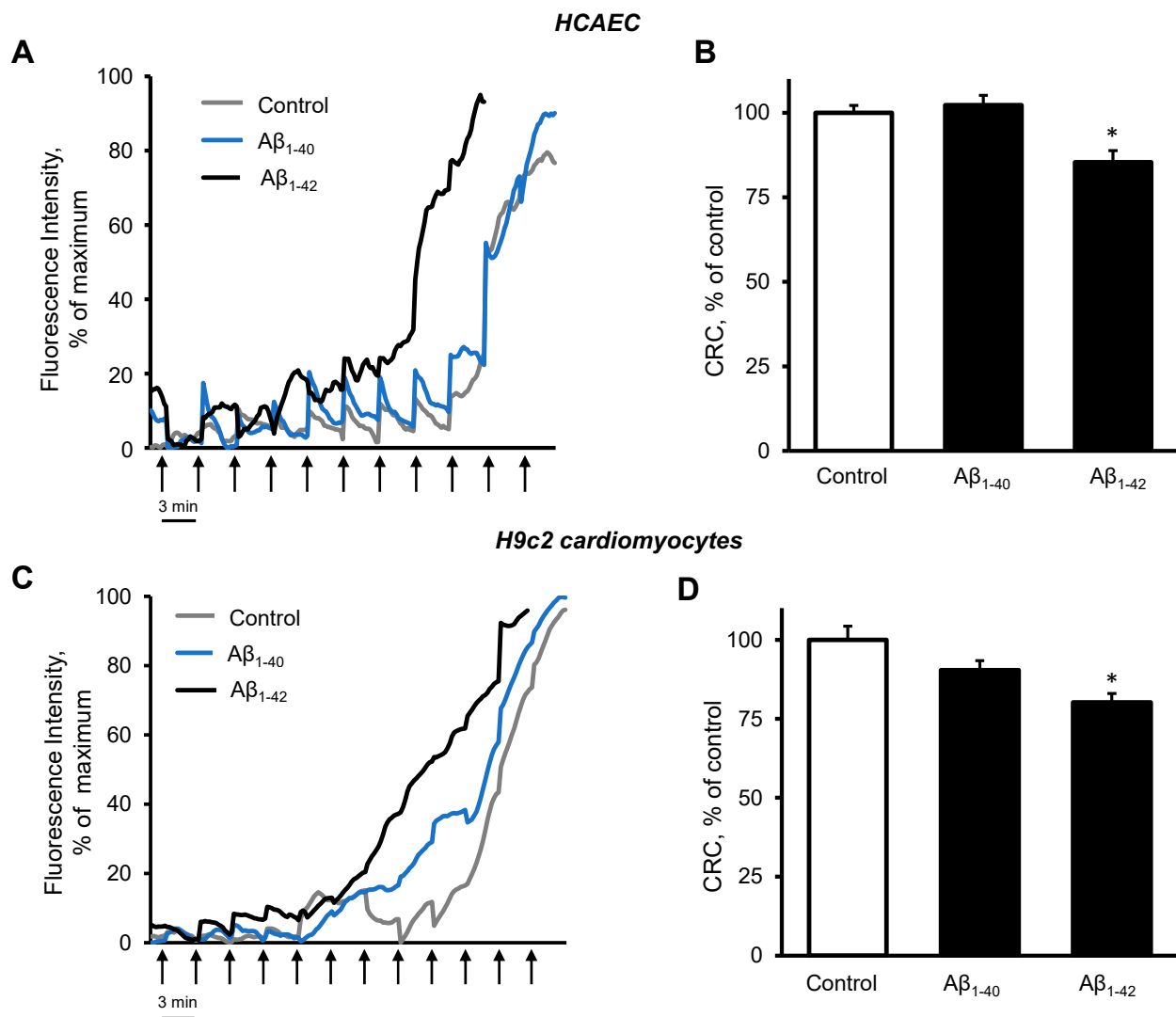


Figure 5. A β_{1-42} decreased CRC. (A,B): Primary HCAEC grown for 48 h in the presence and absence of 10 μ M A β_{1-40} or A β_{1-42} . (A): Representative traces of the fluorescence intensity of Calcium Green 5N. (B): Quantification of CRC calculated from the burst cycles (cycles that showed the highest fluorescence signal increase). (C,D): H9c2 cardiomyocytes were grown for 48 h in the presence and absence of 10 μ M A β_{1-40} or A β_{1-42} . (C): Representative traces of the fluorescence intensity of Calcium Green 5N. (D): CRC was calculated from the burst cycles. The addition of 1 nmol calcium was done every 3 min (one cycle) at 37 °C. Assays without cells were used as a baseline. $n = 6$ per group. * $p < 0.05$ vs. control.

Immunocytochemical analysis of the primary HCAEC incubated with 10 μM $\text{A}\beta_{1-42}$ for 48 h showed aggregated $\text{A}\beta$ throughout the cytoplasm (Figure 6A, green). Mitochondria in $\text{A}\beta_{1-42}$ -treated cells showed mostly fragmented mitochondria, whereas control cells and the cells incubated with $\text{A}\beta_{1-40}$ presented a well-organized mitochondrial network (Figure 6A, white). Likewise, mitochondria were more fragmented in H9c2 cardioblasts incubated with 10 μM $\text{A}\beta_{1-42}$ for 48 h (Figure 6B, white) and showed aggregated $\text{A}\beta$ in the cytoplasm (Figure 6B, green).

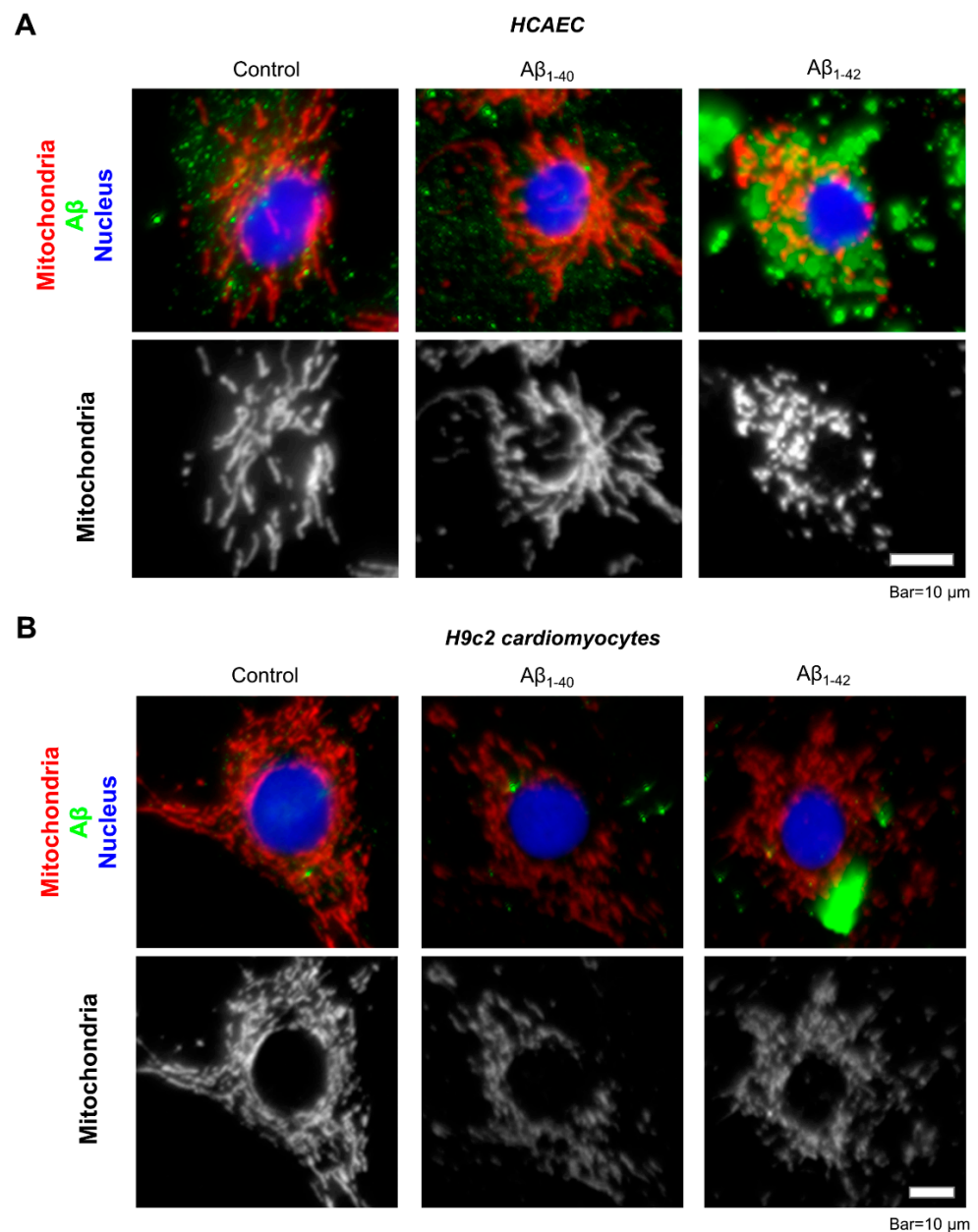


Figure 6. Visualization of $\text{A}\beta$ aggregates and mitochondria by immunostaining. (A): Representative images of primary HCAEC. (B): Representative images of H9c2 cardiomyocytes. Primary HCAEC and H9c2 cardiomyocytes grown for 48 h in the presence and absence of 10 μM $\text{A}\beta_{1-40}$ or $\text{A}\beta_{1-42}$ were fixed and stained with anti-amyloid beta (green) and anti-ATP5A antibody (red). Nuclei were stained by DAPI. Bar = 10 μm .

3.5. $\text{A}\beta$ induced Dysfunction of Cardiac Mitochondria In Vitro

To investigate the possible direct effects of $\text{A}\beta$ on the mitochondria, we measured mitochondrial swelling rates, mtROS, $\Delta\Psi\text{m}$, and ATP levels in mitochondria that were

isolated from healthy rat hearts. The mitochondria were incubated with 10 μ M A β ₁₋₄₀ or A β ₁₋₄₂ for 10 min. Results demonstrated that A β ₁₋₄₀ and A β ₁₋₄₂ further increased the Ca²⁺-induced mitochondrial swelling rate by 34% ($p < 0.05$) and 97% ($p < 0.05$), respectively, in comparison with the control (Figure 7A,B). The swelling of mitochondria apparently was induced by mPTP opening since the addition of sanglifehrin A, a cyclophilin D inhibitor, completely prevented the mitochondrial swelling in control and A β -treated groups (Figure 7B). A β ₁₋₄₂ slightly (5%, $p < 0.05$) increased mtROS (Figure 7C), decreased the $\Delta\Psi_m$ by 12% ($p < 0.05$, Figure 7D), and increased ATP levels (8%, $p < 0.05$), (Figure 7E). A β ₁₋₄₀ did not induce any remarkable changes in mtROS, $\Delta\Psi_m$, and ATP levels.

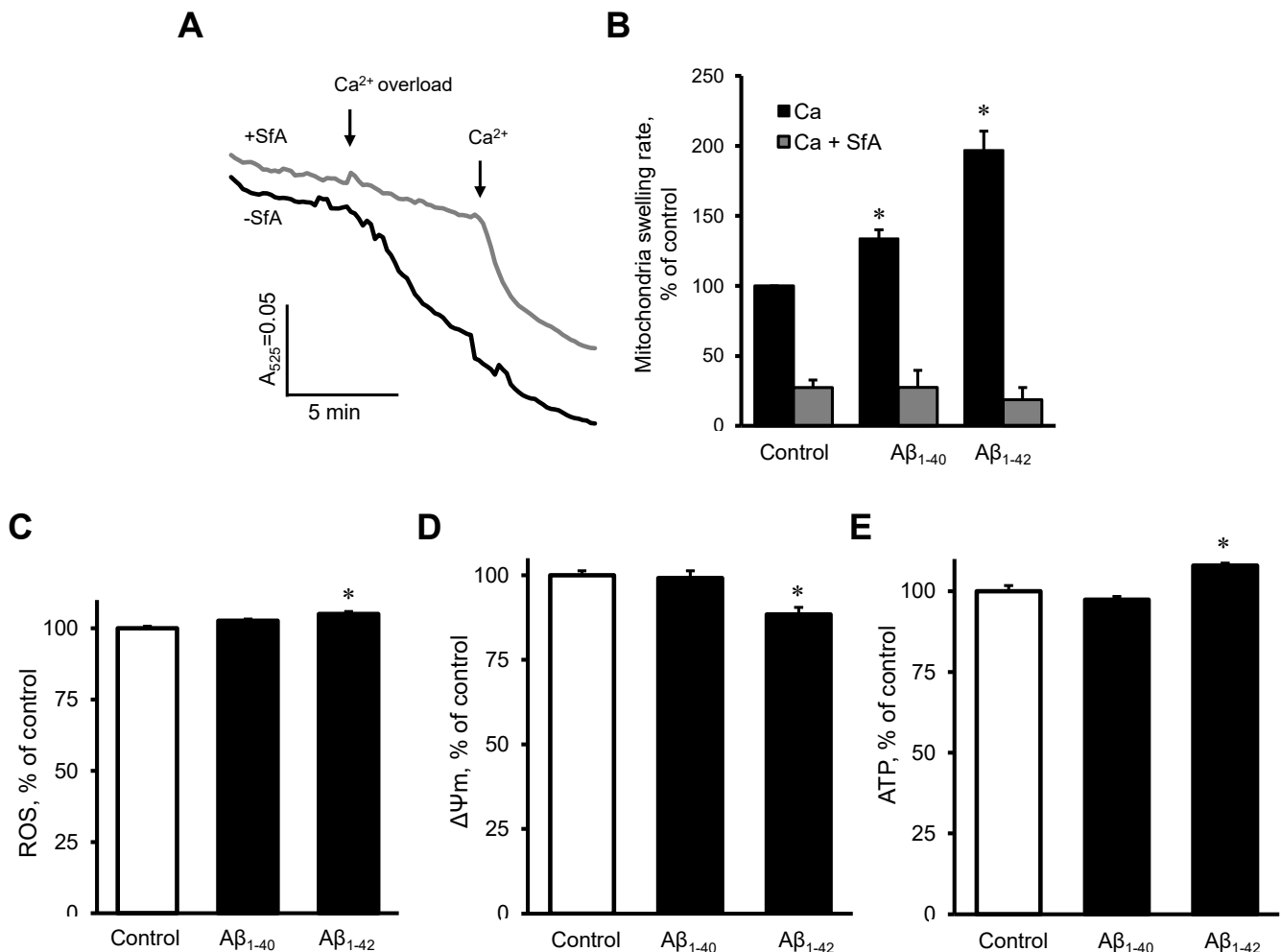


Figure 7. The effect of A β on isolated cardiac mitochondria. (A,B): Isolated rat heart mitochondria were used to analyze mitochondria swelling. (A): Mitochondrial swelling measured as decrement of A₅₂₅ during the first 2 min of the calcium overload. Sanglifehrin A (SfA), a cyclophilin D (an mPTP regulator) inhibitor, was used to demonstrate that the swelling was mediated by mPTP. (B): Quantification of mitochondrial swelling rate presented as a percentile of control. A β ₁₋₄₀ or A β ₁₋₄₂ were added 10 min before the experiment to 10 μ M. 0.1% DMSO was used as a vehicle. $n = 3$ –6 per group. (C): mtROS levels (D): $\Delta\Psi_m$. (E): ATP levels. $n = 4$ per group. * $p < 0.05$ vs. control.

4. Discussion

Results of the present study demonstrate the direct detrimental effects of A β , particularly A β ₁₋₄₂, on cardiomyocytes and primary coronary endothelial cells. On the other hand, the cardiac cells demonstrated a different sensitivity to A β . Comparative analysis of early (48 h) responses for both cells revealed severe effects of A β ₁₋₄₂ on the mitochondria. We

have previously shown that the mitochondrial bioenergetics, metabolism, function, and morphology of H9c2 are similar to primary cardiomyocytes [29].

Several metabolic alterations could be involved in the adverse action of A β ₁₋₄₂ on mitochondria. A β oligomers have been shown to disrupt cell membrane permeability and calcium homeostasis in neurons [33], cerebral endothelial cells [21,24,34], and cardiomyocytes [5]. Our results showed that A β decreased the cell viability of cardiac cells as well as primary HCAEC. Our study and others have reported that mitochondrial dysfunction and mitochondria-mediated apoptosis play a crucial role in the pathogenesis of both AD and heart failure [19,35]. However, despite these findings, it is not clear whether these changes are the cause or just another result of aging in AD and heart failure patients. Our study showed early signs of cellular and mitochondrial dysfunctions because of A β . In addition to cerebral vasculature, A β peptides were found in atherosclerotic lesions and platelets [36]. High blood levels of A β ₁₋₄₀ in patients with coronary heart disease were identified as a marker that could predict a high risk of mortality [37]. Analysis of the hearts and brains of patients with AD demonstrated that the hearts of several patients contain A β deposits (A β ₁₋₄₀ and A β ₁₋₄₂) that are structurally similar to those found in the brain. Similar amounts of A β ₁₋₄₀ and A β ₁₋₄₂ peptides were also found in the hearts and brains of AD patients. [6]. The patients with A β deposition in the heart presented diastolic dysfunction, although none of them had a history of coronary heart diseases. A β induced decreased complex activity, H₂O₂ production, ATP synthesis, state 3 and 4 respiration, and release of cytochrome c in rat muscle mitochondria [38]. Interestingly, similar mitochondrial dysfunction and apoptotic mechanisms were shown by our group in cerebral microvascular endothelial cells challenged with multiple A β variants, including A β ₁₋₄₀ and A β ₁₋₄₂ [19,21,24,26].

Our results demonstrated that A β increases mitochondrial or cellular ROS production, which, in turn, can induce further propagate A β and tau protein production [4]. Analysis of the hearts of AD patients resulted in the discovery of the presence of amyloid aggregates (A β ₁₋₄₀ and A β ₁₋₄₂) in cardiomyocytes and interstitial spaces, and that was associated with myocardial diastolic dysfunction [6]. Although it is not yet clear what is the source of the A β aggregates in the hearts of AD patients, accumulation of A β has been observed not only in the heart but also in other organs of AD patients [39], suggesting that circulating A β can be deposited or that these peptides can be produced in the heart, among other organs. Furthermore, circulating A β oligomers, recognized as the major toxic species for multiple cell types, could participate in the induction of cardiac or vascular endothelial dysfunction through oxidative stress mechanisms. On the other hand, amylin and amyloid deposits were found in the hearts of non-AD patients with diabetic and idiopathic cardiomyopathy [5,40].

Our data demonstrate that A β has a direct effect on isolated mitochondria, although A β is not produced in mitochondria [41]. Existing data on the transportation/localization of A β in mitochondria are still controversial. Accumulation of amyloid precursor protein across mitochondrial import channels (TOM and TIM) was detected in brain mitochondria of AD patients [42]. Alternatively, extracellular oligomeric A β aggregates may affect mitochondrial function by triggering cell membrane receptors (e.g., death receptors) and/or other signal transduction pathways that converge on the mitochondria [19]. A β has been shown by many studies, including ours, to diminish mitochondrial respiration and increase the levels of mtROS in multiple cell types, including neuronal and cerebral endothelial cells [19,21,24,25,34,43], and thus, can induce further A β and tau protein production [4]. However, there are few if any studies that investigate the direct effects of A β on cardiomyocytes and coronary endothelial cells.

A β neurotoxicity is associated with intraneuronal Ca²⁺ dyshomeostasis; increased cytosolic Ca²⁺ levels were detected in AD mice [44] and after application of soluble A β oligomers to the brain of wild-type mice [45]. Our results demonstrated that A β accelerated the mitochondria swelling caused by Ca²⁺ overload, an indicator of the mPTP opening. Studies on the brain tissue and neuronal cells revealed that the mPTP opening induced by

high Ca^{2+} is one of the mechanisms that mediate the effects of $\text{A}\beta$ and tau protein leading to mitochondrial dysfunction and cell death [46]. Our results showed that the cyclophilin D inhibitor sanglifehrin A completely inhibited mitochondrial swelling induced by Ca^{2+} and by $\text{Ca}^{2+} + \text{A}\beta$, suggesting the effect of $\text{A}\beta$ on the swelling depends on the mPTP opening. Cyclophilin D, a major mPTP regulator in the matrix, was found increased in AD-affected brain regions; $\text{A}\beta$ -cyclophilin D complex was detected in $\text{A}\beta$ -rich mitochondria from AD brain and transgenic AD mice, suggesting that the effects of $\text{A}\beta$ to induce mPTP opening are mediated through its interaction with cyclophilin D [47]. Conversely, genetic or pharmacological inhibition of cyclophilin D prevented $\text{A}\beta$ -induced mPTP opening and cell death [48], decreased mitochondrial and neuronal perturbations, and improved learning and memory in AD [49].

Mitochondrial quality control mechanisms, including mitophagy and mitochondrial biogenesis and dynamics, are compromised by aging. A large number of factors and mechanisms maintain the structural and functional integrity of mitochondria, and modulation of their intensity/efficiency with aging apparently impairs structural and functional integrity of mitochondria and diminishes the mitochondrial quality control leading to mitochondrial dysfunction and, eventually, age-related diseases such as AD [50]. In this context, our results demonstrated that, among other factors, $\text{A}\beta$ accumulation in cardiac cells with aging might play a certain role in mitochondrial/cellular dysfunction in the elderly population.

5. Conclusions

This study reveals adverse effects of $\text{A}\beta$, particularly $\text{A}\beta_{1-42}$, on cardiomyocytes and coronary endothelial cells that could be mediated, among other mechanisms, through functional and metabolic alterations of mitochondria. Like neurons and cerebral endothelial cells, mitochondrial abnormalities might stimulate the intrinsic apoptotic pathway leading to cell death. Although the source of $\text{A}\beta$ and mechanisms of accumulation of $\text{A}\beta$ aggregates in the heart in the aged population and AD patients remains undiscovered, this study opens new perspectives for elucidating the potential role of $\text{A}\beta$ in cardiac dysfunction.

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