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Amyloid- β Causes Mitochondrial Dysfunction via a Ca^{2+} -Driven Upregulation of Oxidative Phosphorylation and Superoxide Production in Cerebrovascular Endothelial Cells

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Abstract

Cerebrovascular pathology is pervasive in Alzheimer's disease (AD), yet it is unknown whether cerebrovascular dysfunction contributes to the progression or etiology of AD. In human subjects and in animal models of AD, cerebral hypoperfusion and hypometabolism are reported to manifest during the early stages of the disease and persist for its duration. Amyloid- β is known to cause cellular injury in both neurons and endothelial cells by inducing the production of reactive oxygen species and disrupting intracellular Ca^{2+} homeostasis. We present a mechanism for mitochondrial degeneration caused by the production of mitochondrial superoxide, which is driven by increased mitochondrial Ca^{2+} uptake. We found that persistent superoxide production injures mitochondria and disrupts electron transport in cerebrovascular endothelial cells. These observations provide a mechanism for the mitochondrial deficits that contribute to cerebrovascular dysfunction in patients with AD.

Keywords

Amyloid- β ; calcium; mitochondria; superoxide; vascular endothelial cells

INTRODUCTION

Alzheimer's disease (AD) is the most common form of senile dementia and is characterized by progressive neurodegeneration and cognitive decline. Parenchymal plaques are a hallmark of AD and are formed by the deposition of amyloid- β ($\text{A}\beta$) polymers [1, 2]. The contribution of $\text{A}\beta$ deposition to the manifestation and progression of AD has been the focus of investigation since the discovery of the peptide. Remarkably, 90% of patients with AD demonstrate cerebral amyloid angiopathy, a neuropathological disease characterized by the deposition of $\text{A}\beta$ on the walls of cerebral vasculature [3–6].

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SUPPLEMENTARY MATERIAL

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Cerebrovascular comorbidities are common in AD, as many as 92% of AD patients demonstrate ischemic white matter lesions that resemble arteriosclerosis of small vessels. Microvascular degeneration can have obvious detrimental consequences to cerebral tissue. A number of reports show regional hypoperfusion, hypometabolism, and blood-brain barrier hyperpermeability in subjects with AD [7–9]. To date, a large prevalence of patients with AD are at risk for developing severe vascular conditions that include: hemorrhagic stroke, spontaneous cerebral emboli, cerebral microinfarctions, and microhemorrhages [8, 10–12]. A growing body of evidence has emerged documenting cerebrovascular dysfunction preceding cognitive decline in AD patients [13–15], suggesting that vascular dysfunction may play a causative role in the emergence of AD. However, the mechanism(s) by which A β exerts its cytotoxic effects on the cerebrovasculature are not yet known.

It is known that bioenergetic deficits occur in neurons exposed to A β and that mitochondrial dysfunction precedes the onset of pathology in AD [16, 17]. These observations suggest that mitochondrial dysfunction during the early stages of AD (non-symptomatic) compromise cerebrovascular function resulting in cerebral A β retention, vascular degeneration, and hypoperfusion, which together may initiate the cognitive decline and symptomatic phase of AD. Observations of postmortem brain tissue from AD patients have revealed extensive mitochondrial network disruption, presenting as evidence of excessive mitochondrial fission and fragmented cristae [18–21]. In cultured cortical neurons, A β exposure results in deficits of synaptic mitochondria and reduced mitochondrial transport, likely contributing to the synaptic dysfunction in AD [22]. Together, these observational studies indicate a pivotal role for mitochondrial injury in the cytotoxicity of A β . Cellular production of reactive oxygen species (ROS) has been shown to result in S-nitrosylation of Drp-1 resulting in aberrant mitochondrial fission, synaptic loss, and neuronal damage [23]. Increased production of A β -induced ROS has also been demonstrated in cerebrovascular endothelial cells and causes cell dysfunction, blood-brain barrier disruption, and degeneration [6, 24]. The mechanism by which A β induces the production of ROS in endothelial cells is thought to involve the activation of the NADPH oxidase membrane-bound subunit, Nox [25, 26]. Genetic ablation of vascular endothelial cell Nox2 or inhibiting NADPH assembly abrogates ROS and vascular dysfunction induced by A β [24, 27, 28].

In the present study, we describe a mechanism for A β -induced cerebrovascular endothelial cell injury. We provide experimental evidence from primary cultures of vascular endothelial cells and a bEnd.3 cell line exposed to A β , which showed increased production of mitochondrial superoxide via direct A β ₁₋₄₂ interaction with mitochondria. A β induced a dose-dependent increase in mitochondrial superoxide production that paralleled an increase in mitochondrial oxygen consumption and ATP production, suggesting mitochondrial stress. We found that A β exerts these effects by increasing mitochondrial Ca²⁺ concentration, driving hyper-oxidative metabolism, and increasing ROS production by mitochondria. Furthermore, removal of extracellular Ca²⁺ abrogates the A β ₁₋₄₂-induced increase of mitochondrial oxygen consumption, ATP production, Ca²⁺ accumulation, superoxide production, and cerebrovascular endothelial cell death. These A β -induced effects on mitochondrial function may provide a mechanism for the mitochondrial dysfunction and deficits observed in AD and provide evidence for a therapeutic intervention that targets extracellular calcium.

MATERIALS AND METHODS

Study design

All experiments were performed using either primary cerebrovascular endothelial cells from mice or the bEnd.3 [BEND3] (RRID: CVCL_0170, ATCC® CRL-2299) mouse brain capillary endothelial cell line. The brain endothelial cell line is not listed as a commonly misidentified cell line by the International Cell Line Authentication Committee. Initial authentication of the bEnd.3 cell line was performed by the vendor; no further authentication was performed in the laboratory. Cells were cultured with Dulbecco's modified Eagles medium (DMEM) (Cat. No. SH30022.01 (2019), HyClone GE) supplemented with 10% fetal bovine serum (Cat. No. S12450 (2019), Atlanta Biosciences) and 1% penicillin streptomycin (Cat. No. SV30010 (2019), HyClone GE) (DMEM+). Cells were plated in either 96-well assay plates (1.5×10^4 cells/well), 24-well culture plates (3.5×10^4 cells/well), or 10-cm culture dishes (1.5×10^5 cells/dish) and incubated at 37°C in a humidified incubator under 5% CO₂. Primary cerebrovascular endothelial cells were subcultivated every 4 days (ratio = 1 : 2) and bEnd.3 cells every 3 days (ratio = 1 : 8). The first and second passage of primary cell cultures were retained for cryopreservation. For experimental purposes, primary endothelial cell subcultures were utilized beginning at passage 3 up until a maximum passage number of 6. Experiments that used the bEnd.3 cell line were performed on cells at passage 15–20. This study was not pre-registered or blinded. Stratified randomization was employed in all experiments that utilized cultured cells to prevent any experimental condition to be statically positioned across all experiments.

Animal usage

Three-month-old C57BL6 (National Institutes on Aging) pregnant female mice ($n = 4$) were used to produce primary cerebrovascular endothelial cell cultures for the purposes of this study. Primary cell cultures were prepared from embryos at gestational day 19 from each of the pregnant mice. Primary cells cultures were prepared from each mouse over four separate occasion, around 12 : 00pm eastern time. Pregnant mice produced an average of 6 embryos that were harvested and pooled together to produce a culture of primary cerebrovascular endothelial cells. Therefore, each mouse produced a single culture of primary cells and each of the four instances of prepared primary cultures was used to replicate the experiments described in the study. Power analysis for ANOVA designs indicated a sample size of 3 pregnant mice (power = 0.999) for an effect size of $f = 1.25$. Mice were housed in accordance to IACUC guidelines of West Virginia University (protocol #: 13-0704). Animals were maintained under a light/dark cycle (12 : 12 h) with food and water available *ad libitum*. For all procedures performed in this study that involved the use of laboratory animals were carried out in accordance with the National Institutes of Health guide for the care and use of laboratory animals (NIH Publication No. 8023, revised 1978) and in compliance with the ARRIVE guidelines.

Preparation of primary cerebrovascular endothelial cell cultures

Primary cerebrovascular endothelial cells were prepared from embryonic mice. At embryonic day 19, pregnant mice were deeply anesthetized with 4% isoflurane diffused into 70% nitrogen and 30% oxygen mixture. After confirming deep anesthetization via tail pinch,

mice were euthanized by cervical dislocation. Mice received a vertical incision to the abdomen. Through the incision, each fetus was extracted, removed from the amniotic sack, and decapitated. The collected fetal heads were immediately immersed in ice-cold HBSS buffer (Cat. No. 14025-092 (2019), Gibco). Each brain was extracted through an incision made along the superior sagittal suture followed by the removal of the cerebellum for discard. The remaining tissue was placed into a 50 mL centrifuge tube containing ice-cold DMEM. The brain tissue from each fetus was consolidated into a single 50 mL centrifuge tube containing 20 mL of ice-cold DMEM then centrifuged at 2500 x g for 3 min. The supernatant was decanted and the tissue pellet was resuspended in 1 mL of ice-cold DMEM. The suspended tissue was then transferred to a Dounce homogenizer that was kept on ice. After homogenization, the tissue was transferred into a 50 mL tube containing 1.5 % HEPES 1M (Cat. No. H0887-100ML (2019), Sigma Aldrich) in 20 mL HBSS. The tissue was briefly vortexed followed by centrifugation at 2000 x g for 10 min at 4°C. Supernatant was decanted and the pellet was resuspended in a solution of 18% (w/v) dextran (Cat. No. 31392-50G (2018), Sigma Aldrich) and 1.5% HEPES 1M in 20 mL HBSS. The tube was mixed by inverting then centrifuged at 5000 x g for 10 min at 4°C. Once centrifuged, the supernatant along with the cholesterol interphase was discarded; the pellet was then resuspended in 20 mL of warm DMEM containing 10% FBS and 1% penicillin streptomycin. Vessels were isolated from the suspension by passing the solution through a 50 µm filter, and rinsed with 3 mL of DMEM. Isolated vessels were removed from the filter by immersion in warm DMEM containing 10% FBS and 1% penicillin streptomycin. Collected vessels were centrifuged in a 50 mL tube at 2000 x g for 3 min, resuspended in 1 mL of DMEM and plated in 10 cm culture dishes. Vessels were allowed to reach 65% confluence before passaging and several passages were performed before experimental use. This procedure yields cerebrovascular endothelial cell cultures primarily of capillary in origin.

Mitochondria isolation from cerebrovascular endothelial cells

Mitochondrial isolation was performed on primary cerebrovascular endothelial cells via a previously reported technique [29]. Briefly, cells were cultured to 85% confluence then harvested with a cell scraper and collected into a 50 mL centrifuge tube. Endothelial cells were pelleted by centrifugation at 2000 x g for 3 min, supernatant was discarded and pellet resuspended in 8 mL. With a Dounce homogenizer, cell membranes were fractured, releasing the mitochondria into the medium. The homogenate was transferred into a 15 mL centrifuge tube then aspirated several times using a 10 mL syringe with a 27-gauge needle. The homogenate was centrifuged at 800 x g to pellet the cell debris. After centrifugation, the supernatant was decanted into a new 15 mL tube then distributed into 2 mL microcentrifuge tubes. Microcentrifuge tubes were centrifuged at 10000 x g for 5 min at 5°C to pellet the mitochondria. Once centrifugation completed, the supernatant was discarded and the mitochondrion pellet was resuspended in a solution of 10 mM Tris-MOPS (Cat. No. M-8899 (2019), Sigma Aldrich) at pH 7.4, 1 mM EGTA-Tris (Cat. No. E8145-10G (2019), Sigma Aldrich), and 200 mM sucrose (Cat. No. S5-12 (2018), Fisher Scientific).

A β ₁₋₄₂ preparation

Four separate preparations of A β ₁₋₄₂ (Cat. No. 03112 (2019), Thermo Fisher Scientific) were performed over the duration of the study. For each preparation, 1 mg of lyophilized A β ₁₋₄₂ monomers was suspended in 167 μ L of HPLC grade water (Cat. No. AM9937 (2018), Ambion) and incubated at room temperature (RT) for 5 min. The dissolved A β ₁₋₄₂ was then diluted to 230 μ M by adding 833 μ L of Ca²⁺-free phosphate buffered saline (PBS) (Cat. No. P4417-100TAB (2019), 0.01M, Sigma Aldrich) and incubated for 48 h at 37°C for polymerization. After polymerization, A β ₁₋₄₂ was separated into 50 μ L aliquots and stored at -80°C. The final product produced 230 μ M A β ₁₋₄₂ at mixed polymerization states.

Endothelial cell death assay

To assess A β ₁₋₄₂-induced cell death, bEnd.3 cells were seeded in 24-well plates at a density of 2.5×10^4 and allowed to reach 70–85% confluency. Then, vascular endothelial cells were rinsed with pre-warmed DMEM containing 10% FBS and 1% penicillin streptomycin. A β ₁₋₄₂ was prepared in 5-, 9-, and 18- μ M solutions diluted in DMEM+then added to the wells containing the vascular endothelial cells. Cerebrovascular endothelial cells were exposed to A β ₁₋₄₂ for a duration of 24 h. After exposure, cells were rinsed 3 times with 0.1 M PBS, and then fixed with paraformaldehyde (PFA) 4% (Cat. No. 15714 (2018), Electron Microscopy Sciences) for 15 min at room temperature (RT). After fixation, endothelial cells were washed 2 times for 5 min with 0.1 M PBS containing 0.045% Tween-20 (Cat. No. P1379-500ML (2018), Sigma Aldrich). Residual detergent was removed with an additional 2 washes, each for 5 min. Nuclei were stained with 50 nM 4', 6-diamidino-2-phenylindole (DAPI) (Cat. No. D-21490 (2017), Molecular Probes) counter stain. After nuclear staining, cells were washed 3 times for 5 min with 0.1 M PBS and imaged on an EVOS2 FL Auto microscope (Invitrogen). Cells were imaged at 10x magnification and fluorescence excitation using a mercury arc lamp and a DAPI filter set. Three images were acquired for each well and saved as tiff-formatted images. The number of nuclei in each of the three images were averaged and the average number of nuclei was multiplied by the area of the well floor divided by the microscope frame size, yielding the total number of cells per well. Cell death by chronic exposure to A β ₁₋₄₂ was assessed using the live-dead probe (Cat. No. R37609 (2019), Invitrogen). Cells were plated on 96-well assay plates and allowed 24 h of growth before receiving 9 μ M A β ₁₋₄₂ in DMEM+for 48, 72, 96, and 122 h. After exposure, dead and live cells were labeled and quantified.

Mitochondrial superoxide assay

Mitochondrial superoxide production in response to A β ₁₋₄₂ exposure was quantified using MitoSOX Red (Cat. No. M36008 (2019), Molecular Probes) according to the manufacturer's guidelines. Cerebrovascular endothelial cells were seeded on 96-well plates at a density of 1.5×10^4 cells per well. Endothelial cells were allowed to proliferate for 24 h (85% confluency). Cells were rinsed twice with pre-warmed DMEM and then exposed to 0-, 5-, 9-, and 18- μ M A β ₁₋₄₂ for 24 h. After exposure to A β ₁₋₄₂, cells were rinsed 3 times with warm DMEM+and endothelial cells were loaded with 5 μ M MitoSOX Red diluted in DMEM+for 10 min at 37°C. After incubation, cells were rinsed 3 times with warm DMEM +. MitoSOX fluorescence was measured with a spectrophotometer by endpoint kinetic set to

an excitation of 510 nm and an emission of 580 nm. Each well was measured in relative fluorescence units (RFU). The RFU measurement from each well per group ($n=12$) was normalized to the average RFU of the control (0 μM $\text{A}\beta_{1-42}$) group to produce a fold-change value.

Endothelial cell ATP content assay

Endothelial cell ATP content was quantified using the CellTiter-Glo luminescent assay kit (Cat. No. G7571 (2018), Promega) according to the manufacturer's guidelines. The bEnd.3 cells were seeded on 96-well plates at a density of 1.5×10^4 cells per well then allowed to proliferate for 24 h (85% confluency). Endothelial cells were rinsed twice with warm DMEM+ then exposed to 5-, 9-, and 18- μM $\text{A}\beta_{1-42}$ for 24 h. Cells were allowed to equilibrate to RT for 30 min before the ATP assay was implemented. The luciferase-based solution was measured with a spectrophotometer (BioTek) by endpoint kinetic set to measure luminescence. ATP was estimated in nM per well using a standard curve. To prevent ATP degradation, the standard curve was created immediately prior to running the assays. The standard curve was prepared by serial tenfold dilutions of ATP in DMEM+.

Mitochondrial respiration assay

Mitochondrial respiration was measured using a Seahorse XFe (Seahorse Biosciences) bioanalyzer using the MitoStress Kit (Cat. No. 101706-100 (2018), Seahorse Bioscience) according to the manufacturer's guidelines. Cultured bEnd.3 cells were seeded on 96-well XF Cell Culture Microplates at a density of 1.5×10^4 cells per well and allowed to proliferate for 24 h. Endothelial cells were exposed to 5-, 9-, and 18- μM $\text{A}\beta_{1-42}$ for 24 h. Three hours prior to running the procedure, the sensor cartridges were hydrated and the instrument parameters set to run the assay. Endothelial cells were rinsed with XF assay medium and placed in the bioanalyzer for calibration. After calibration, each well was measured via the MitoStress Test protocol that used a base measurement of oxygen consumption (pmol/min) to calculate basal respiration, maximal respiration, ATP production, spare capacity, proton leak, and non-mitochondrial respiration.

Cellular calcium assay

Cytosolic and mitochondrial Ca^{2+} levels were measured with Fluo-4AM (Cat. No. F14201 (2018), Molecular Probes) or Rhod-2AM (Cat. No. R1245MP (2018), Molecular Probes) according to the manufacturer's guidelines, respectively. Endothelial cells were seeded on either 96-well assay plates at a density of 1.5×10^4 or glass-bottom culture dishes at a density of 3.5×10^4 cells. Cells were exposed to 5-, 9-, and 18- μM $\text{A}\beta_{1-42}$ for 24 h. Before conducting the experiment, cells were rinsed twice with DMEM+. To enhance dispersion of the Ca^{2+} indicators, an equal volume of dimethyl sulfoxide (DMSO) (Cat. No. D2650-100ML (2019), Sigma Aldrich) containing 20% pluronic F-127 (Cat. No. P2443-1KG (2018), Sigma Aldrich) was added to the Fluo-4AM or Rhod-2AM stock solution. The Fluo-4AM or Rhod-2AM solution was diluted to 5 μM in DMEM and added to each well. Cells were loaded for 45 min at 37°C. After loading, endothelial cells were washed 3 times with DMEM+ and allowed to incubate for an additional 30 min for complete de-esterification. Rhod-2AM was incubated for an additional 24 h to allow for mitochondrial (non-cytosolic) Ca^{2+} indication. Fluorescence was measured with a spectrophotometer (BioTek) by area

scan routine using an excitation of 494 or 552 and an emission of 506 or 581 to measure Fluo-4AM or Rhod-2AM, respectively. Each well was measured in RFU and normalized to the average of the control group to produce a fold change value. For cells that were plated on glass-bottom culture dishes, dynamic Ca^{2+} indication was measured via confocal microscopy. Cells were imaged over a period of 94 min at 30-s intervals.

Calcium pathway inhibitors

The IP_3 R inhibitor, 2-aminoethoxydiphenylborate (2-APB) (Cat. No. 100065-100MG (2020), Sigma Aldrich) was prepared as a 100 mM stock solution in DMSO; subsequent dilution to 50 μM working concentration was prepared in DMEM+. The MCU antagonist, 4-Quinolincarboxamide (ER-000444793) (Cat. No. 50-193-1971 (2020), Fisher Scientific) was prepared as a 5 mM stock solution in DMSO; working concentration was prepared in DMEM+ at 5 μM . The mNCX inhibitor, 2-[4-[(4-nitrophenyl)methoxy]phenyl]ethyl carbamimidothioate (KB-R7943) (Cat. No. 12-441-0 (2020), Fisher Scientific) stock solutions was prepared in DMSO at a concentration of 5 mM; subsequent dilution to 7 μM working concentration was prepared in DMEM+. All compounds used in the following studies were prepared at working concentration immediately before use.

Statistics

All experiments were performed on either primary cerebrovascular endothelial cells from mice or the bEnd.3 cell line. Experiments involving isolated mitochondria were derived from primary cerebrovascular endothelial cells. Quantitative experiments were repeated a minimum of three times. Qualitative experiments were repeated a minimum of two times. Data are reported as the mean value of the experimental group across all replicates plus or minus the standard deviation (mean \pm SD). The criteria for sample exclusion consisted of human error or instrument failure. No data was excluded from the statistical analysis of the study. Statistical comparison was performed on GraphPad Prism version 8.0 (RRID: SCR_002798) by one-way ANOVA (Supplementary Table 1) to assess normality of data and with either Bonferroni's multiple comparison analysis (to compare each group to all other groups), Dunnett's post hoc analysis (to compare all groups to control group), or linear trend analysis (linear trend to determine dose-dependency) (Supplementary Table 2). All p -values less than 0.05 were considered significant ($*p < 0.05$; $**p < 0.001$; $***p < 0.0001$).

RESULTS

Acute and chronic exposure to $\text{A}\beta$ is cytotoxic to cerebrovascular endothelial cells

Previous studies have shown that, in neuronal cultures, exposure to $\text{A}\beta_{1-42}$ is cytotoxic, causing cellular death at relatively high concentrations [30–32]. However, these experiments describe an unsustainable cellular environment at late stages of AD, when $\text{A}\beta_{1-42}$ is found at significantly elevated concentrations in the brain compared to the early stages of AD. We constructed an experimental paradigm that describes the early changes to cells during the disease progression, when $\text{A}\beta_{1-42}$ is found at lower levels in the brain. To determine the cytotoxicity of acute exposure of $\text{A}\beta_{1-42}$ to cerebrovascular endothelial cells, we used brightfield microscopy to visualize live bEnd.3 cells exposed to 18 μM $\text{A}\beta_{1-42}$. We observed marked changes in cell morphology that resembled apoptosis, characterized by an initial

retraction of cellular processes after 120 min that progressed and resulted in a spherical cellular morphology after only 240 min of exposure to 18 μM $\text{A}\beta_{1-42}$ (Fig. 1A). To provide a quantitative measure of the bEnd.3 cell death we observed via live cell imaging, we counted the number of nuclei of cells exposed to 5-, 9-, and 18- μM $\text{A}\beta_{1-42}$ for 24 h and compared this to the number of nuclei of bEnd.3 cells exposed to vehicle control. We found no significant difference in the number of nuclei between bEnd.3 cells that were exposed to $\text{A}\beta_{1-42}$ and cells that were not (Fig. 1B). Although, we found some evidence of cytotoxicity for acute exposure to $\text{A}\beta_{1-42}$, we next evaluated whether prolonged exposure of cerebrovascular endothelial cells to $\text{A}\beta_{1-42}$ could induce cytotoxicity. We assessed this by exposing bEnd.3 cells to 9 μM $\text{A}\beta_{1-42}$ over a period of 48–144 h. In order to account for differences in the number of cells between the latency-groups, we normalized our cell death measurement by counting the total number of nuclei of both dead and live cells to produce a dead-to-live ratio ($F_{\text{D}}/F_{\text{L}}$), and this value was converted to fold-change from the latency-matched non-exposed control group. We found that cell death caused by 9 μM $\text{A}\beta_{1-42}$ begins after 96 h of exposure. Specifically, cell death was increased by 1.56 ± 0.15 ($p < 0.0001$)-fold and 1.52 ± 0.21 ($p < 0.0001$)-fold by 96 h and 120 h of exposure, respectively (Fig. 1C).

Cerebrovascular endothelial cell exposure to $\text{A}\beta$ results in mitochondrial dysregulation characterized by accelerated mitochondrial oxidative phosphorylation

Most cells respond to stress by increasing oxidative phosphorylation. For example, cellular damage originating from starvation [33, 34], inflammation [35], membrane rupture [36], thermal limitations [37, 38], and ionic imbalance [39, 40] can result in increased oxidative phosphorylation, which may provide additional metabolic resources that are needed to repair damage incurred. We first assessed mitochondrial oxidative phosphorylation after exposure to $\text{A}\beta_{1-42}$. We used an extracellular flux bioanalyzer XF^e 96 to assess changes in mitochondrial oxygen consumption following $\text{A}\beta_{1-42}$ exposure. Oxygen consumption in cultured bEnd.3 cell was measured for 14 min to obtain a baseline metabolic oxygen consumption rate. After 14 min, bEnd.3 cells were exposed to either 9 μM $\text{A}\beta_{1-42}$ or DMEM (control) for a total of 589 min, to determine if $\text{A}\beta_{1-42}$ modulates mitochondrial oxygen consumption. Our analysis revealed a progressive increase in mitochondrial oxygen consumption that began 397 min post- $\text{A}\beta_{1-42}$ exposure, which increased 1.81 ± 1.77 -fold ($p < 0.001$) above the control group after 461 min and continued to increase up to 3.76 ± 1.41 -fold ($p < 0.001$) above the control group over the duration of the experiment (Fig. 2A). These observations were in contrast to bEnd.3 cells maintained in DMEM without $\text{A}\beta_{1-42}$, which demonstrated an essentially unchanged oxygen consumption rate for the duration of the experiment (Fig. 2A, black).

Since our results demonstrated a clear increase in mitochondrial oxygen consumption in response to $\text{A}\beta_{1-42}$ exposure, we characterized specific indices of mitochondrial oxygen consumption. To determine this, we measured cellular basal respiration, maximum respiration, spare capacity, and proton leak after bEnd.3 cells were exposed for 24 h to 5-, 9-, and 18- μM $\text{A}\beta_{1-42}$. We found that $\text{A}\beta_{1-42}$ dose-dependently increased mitochondrial basal respiration, maximal respiration, spare capacity, and proton leak after 24 h of exposure. For all evaluated parameters of oxidative phosphorylation, the maximum effect was measured for bEnd.3 cells exposed to 18 μM $\text{A}\beta_{1-42}$. Specifically, exposure to $\text{A}\beta_{1-42}$

resulted in an increased oxygen consumption to 67 ± 5.6 ($p < 0.0001$), 159 ± 13.5 ($p < 0.0001$), 88 ± 14.9 ($p < 0.0001$), and 15 ± 1.6 ($p < 0.0001$) pmol/min of oxygen used for basal respiration, maximum respiration, spare capacity, and proton leak, respectively (Fig. 2B–E).

We then repeated this experiment on primary cerebrovascular endothelial cells to determine if $A\beta_{1-42}$ exposure results in similar mitochondrial bioenergetic hyperactivity. We observed a similar dose-dependent upregulation of mitochondrial oxygen consumption. The maximum effect was measured from cells exposed to $18 \mu\text{M}$ $A\beta_{1-42}$, which caused oxygen consumption to increase to 167 ± 11.2 ($p < 0.0001$), 311 ± 45.3 ($p < 0.0001$), 144 ± 44.9 ($p < 0.001$), and 48 ± 11.5 ($p < 0.0001$) pmol/min of oxygen used for basal respiration, maximum respiration, spare capacity, and proton leak, respectively (Fig. 2F–I). Thus, both primary cerebrovascular endothelial cells and bEnd.3 cells responded to $A\beta_{1-42}$ by increasing cellular respiration in a dose-dependent manner.

$A\beta$ exposure results in an increased mitochondrial Ca^{2+} concentration

It is firmly established that mitochondrial activity is modulated by Ca^{2+} . Since mitochondrial oxygen consumption was increased in both primary cerebrovascular endothelial cells and in bEnd.3 cells that were exposed to $A\beta_{1-42}$, we hypothesized that mitochondrial Ca^{2+} dyshomeostasis participated in the $A\beta_{1-42}$ -induced mitochondrial dysregulation. We next measured mitochondrial matrix Ca^{2+} concentration in bEnd.3 cells after exposure to 5-, 9-, and $18\text{-}\mu\text{M}$ of $A\beta_{1-42}$ for 24 h and compared this to the Ca^{2+} concentration in cells that were not exposed to $A\beta_{1-42}$. We found a dose-dependent increase of mitochondrial matrix Ca^{2+} concentrations after 24 h exposure to $A\beta_{1-42}$. Compared to vehicle-treated controls (100%), our data demonstrated that $A\beta_{1-42}$ increased mitochondrial Ca^{2+} levels to $119 \pm 30.1\%$ ($p < 0.001$) at $9 \mu\text{M}$ $A\beta_{1-42}$ and $153 \pm 24.7\%$ ($p < 0.0001$) at $18 \mu\text{M}$ $A\beta_{1-42}$ (Fig. 3A).

Because mitochondrial uptake of Ca^{2+} is a critical process required to clear cytosolic Ca^{2+} , we assessed the concentration of cytosolic Ca^{2+} after exposure to $A\beta_{1-42}$. Cytosolic Ca^{2+} was measured via Fluo-4AM fluorescence intensity after exposure to 5-, 9-, and $18\text{-}\mu\text{M}$ $A\beta_{1-42}$ for 24 h. We found a dose-dependent decrease in cytosolic Ca^{2+} concentration following $A\beta_{1-42}$ exposure. In bEnd.3 cells exposed to $18 \mu\text{M}$ $A\beta_{1-42}$, cytosolic Ca^{2+} declined to nearly $64 \pm 4.9\%$ ($p < 0.0001$) of the Ca^{2+} measured in non-exposed cells (Fig. 3B).

Cellular ATP content increases dose-dependently following exposure to $A\beta_{1-42}$

Cellular ATP production is stringently coupled to the cell's ATP requirements. Following the observation, that $A\beta_{1-42}$ exposure leads to mitochondrial Ca^{2+} accumulation and because intra-mitochondrial Ca^{2+} can modulate ATP production, we questioned whether changes to ATP production result from exposure to $A\beta_{1-42}$. We evaluated mitochondrial ATP production by calculating the oxygen consumption used to produce ATP by summing the oxygen consumed by proton leak and mitochondrial respiration then subtracting this value from the oxygen consumed by the cell's basal respiration. This produced an estimate for oxygen in pmol/min that was used for ATP production. Our analysis indicated that cells

exposed to A β ₁₋₄₂ dose-dependently increased ATP production after 24 h of exposure. bEnd.3 cells exposed to 18 μ M A β ₁₋₄₂ consumed as much as 51 ± 4.6 ($p < 0.0001$) pmol/min of oxygen for ATP production compared to only 33 ± 3.0 pmol/min of oxygen by non-exposed cells (Fig. 4A).

We next repeated the experiment on primary cerebrovascular endothelial cells to determine if A β ₁₋₄₂ mediates similar alterations to oxygen consumption for ATP production. Primary cerebrovascular endothelial cells demonstrated a dose-dependent increase of oxygen consumed for ATP production. Specifically, cells exposed to 9 μ M and 18 μ M A β ₁₋₄₂ consumed 121 ± 13.8 ($p < 0.05$) and 130 ± 7.6 ($p < 0.0001$) pmol/min of oxygen, respectively (Fig. 4B). These data indicate that primary cerebrovascular endothelial cells respond to A β ₁₋₄₂ exposure similarly to bEnd.3 cells.

An increase in mitochondrial oxidative phosphorylation is often indicative of a cellular ATP deficit. This may result from cellular injury, increased cellular activity, or as a response to cell signaling. Because we observed an increase in mitochondrial oxygen consumption for ATP production, we next measured cellular ATP content in bEnd.3 cells to determine if the changes in oxygen consumption reflect ATP availability. We found that 24 h of exposure to A β ₁₋₄₂ resulted in a dose-dependent increase in intracellular ATP content. Exposure to 18 μ M A β ₁₋₄₂ resulted in an increase in the intracellular ATP concentration to 1922 ± 41.4 ($p < 0.0001$) nM compared to 1683 ± 46.2 nM in cells that were not exposed to A β ₁₋₄₂ (Fig. 4C). Interestingly, the dose-dependent increase of intracellular ATP content occurred alongside a dose-dependent decrease in extracellular ATP. We found that the medium from cell that were not exposed to A β ₁₋₄₂ contained 7.8 ± 1.3 nM of ATP which decreased as low as 5.6 ± 0.5 ($p < 0.0001$) nM of ATP in the medium from cell exposed to 18 μ M A β ₁₋₄₂ (Fig. 4D). These data suggest that dysregulated mitochondria metabolism is characterized by aberrant ATP production that surpasses ATP utilization.

Mitochondrial superoxide production is driven by A β -induced acceleration of mitochondrial oxygen consumption

Mitochondrial production of superoxide is the primary contributor to the oxidative damage during aging and degenerative diseases [41]. We next assessed whether the A β ₁₋₄₂-induced increase in mitochondrial oxygen consumption in both primary cerebrovascular endothelial cells and bEnd.3 cells results in elevated mitochondrial superoxide production. Mitochondrial superoxide was probed with a fluorescence indicator after bEnd.3 cells were exposed to A β ₁₋₄₂ for 24 h. The fluorescence intensity of the superoxide indicator was converted into fold-change relative to vehicle treated cells. Cultured bEnd.3 cells demonstrated a remarkable dose-dependent increase in mitochondrial superoxide production following 24 h exposure to A β ₁₋₄₂. Mitochondrial superoxide production increased 9.2 ± 3.9 ($p < 0.0001$)-fold and 12.1 ± 3.8 ($p < 0.0001$)-fold above vehicle treated cells after 24 h exposure to 9 μ M and 18 μ M A β ₁₋₄₂, respectively (Fig. 5D).

We next assessed the effects of increased extracellular Ca²⁺ on mitochondrial superoxide production and compared this with superoxide production following A β ₁₋₄₂ exposure. To determine this, bEnd.3 cells were probed with a fluorescent superoxide indicator then imaged via confocal microscope after 5 min to obtain a baseline and measurement (Fig. 5A).

After baseline measurement, cells were exposed to either 5 mM of Ca^{2+} or 9 μM of $\text{A}\beta_{1-42}$ and incubated for 30 min before obtaining an outcome measurement. Our time series image analysis revealed a 1.6 ± 0.2 ($p < 0.0001$)-fold increase above baseline in mitochondrial superoxide production after 30 min exposure to 9 μM $\text{A}\beta_{1-42}$ (Fig. 5B). Similarly, cells that were exposed to an additional 5 mM of Ca^{2+} demonstrated a 1.4 ± 0.1 ($p < 0.0001$)-fold increase above baseline in superoxide production after 30 min of exposure (Fig. 5C). These data suggest that increased Ca^{2+} concentration can drive superoxide production in bEnd.3 cells at a magnitude similar to cells exposed to $\text{A}\beta_{1-42}$ for the same amount of time.

We were unsure if direct interaction between $\text{A}\beta_{1-42}$ and mitochondria produce the mitochondrial changes we observed. To answer this question, we isolated the mitochondria from primary cerebrovascular endothelial cells and measured superoxide production following $\text{A}\beta_{1-42}$ exposure for 1 h. Isolated mitochondria exposed to $\text{A}\beta_{1-42}$ for 1 h resulted in a dose-dependent increase in mitochondrial superoxide production. Exposure to 1-, 3-, 6-, and 9- μM $\text{A}\beta_{1-42}$ demonstrated a 1.35 ± 0.07 ($p < 0.001$), 1.95 ± 0.10 ($p < 0.0001$), 3.10 ± 0.13 ($p < 0.0001$), and 4.29 ± 0.27 ($p < 0.0001$)-fold increase in superoxide production relative to control, respectively (Fig. 5E).

$\text{A}\beta$ mediates mitochondrial dysregulation by increasing mitochondrial matrix Ca^{2+} that accelerates mitochondrial oxidative phosphorylation and superoxide production

Mitochondrial influx of Ca^{2+} has been shown to accelerate oxidative phosphorylation and ATP production by increasing the activity of mitochondrial dehydrogenase enzymes. Therefore, we hypothesized that the increased oxygen consumption and production of superoxide by cerebrovascular endothelial cell mitochondria following exposure to $\text{A}\beta_{1-42}$ is mediated by the $\text{A}\beta_{1-42}$ -induced Ca^{2+} accumulation in mitochondria. To determine if the increased Ca^{2+} in the mitochondrial matrix following the exposure of bEnd.3 cells to $\text{A}\beta_{1-42}$ mediates the hyperoxidative state of mitochondria, we reduced the Ca^{2+} available to mitochondria by chelation. Extracellular Ca^{2+} was chelated with 1 mM ethylenediaminetetraacetic acid (EDTA) supplemented in cell culture medium with or without exposure to $\text{A}\beta_{1-42}$. We first assessed whether chelation of extracellular Ca^{2+} results in reduced $\text{A}\beta_{1-42}$ -induced mitochondrial Ca^{2+} accumulation. Cultured bEnd.3 cells were plated in 96-well assay plates at a density of 1.5×10^4 cells per well and grown for 24 h. Cells were then exposed to either vehicle or 9 μM $\text{A}\beta_{1-42}$ with or without 1 mM EDTA for 24 h and then assessed for changes in mitochondrial Ca^{2+} levels. Our results indicated that 1 mM EDTA effectively abolished the $\text{A}\beta_{1-42}$ -induced accumulation of Ca^{2+} in the mitochondrial matrix (Fig. 6A). Specifically, cells exposed to 9 μM $\text{A}\beta_{1-42}$ without EDTA (9 μM $\text{A}\beta_{1-42}$ /0 mM EDTA) increased mitochondrial Ca^{2+} to 1.8 ± 0.4 -fold (versus control, $p < 0.0001$) above cells exposed to vehicle (control). Whereas, cells exposed to 9 μM $\text{A}\beta_{1-42}$ with 1 mM EDTA (9 μM $\text{A}\beta_{1-42}$ /1 mM EDTA) completely abolished the $\text{A}\beta_{1-42}$ -induced accumulation of mitochondrial Ca^{2+} to 0.9 ± 0.4 (mean \pm SD)-fold (versus $\text{A}\beta$ only, $p < 0.0001$) compared to control cells ($p = 0.65$) (Fig. 6A). These data indicate that extracellular Ca^{2+} chelation with EDTA is effective at reducing mitochondrial Ca^{2+} .

We next assessed the effectiveness of reducing mitochondrial Ca^{2+} accumulation with EDTA at ameliorating elevated mitochondrial oxygen consumption after exposure to $\text{A}\beta_{1-42}$. To

determine this, we measured the oxygen consumption rate of bEnd.3 cells exposed to either 0 or 9 μM $\text{A}\beta_{1-42}$ with or without 1 mM EDTA. As before, cells exposed to 9 μM $\text{A}\beta_{1-42}$ without EDTA for 24 h demonstrated increased oxygen consumption by basal respiration (114 ± 4.0 pmol/min, $p < 0.0001$), maximum respiration (296 ± 8.4 pmol/min, $p < 0.0001$), and spare capacity (182 ± 8.0 pmol/min, $p < 0.0001$). However, treatment with 1 mM EDTA completely ameliorated the $\text{A}\beta_{1-42}$ -induced increase in oxygen consumption. In cells exposed to 9 μM $\text{A}\beta_{1-42}$ with 1 mM EDTA oxygen consumption returned to near vehicle control values for basal respiration (101 ± 3.3 pmol/min, $p < 0.02$), maximum respiration (227 ± 8.8 pmol/min, $p < 0.0001$), and spare capacity (126 ± 7.2 pmol/min, $p < 0.0001$) (Fig. 6B–D). Interestingly, proton leak was the only oxidative phosphorylation parameter that increased further with EDTA treatment. Cells exposed to $\text{A}\beta_{1-42}$ alone consumed 39 ± 1.2 ($p < 0.0001$) pmol/min of oxygen by proton leak while cells exposed to both $\text{A}\beta_{1-42}$ and EDTA consumed 49 ± 1.6 ($p < 0.0001$) pmol/min of oxygen (Fig. 6E). In support of our hypothesis, reduction of mitochondrial Ca^{2+} ameliorates the increased oxygen consumption mediated by mitochondrial Ca^{2+} accumulation following exposure to $\text{A}\beta_{1-42}$. These data suggest that $\text{A}\beta_{1-42}$ exposure results in mitochondrial dysfunction by increasing mitochondrial Ca^{2+} levels.

Mitochondrial Ca^{2+} content is stimulatory to ATP synthesis by the ATP synthase respiratory complex. Therefore, we predicted that by blocking Ca^{2+} from entering mitochondria, the $\text{A}\beta_{1-42}$ -induced upregulation of ATP production should be prevented. To determine the interplay between elevated mitochondrial oxygen consumption and matrix Ca^{2+} accumulation on the production of ATP, we exposed bEnd.3 cells to 9 μM $\text{A}\beta_{1-42}$ with or without 1 mM EDTA for 24 h and then estimated the production of ATP by mitochondria. Consistently, bEnd.3 cells exposed to 9 μM $\text{A}\beta_{1-42}$ responded by increasing oxygen consumption for ATP production to 75 ± 3.5 ($p < 0.0001$) pmol/min. Oxygen consumption in cells that were exposed to 9 μM $\text{A}\beta_{1-42}$ with 1 mM EDTA restored ATP production to near vehicle control values, 51 ± 2.6 ($p < 0.0001$) (Fig. 6F).

Since mitochondrial Ca^{2+} accelerates electron transport by increasing the activity of mitochondrial dehydrogenase enzymes, we hypothesized that blocking $\text{A}\beta_{1-42}$ -induced mitochondrial Ca^{2+} accumulation would mitigate the increased production of superoxide by mitochondria following exposure to $\text{A}\beta_{1-42}$. To determine if mitochondrial Ca^{2+} accumulation mediates the $\text{A}\beta_{1-42}$ -induced increase in mitochondrial superoxide production, we exposed bEnd.3 cells to $\text{A}\beta_{1-42}$ in the presence or absence of EDTA then measured mitochondrial superoxide levels. Superoxide production by bEnd.3 cells exposed to 9 μM $\text{A}\beta_{1-42}$ /0mM EDTA increased superoxide production 5.5 ± 1.3 (versus control, $p < 0.0001$)-fold above control cells, whereas bEnd.3 cells exposed to 9 μM $\text{A}\beta_{1-42}$ /1 mM EDTA reduced superoxide production to 1.8 ± 0.2 (versus $\text{A}\beta$ only, $p < 0.0001$)-fold of control cells (Fig. 6G).

Mitochondrial fission-fusion dynamics are a transient process that serves a physiologically important function, i.e., regulation of mitochondrial quality control and metabolic homeostasis. Mitochondrial fragmentation often precedes mitophagy of damaged mitochondria that occurs during states of metabolic and oxidative stress [42]. Mitochondrial fragmentation is well-documented in AD animal models and in human AD patients [43, 44].

It is clear that oxidative stress can cause transient changes to the morphology of mitochondria while prolonged oxidation can result in mitochondrial network fragmentation leading to apoptosis [45–47]. Since our results demonstrated that chelation of extracellular Ca^{2+} was exceptionally effective at ameliorating the $\text{A}\beta_{1-42}$ -induced dysregulation of mitochondrial activity, we asked whether the chelation of extracellular Ca^{2+} can rescue mitochondria from aberrant fragmentation. We answered this by labeling mitochondria in bEnd.3 cells with MitoTracker CMXRos Red and the nuclei with DAPI counterstain following 24 h exposure to 9 μM $\text{A}\beta_{1-42}$ with or without 1 mM EDTA. Indeed, we observed fragmentation of the mitochondrial network in bEnd.3 cells exposed to $\text{A}\beta_{1-42}$ alone and found that by chelation of extracellular Ca^{2+} completely prevented the morphological changes to the mitochondrial network (Fig. 6H–K). These observations demonstrate the role of $\text{A}\beta_{1-42}$ -induced mitochondrial Ca^{2+} accumulation in the dysregulation of mitochondrial bioenergetics, resulting in mitochondrial fragmentation following oxidative damage.

Our data describes a mechanism in which mitochondrial Ca^{2+} accumulation following cell exposure to $\text{A}\beta_{1-42}$ mediates cell damage by dysregulating mitochondrial function. We next asked whether preventing the $\text{A}\beta_{1-42}$ -induced mitochondrial dysfunction could rescue cerebrovascular endothelial cells from $\text{A}\beta_{1-42}$ mediated death. To assess the cytoprotective effects of blocking $\text{A}\beta_{1-42}$ -induced mitochondrial Ca^{2+} accumulation against $\text{A}\beta_{1-42}$ mediated cell death, we exposed bEnd.3 cells to 9 μM $\text{A}\beta_{1-42}$ with or without 1 mM EDTA for 24 h and then imaged cellular changes in morphology. Exposure to either vehicle control (Fig. 6L) or 1 mM EDTA alone (Fig. 6M) caused no apoptotic cells upon visualization. While, bEnd.3 cells that were exposed to 9 μM $\text{A}\beta_{1-42}$ /0mM EDTA (Fig. 6N, P) demonstrated numerous apoptotic cells. However, bEnd.3 cells that were exposed to 9 μM $\text{A}\beta_{1-42}$ with 1 mM EDTA (Fig. 6O) completely prevented the apoptotic cells observed in cultures exposed to $\text{A}\beta_{1-42}$ alone. We observed a reduced number of apoptotic cells in cultures exposed to $\text{A}\beta_{1-42}$ with 1 mM EDTA. We observed no apoptotic cells in cultures that were not exposed to $\text{A}\beta_{1-42}$ with and without 1 mM EDTA.

$\text{A}\beta$ associated calcium entry into the mitochondria occurs via multiple intracellular calcium regulation pathways

Since we found that the $\text{A}\beta_{1-42}$ -induced elevated mitochondrial calcium could be attenuated by extracellular calcium chelation, we next assessed intracellular calcium regulation pathways involved in calcium uptake by mitochondria. To assess the contribution of intracellular calcium regulation pathways, we used pharmacological inhibitors, which prevent calcium entry into the mitochondria from the endoplasmic reticulum, mitochondrial permeability transition pore (mPTP), and mitochondrial membrane associated calcium exchangers. Mitochondrial calcium was measured with Rhod-2AM as previously stated, then exposed to 9 μM $\text{A}\beta_{1-42}$ or DMEM+control containing 50 μM 2-APB, 5 μM ER-000444793, 7 μM KB-R7943, or vehicle (DMSO) for 24 h. We found that bEnd.3 cells exposed to 9 μM $\text{A}\beta_{1-42}$ -vehicle resulted in a 3.2 ± 0.46 ($p < 0.0003$)-fold increase in mitochondrial calcium above DMEM control 1 ± 0.35 -fold. Whereas, treatment with the IP3 receptor antagonist 2-APB (50 μM) attenuated the elevated mitochondrial matrix calcium to 1.2 ± 0.27 ($p < 0.00007$)-fold. Inhibition of the mPTP with ER-000444973 (5 μM) also reduced the $\text{A}\beta_{1-42}$ -induced mitochondrial calcium accumulation to 1.8 ± 0.24 ($p < 0.002$)-

fold. Treatment with an inhibitor of mitochondrial reverse mode Na^+/Ca^+ exchanger and calcium uniporter, KB-R7943 (7 μM) similarly reduced calcium in the mitochondrial matrix upon exposure to $\text{A}\beta_{1-42}$ to 1.4 ± 0.15 ($p < 0.0002$) (Fig. 7).

DISCUSSION

Although $\text{A}\beta$ is heavily studied in AD research, its precise role in the disease pathogenesis remains unknown. In AD brains, $\text{A}\beta$ plaques are found clustered around regions with damaged and degenerating mitochondria [48, 49]. In culture, $\text{A}\beta$ causes ROS production and cell death [6, 23]. The mechanism that drives the $\text{A}\beta$ -induced production of ROS is thought to involve the Nox subunit of NADPH oxidase that catalyzes the S-nitrosylation of cellular components [25, 26]. S-nitrosylation of mitochondrial Drp-1 is shown to cause mitochondrial fission and may contribute to the mitochondrial fragmentation found in AD both in humans and in animal models [23]. In addition, $\text{A}\beta$ has been shown to cause intracellular Ca^{2+} dyshomeostasis in culture [50]. However, the interplay between $\text{A}\beta$ -induced ROS production, mitochondrial effects, and intracellular Ca^{2+} levels has not been elucidated previously.

We demonstrate in brain vascular endothelial cells that exposure to $\text{A}\beta$ causes a rapid change in cellular bioenergetics, characterized by accelerated mitochondrial oxygen consumption and increased superoxide production. We find that these changes were concurrent with the accumulation of calcium in mitochondria and could be prevented by blocking calcium from entering mitochondria by chelation.

In the present study, we investigate the early changes to endothelial cells following exposure to $\text{A}\beta$ that contribute to the cellular death observed after prolonged exposure to the peptide. We noticed a rapid change to mitochondrial oxygen consumption after 7 h of exposure to $\text{A}\beta$, and in both bEnd.3 cells and in primary cerebrovascular endothelial cells, $\text{A}\beta$ exposure increased basal respiration, maximum respiration, spare capacity, and proton leak by 24 h. Mitochondrial basal respiration is strongly influenced by the turnover of ATP and partially by the oxidation of substrates and by the leakage of protons from the intermembrane space [51, 52]. Therefore, the rate of basal respiration reflects the cellular demand for ATP. In the present study, elevated basal respiration may indicate an increased endothelial demand for ATP in response to the early cellular injury following exposure to $\text{A}\beta$. The spare respiratory capacity of a cell is defined by the ability to utilize substrate and electron transport to meet cellular energy demand before reaching its biological limit [53, 54]. We observed a dose-dependent increase in spare respiratory capacity for both bEnd.3 cells and primary cerebrovascular endothelial cells. Because we observed both an elevated basal respiration and spare respiratory capacity in cells exposed to $\text{A}\beta$, we interpret this to suggest that both an enhanced mitochondrial respiratory competency and an elevated respiratory rate participate in the mechanism of $\text{A}\beta$ -induced cellular injury.

It is firmly established that mitochondrial activity is modulated by Ca^{2+} . The influx of Ca^{2+} to the mitochondrial matrix can exert both stimulatory and inhibitory effects on mitochondrial function. Mitochondrial influx of Ca^{2+} results in upregulated oxidative phosphorylation by activating mitochondrial dehydrogenase enzymes, glycerophosphate

dehydrogenase, isocitrate dehydrogenase, pyruvate dehydrogenase, and oxoglutarate dehydrogenase [55–57]. Activation of these enzymes by Ca^{2+} increases mitochondrial oxidative phosphorylation and ATP production [58]. We found that $\text{A}\beta$ causes increased mitochondrial matrix calcium while decreasing cytosolic calcium in bEnd.3 cells. The stimulatory activity of calcium to resident dehydrogenase enzymes in the mitochondrial matrix may account for the elevated respiration we observed in both bEnd.3 cells and in primary cerebrovascular endothelial cells. In addition, increased concentrations of mitochondrial matrix Ca^{2+} have been shown to cause the formation of the permeability transition pore (PTP) on the mitochondrial membrane, allowing the release of apoptotic signaling molecules which facilitate mitochondrial-mediated apoptosis [59–61]. Thus, the cytotoxicity we observed following prolonged exposure to $\text{A}\beta$ may be driven by the PTP following $\text{A}\beta$ -induced accumulation of mitochondrial matrix calcium.

Exposure to $\text{A}\beta$ in both bEnd.3 cells and in primary endothelial cells resulted in increased oxygen consumption for ATP production and intracellular ATP content. This upregulated ATP synthesis may be an effect of the increased calcium levels in the mitochondrial matrix and its stimulatory role with mitochondrial dehydrogenase enzymes [55–57]. The elevated activity of mitochondrial dehydrogenase enzyme enhances the production of NADH. Catabolism of NADH to NAD^+ increases the proton-motive force by serving as a proton donor substrate, increasing the throughput of hydrogen ions pumped into the mitochondrial intermembrane space at complex I. The increased hydrogen ions pumped across the mitochondrial inner membrane drives the accelerated ATP synthesis by mechanically powering the coupling of ADP to inorganic phosphate by ATP synthase.

Mitochondrial superoxide production is sensitive to the proton-motive force [41]. High rates of electron transport can cause its reversal, delivering electrons to complex I; these electrons are coupled to molecular oxygen and produce superoxide radicals [62]. Indeed, exposure to $\text{A}\beta$ resulted in an increased production of mitochondrial superoxide in intact cells and in isolated mitochondria. These observations indicate that the aberrant elevated mitochondrial respiration following $\text{A}\beta$ exposure is inefficient, resulting in increased production of ROS where it can potentially damage the cell and subsequently lead to cell death.

Since mitochondrial calcium is implicated in many of the cellular changes following amyloid- β exposure, we hypothesized that blocking the $\text{A}\beta$ -induced mitochondrial calcium accumulation would mitigate the elevated mitochondrial respiration, superoxide production, and cell death following exposure to $\text{A}\beta$. Chelation of extracellular calcium with EDTA was effective at preventing the $\text{A}\beta$ -induced accumulation of calcium in the mitochondrial matrix. Thus, we evaluated the role of mitochondrial calcium accumulation as central to the mechanism underlying $\text{A}\beta$ mediated endothelial cell injury. We found that by preventing calcium accumulation in the mitochondria, the elevated respiration following $\text{A}\beta$ could be mitigated. In addition, blocking the $\text{A}\beta$ associated calcium in the mitochondrial matrix, the elevated ATP production and superoxide generation could be halted. These data suggest that calcium influx to mitochondria is a central component to the mechanism mediated by $\text{A}\beta$ during the early phase of cellular injury in cerebrovascular endothelial cells.

We further investigated the route by which calcium influx through the plasma membrane enters mitochondria. Specifically, we assessed the contribution of calcium to the mitochondria through the endoplasmic reticulum IP₃ R, reverse mode mNCX, and the PTP. Each pathway was evaluated by exposing cells to A β ₁₋₄₂ in the presence of specific antagonists then measuring mitochondrial calcium content.

Our data describe a major role for extracellular calcium in the mechanism mediating the cytotoxicity of A β ₁₋₄₂. We found that chelation of extracellular calcium prevented its accumulation in mitochondria (Fig. 6) and the increased oxidative phosphorylation and superoxide production by mitochondria. Recent studies have demonstrated that exposure of the cell membrane to A β results in increased permeability of the bilayer, caused by the insertion of A β into the membrane, forming ion conductive channels [63, 64]. In principal, these channels would allow the influx of calcium into the cell. Furthermore, we used agents that antagonize mPTP formation, mNCX, and IP₃ R to evaluate the contribution of intracellular calcium regulatory pathways in the accumulation of calcium in mitochondria. In our experiments, inhibition of any of these three calcium pathways resulted in attenuated calcium influx into mitochondria. Therefore, these data suggest that exposure to A β ₁₋₄₂ causes the cell membrane to become conductive to calcium, the resulting influx of calcium is shunted into mitochondria via multiple calcium regulatory pathways that cause the hyperoxidative state of mitochondria observed in endothelial cell mitochondria.

It is well known that mitochondria possess a sodium/calcium exchanger (mNCX) that functions in the efflux of calcium from mitochondria [65, 66]. It is possible for mNCX to function in reverse mode during conditions of metabolic and calcium dysregulation [67, 68], transporting calcium into the mitochondria. To assess this possibility, we used KB-R7943, an inhibitor of reverse mode mNCX at concentrations previously used in experiments [69], and observed a profound reduction in A β ₁₋₄₂-induced mitochondrial calcium influx (Fig. 7).

Mitochondrial calcium uptake is a critical cellular function that maintains low cytosolic calcium concentrations. Calcium influx into the mitochondrial matrix also participates in a regulatory role facilitating mitochondrial dehydrogenase enzyme activity. However, excessive accumulation of calcium by mitochondria may lead to mitochondrial damage via the induction of the mPTP. Calcium-dependent induction of the mPTP is a major mechanism of calcium-induced damage to mitochondria.

Calcium uptake into mitochondria primarily occurs via diffusion across the outer mitochondrial membrane through the voltage-dependent anion channel (VDAC). Translocation of hydrogen by the respiratory chain forms the mitochondrial membrane potential that serves as the driving force to transport calcium across the inner mitochondrial membrane, down its electrochemical gradient. Calcium influx into the mitochondrial matrix is mediated by the mitochondrial calcium uniporter (MCU) at the inner mitochondrial membrane. The MCU channel conducts calcium ions from the outer membrane space into the mitochondrial matrix. Antagonism of MCU substantially reduced A β ₁₋₄₂-induced mitochondrial calcium influx (Fig. 7).

The efflux of calcium from the endoplasmic reticulum via IP₃ R produce localized microdomains of elevated calcium concentration that are crucial for calcium uptake by the mitochondria [70]. The contact sites that hold mitochondria stationary and within the calcium-rich microdomain occurs at regions termed mitochondrial associated membranes (MAMs). MAMs are sites involved in the exchange of biomolecules, such as lipids, ROS, and calcium between the endoplasmic reticulum and mitochondria. At MAM sites, calcium is conducted into the mitochondrial intermembrane space via a large protein complex involving VDAC. To determine whether the influx of calcium into mitochondria following Aβ₁₋₄₂ exposure involves the IP₃ R pathway, we used 2-APB, an IP₃ R antagonist. Inhibition of IP₃ R eliminated Aβ₁₋₄₂-induced mitochondrial calcium influx (Fig. 7).

Together, these data describe a role for calcium in the cytotoxicity exerted by Aβ₁₋₄₂, where the cell membrane becomes conductive to extracellular calcium, therefrom the influx of calcium is shunted into the mitochondria via multiple calcium regulatory pathways, causing the hyperoxidative state of mitochondria observed following exposure to Aβ₁₋₄₂.

Consistent with the literature, we observed morphological changes to the mitochondrial network, characterized by punctated mitochondria via excessive fission [23, 43, 71–73]. Mitochondrial fission and fusion is a dynamic process modulated by the energy demand of the cell and participates in the quality control of the mitochondrial network. Damage to mitochondria, such as from ROS can cause mitochondrial fragmentation, similar to our observation in vascular endothelial cells [74–77]. Thus, we interpret these data to suggest that the fragmentation of mitochondria in bEnd.3 cells is mediated by the elevated mitochondrial superoxide production following Aβ exposure. Therefore, because we found that the Aβ associated elevation of mitochondrial superoxide production could be mitigated by preventing mitochondrial matrix calcium. As a result, we expect that the fragmentation of mitochondria could be abrogated. Indeed, we found that by reducing mitochondrial calcium via chelation halted the aberrant mitochondrial fission following Aβ exposure, lending evidence to a mechanism involving aberrant mitochondrial fission driven by the oxidative damage to mitochondria in cerebrovascular endothelial cells exposed to Aβ. Lastly, when we prevented the Aβ-induced accumulation of mitochondrial matrix calcium, the apoptotic morphology of endothelial cells was completely prevented.

Overall, our study provides evidence of early events of cellular injury that involves the elevation of mitochondrial respiration, calcium accumulation in mitochondria, and the production of superoxide. We found that calcium influx into the mitochondria is a central component to the mechanism mediating cell injury. Furthermore, our data indicates that blocking mitochondrial calcium can completely halt the pathological mechanism mediating cell death. In addition, these findings provide a mechanism for the mitochondrial dysfunction and deficits observed in AD and provide evidence for a therapeutic strategy that targets mitochondrial matrix calcium.

Supplementary Material

Refer to Web version on PubMed Central for supplementary material.

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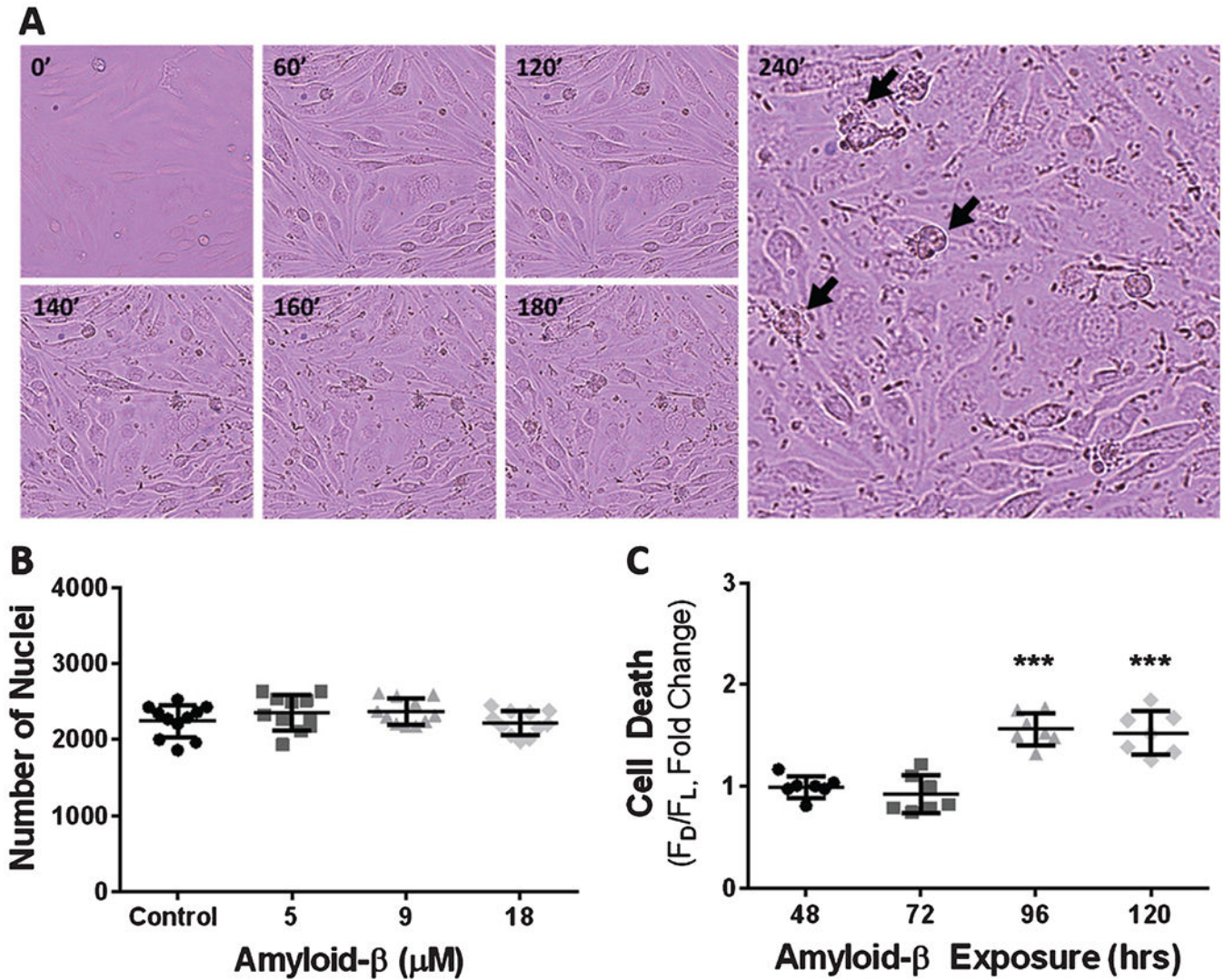


Fig. 1. Cerebrovascular endothelial cell exposure to A β results in cell death. A) Brightfield micrographs depicting bEnd.3 cell death (black arrows) following exposure to 18 μ M A β_{1-42} for 0, 60, 120, 140, 160, 180, and 240 min. B) Scatter plot (mean \pm SD) demonstrating the average number of nuclei per well after exposure to vehicle, 5 μ M A β_{1-42} , 9 μ M A β_{1-42} , and 18 μ M A β_{1-42} ($n = 11$ wells per group). C) Scatter plot (mean \pm SD) depicting endothelial cell death after chronic exposure to 9 μ M A β_{1-42} for 48, 72, 96, 120, and 144 h ($n = 7$ wells per group). One-way ANOVA with Dunnett's post analysis was used to determine the level of significance between the exposure groups (***) $p < 0.0001$.

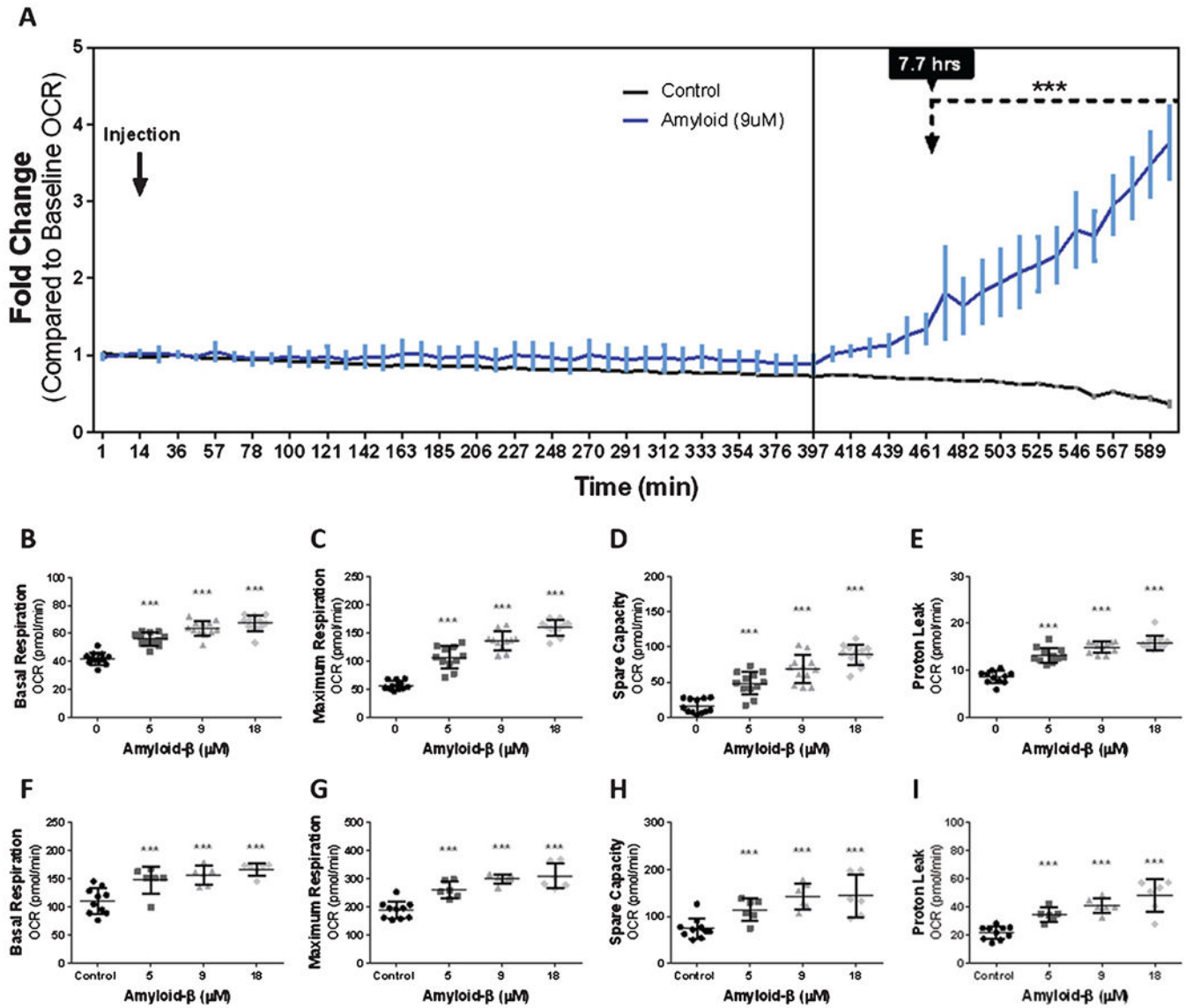


Fig. 2. Exposure to Aβ for 24-h results in a dose-dependent increase in mitochondrial oxygen consumption. **A**) Line graph (mean ± SD) depicting the fold-change of oxygen consumption to baseline as a function of time from bEnd.3 cells exposed to vehicle and 9 μM Aβ₁₋₄₂. Scatter plot(s) (mean ± SD) demonstrating oxygen consumption (pmol/min) by **(B, F)** basal respiration, **(C, G)** maximum respiration, **(D, H)** spare capacity, and **(E, I)** proton leak from **(B–E)** bEnd.3 cells and **(F–H)** primary cerebrovascular endothelial cells after 24 h exposure to vehicle, 5 μM Aβ₁₋₄₂, 9 μM Aβ₁₋₄₂, and 18 μM Aβ₁₋₄₂. One-way ANOVA with Dunnett’s post analysis and linear trend analysis was used to determine the level of significance between the treatment groups (**p* < 0.05; ***p* < 0.001; ****p* < 0.0001), **(B–E)**, *n* = 12 wells per group; **F–I**, control = 10, all other conditions = 6 wells per group).

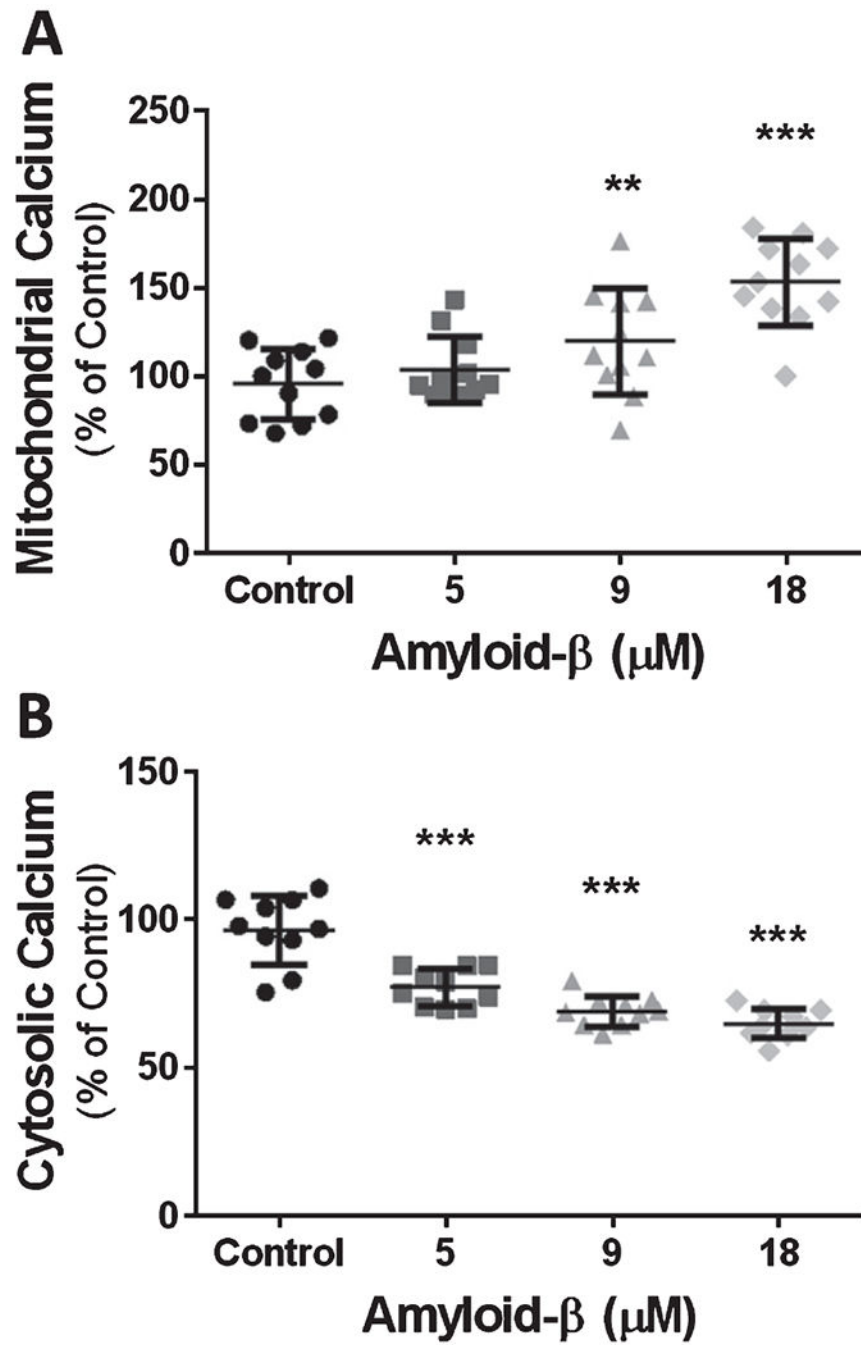


Fig. 3. Exposure to Aβ for 24 h causes a dose-dependent increase in mitochondrial Ca²⁺ content and a dose-dependent decrease of cytosolic Ca²⁺. A) Scatter plot (mean ± SD) depicting mitochondrial Ca²⁺ content as percentage of vehicle treated cells, after 24 h exposure to 5 μM Aβ₁₋₄₂, 9 μM Aβ₁₋₄₂, and 18 μM Aβ₁₋₄₂ (n = 11 wells per group). B) Scatter plot (mean ± SD) demonstrating cytosolic Ca²⁺ content as percentage of vehicle treated cells after 24 h exposure to 5 μM Aβ₁₋₄₂, 9 μM Aβ₁₋₄₂, and 18 μM Aβ₁₋₄₂ (n = 10 wells per group). One-

way ANOVA with Dunnett's post analysis and linear trend analysis was used to determine the level of significance between the treatment groups (** $p = 0.001$; *** $p = 0.0001$).

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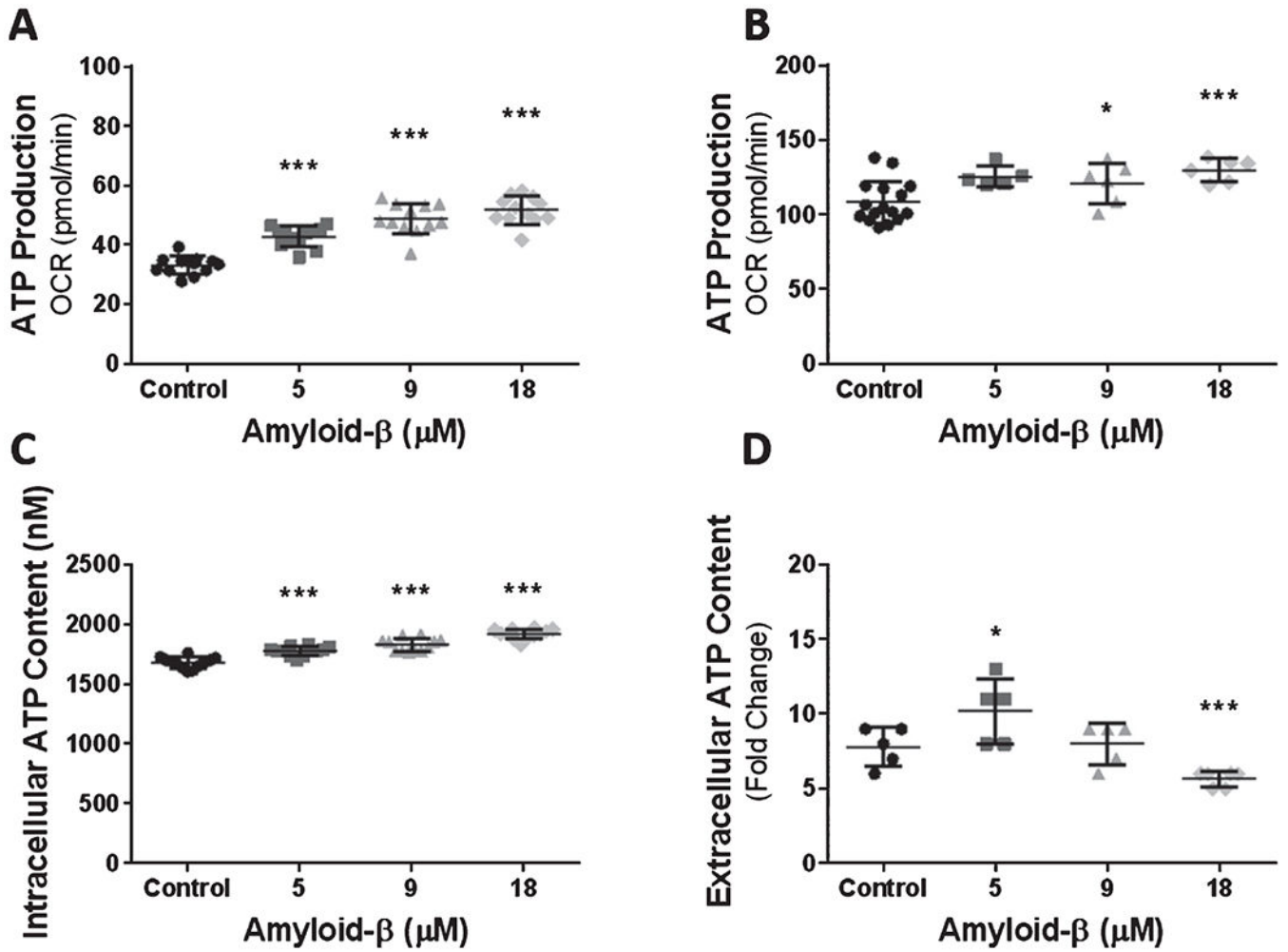


Fig. 4. Exposure to Aβ causes a dose-dependent increase in cellular ATP production and cytosolic ATP content. Scatter plot(s) (mean ± SD) depicting mitochondrial oxygen consumption (pmol/min) for ATP production in (A) bEnd.3 (*n* = 12 wells per group) and (B) primary cerebrovascular endothelial cells after 24 h exposure to vehicle control, 5 μM Aβ₁₋₄₂, 9 μM Aβ₁₋₄₂, and 18 μM Aβ₁₋₄₂ (control = 15, else = 6 wells per group). Scatter plot(s) (mean ± SD) demonstrating the (C) intracellular (*n* = 12 wells per group) and (D) extracellular (*n* = 5 wells per group) ATP content (nM) in bEnd.3 cells after exposure to vehicle control, 5 μM Aβ₁₋₄₂, 9 μM Aβ₁₋₄₂, and 18 μM Aβ₁₋₄₂. Statistical analysis was performed by One-way ANOVA with Dunnett’s post analysis and linear trend analysis to determine the level of significance (**p* = 0.01; ***p* = 0.001; ****p* = 0.0001).

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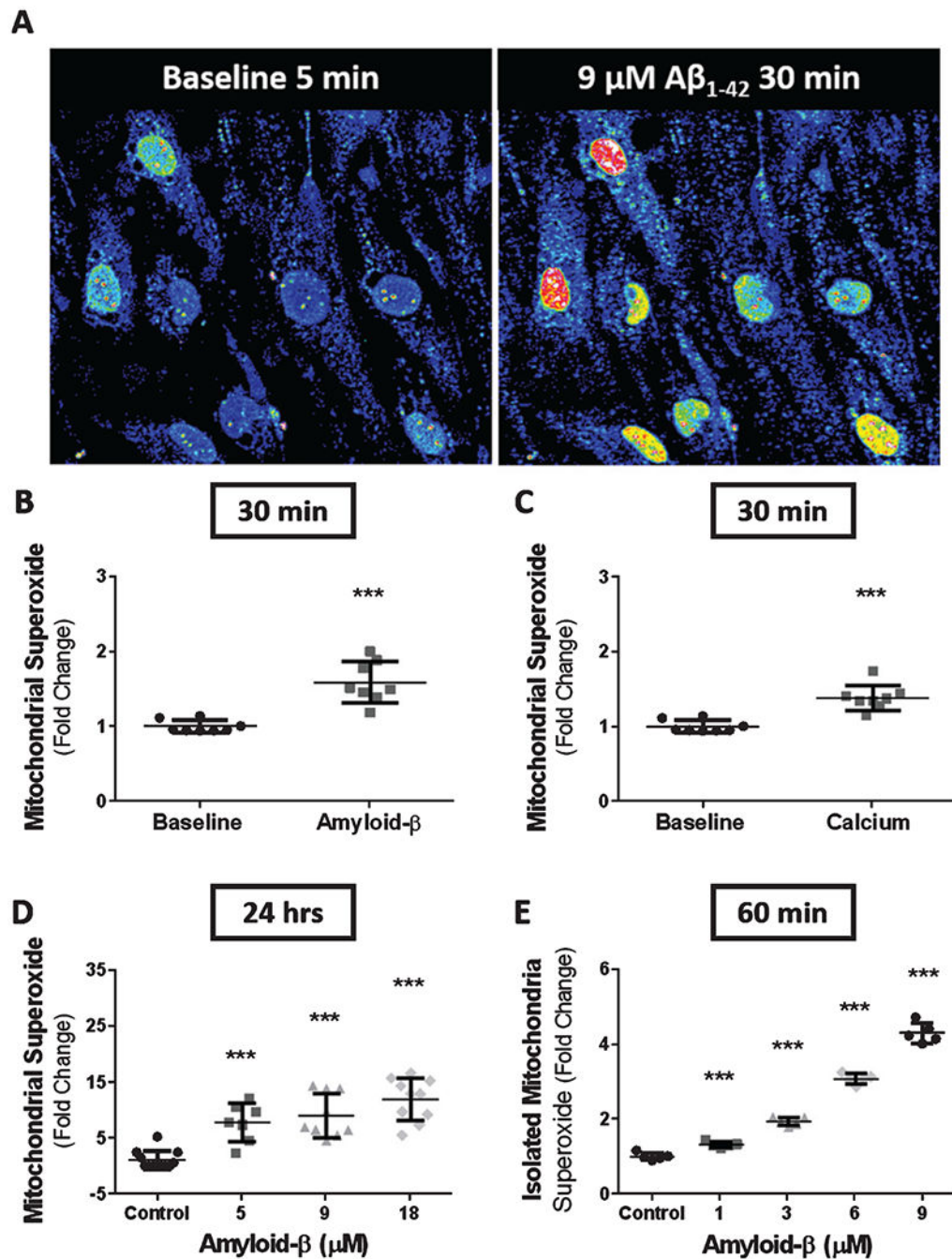


Fig. 5. Exposure to Aβ or Ca²⁺ results in an increase in mitochondrial superoxide production. A) Time-lapse confocal micrographs depicting mitochondrial superoxide production in bEnd.3 cells at baseline (left) and after 30 min of exposure to 9 μM Aβ₁₋₄₂ (right). Scatter plot(s) (mean ± SD) demonstrating superoxide production from time-lapse images as fold-change from baseline for bEnd.3 cells exposed to (B) 9 μM Aβ₁₋₄₂ (*n* = 8 cells) and (C) 5mM CaCl₂ (*n* = 8 cells) for 30 min. D) Scatter plot (mean ± SD) demonstrating mitochondrial superoxide production as fold-change in bEnd.3 cells after 24 h of exposure to vehicle, 5 μM

A β_{1-42} , 9 μM A β_{1-42} , and 18 μM A β_{1-42} ($n = 10$ wells per group). E) Scatter plot (mean \pm SD) depicting superoxide production as fold-change by isolated mitochondria from primary cerebrovascular endothelial cells after 1 h exposure to 0, 1, 3, 6, and 9 μM A β_{1-42} ($n = 5$ wells per group). Statistical analysis was performed by one-way ANOVA with Dunnett's post analysis and linear trend analysis to determine the level of significance (***) ($p = 0.0001$).

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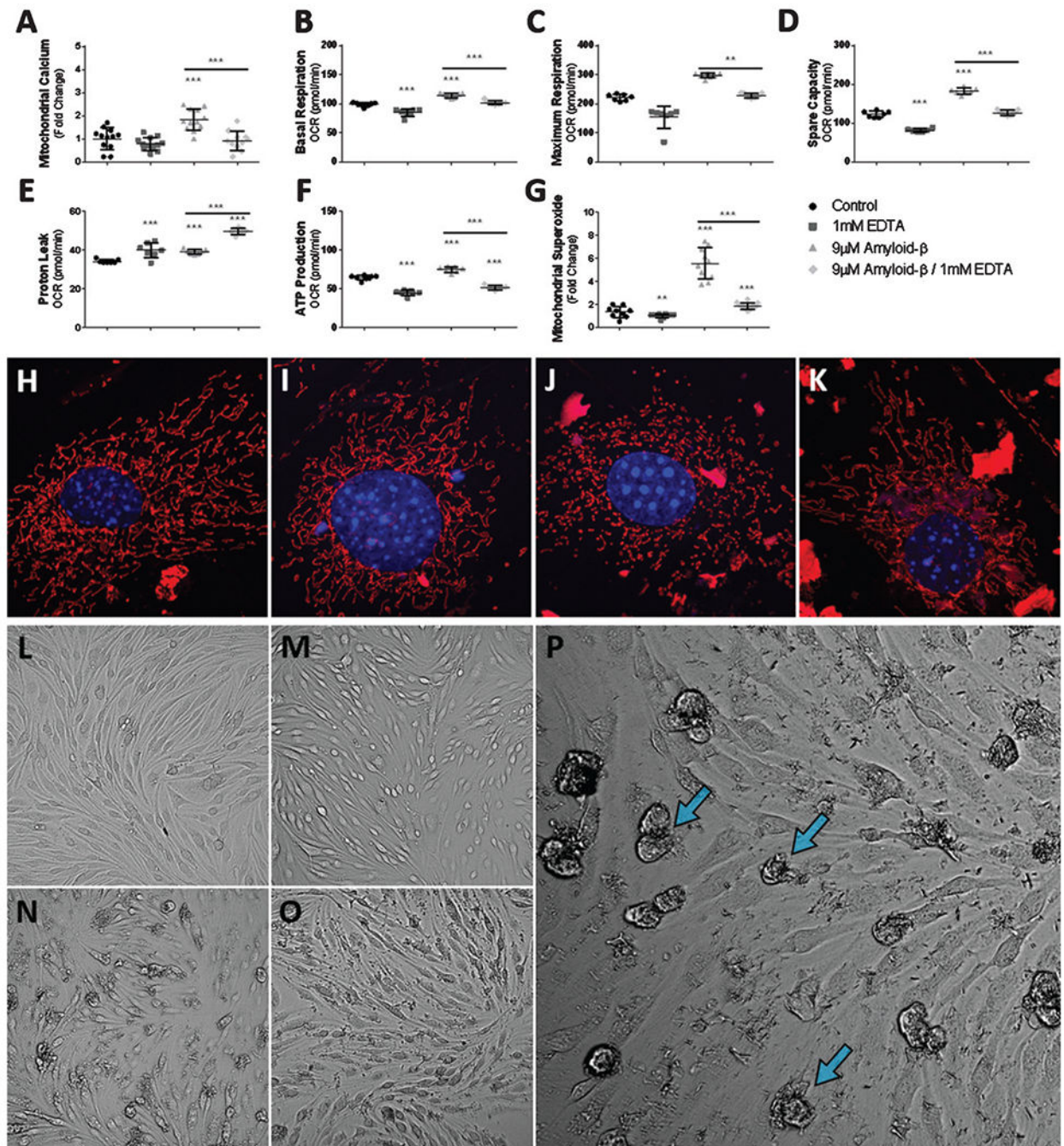


Fig. 6. Ca^{2+} chelation reduces the $\text{A}\beta$ induced hyper-oxidative state of mitochondria and superoxide production via reduced mitochondrial Ca^{2+} uptake. A) Scatter plot (mean \pm SD) demonstrating mitochondrial Ca^{2+} accumulation as fold-change in bEnd.3 cells after 24 h exposure to vehicle control, 1 mM EDTA, 9 μM $\text{A}\beta_{1-42}$, and 9 μM $\text{A}\beta_{1-42}$ with 1 mM EDTA ($n = 12$ per group). Scatter plot (mean \pm SD) demonstrating mitochondrial oxygen consumption (pmol/min) by (B) basal respiration, (C) maximum respiration, (D) spare capacity, (E) proton leak, and (F) ATP production from bEnd.3 cells after 24 h exposure to

vehicle control, 1 mM EDTA, 9 μM $\text{A}\beta_{1-42}$, and 9 μM $\text{A}\beta_{1-42}$ with 1 mM EDTA ($n = 7$ wells per group). G) Scatter plot (mean \pm SD) demonstrating mitochondrial superoxide production as fold-change in bEnd.3 cells after 24 h exposure to vehicle control, 1 mM EDTA, 9 μM $\text{A}\beta_{1-42}$, and 9 μM $\text{A}\beta_{1-42}$ with 1 mM EDTA ($n = 10$ wells per group). Confocal micrographs at 63x magnification depicting mitochondrial fragmentation (red) and nuclei (blue) in bEnd.3 cell exposed to (H) vehicle, (I) 1 mM EDTA, (J) 9 μM $\text{A}\beta_{1-42}$, and (K) 9 μM $\text{A}\beta_{1-42}$ with 1 mM EDTA. Note mitochondria morphology depicted in panel J, demonstrating numerous fragmented and punctated mitochondria while panel(s) H-I, K depict(s) complex and elongated mitochondria. Brightfield micrographs depicting bEnd.3 cell death (blue arrows) following exposure to 9 μM $\text{A}\beta_{1-42}$ after 24 h exposure to (L) vehicle control, (M) 1 mM EDTA, (N, P) 9 μM $\text{A}\beta_{1-42}$, and (O) 9 μM $\text{A}\beta_{1-42}$ with 1 mM EDTA. One-way ANOVA with Dunnett's post analysis was used to determine significance compared to control while a student's 2-tailed t-test was used to compare the mean(s) of the 9 μM $\text{A}\beta_{1-42}$ group and the 9 μM $\text{A}\beta_{1-42}$ with 1 mM EDTA experimental group (* $p < 0.05$; ** $p < 0.001$; *** $p < 0.0001$).

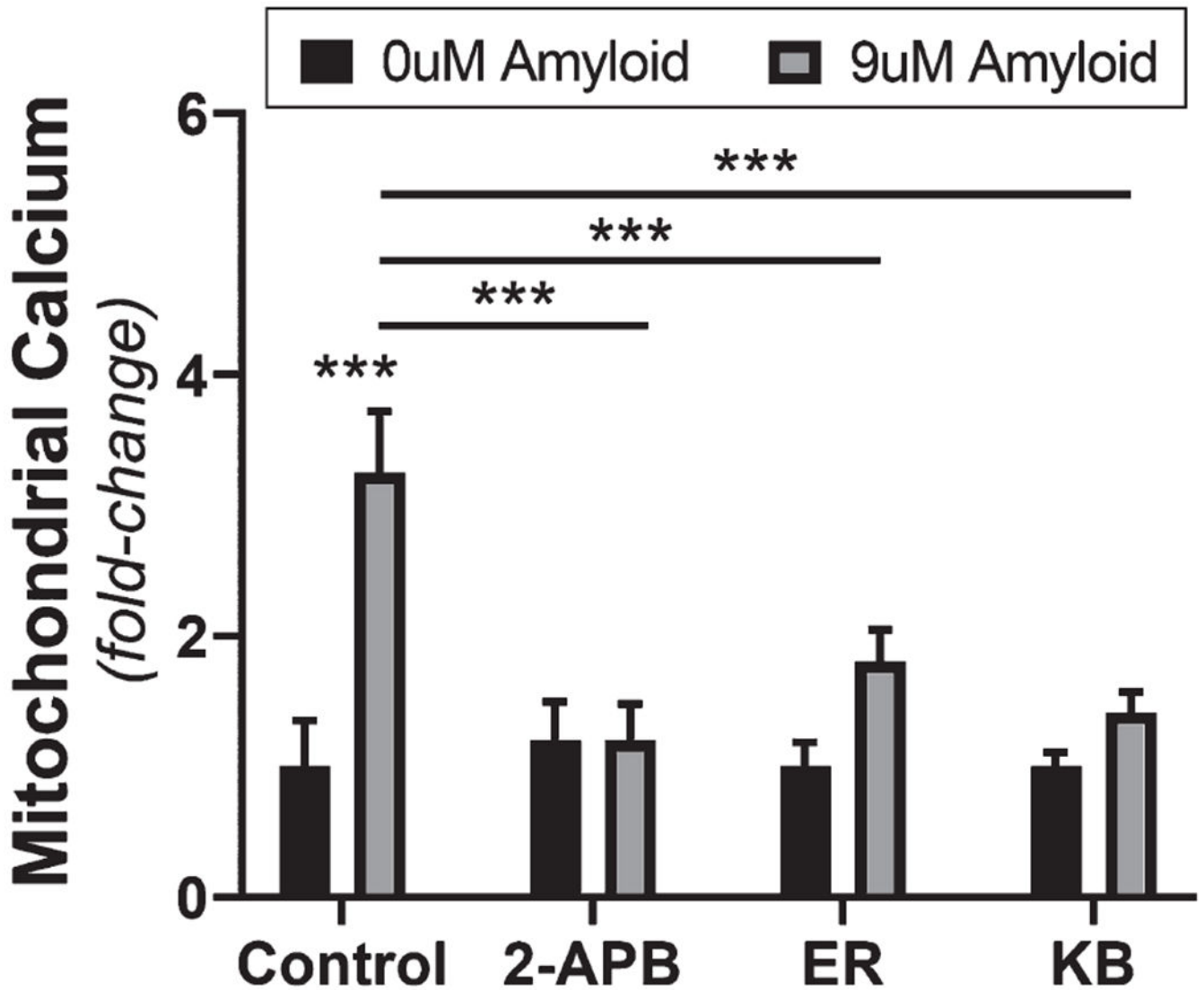


Fig. 7. Antagonism of regulatory intracellular calcium pathways attenuates the A β -induced mitochondrial calcium influx. Grouped bar graph (mean \pm SD) demonstrating mitochondrial Ca²⁺ accumulation as fold-change in bEnd.3 cells after 24 h exposure to vehicle control, 9 μ M A β ₁₋₄₂, and 9 μ M A β ₁₋₄₂ with either 50 μ M 2-APB, 5 μ M ER-000444793, or 7 μ M KB-R7943 ($n = 12$ per group). Student's 2-tailed t-test was used to compare the mean(s) of each experimental group (***) $p < 0.0001$.