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Inhibiting amyloid beta (1–42) peptide-induced mitochondrial dysfunction prevents the degradation of synaptic proteins in the entorhinal cortex

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Increasing evidence suggests that mitochondrial dysfunction and aberrant release of mitochondrial reactive oxygen species (ROS) play crucial roles in early synaptic perturbations and neuropathology that drive memory deficits in Alzheimer's disease (AD). We recently showed that solubilized human amyloid beta peptide 1-42 (hA β_{1-42}) causes rapid alterations at glutamatergic synapses in the entorhinal cortex (EC) through the activation of both GluN2A- and GluN2B-containing NMDA receptors. However, whether disruption of mitochondrial dynamics and increased ROS contributes to mechanisms mediating $hA\beta_{1-42}$ -induced synaptic perturbations in the EC is unknown. Here we assessed the impact of $hA\beta_{1-42}$ on mitochondrial respiratory functions, and the expression of key mitochondrial and synaptic proteins in the EC. Measurements of mitochondrial respiratory function in wild-type EC slices exposed to 1 μ M hA β_{1-42} revealed marked reductions in tissue oxygen consumption and energy production efficiency relative to control. $hA\beta_{1-42}$ also markedly reduced the immunoexpression of both mitochondrial superoxide dismutase (SOD2) and mitochondrial-cytochrome c protein but had no significant impact on cytosolic-cytochrome c expression, voltage-dependent anion channel protein (a marker for mitochondrial density/integrity), and the immunoexpression of protein markers for all five mitochondrial complexes. The rapid impairments in mitochondrial functions induced by $hA\beta_{1-42}$ were accompanied by reductions in the presynaptic marker synaptophysin, postsynaptic density protein (PSD95), and the vesicular acetylcholine transporter, with no significant changes in the degradative enzyme acetylcholinesterase. We then assessed whether reducing $hA\beta_{1-42}$ induced increases in ROS could prevent dysregulation of entorhinal synaptic proteins, and found that synaptic impairments induced by $hA\beta_{1-42}$ were prevented by the mitochondria-targeted antioxidant drug mitoquinone mesylate, and by the SOD and catalase mimetic EUK134. These findings indicate that $hA\beta_{1-2}$ can rapidly disrupt mitochondrial functions and increase ROS in the entorhinal, and that this may contribute to synaptic dysfunctions that may promote early AD-related neuropathology.

KEYWORDS

Alzheimer's disease, mitochondria, acetylcholine, entorhinal cortex, oxidative stress, reactive oxygen species, synaptic proteins

Introduction

Alzheimer's disease (AD) is a debilitating neurodegenerative disorder that results from the progressive loss of neurons in selective brain areas, and is the primary cause of dementia worldwide (Fan et al., 2020; Guo et al., 2020; Olajide et al., 2021). Neuropathology in AD is hallmarked by abnormal accumulation of amyloid beta peptide (A β), a highly neurotoxic derivative of the amyloid precursor protein (APP) which is a transmembrane protein that is particularly concentrated in synapses (Haass and Selkoe, 2007; Masters and Selkoe, 2012; Mucke and Selkoe, 2012; Hampel et al., 2021). AB is thought to interact with multiple biological mechanisms to incite AD-related modifications in neurons, and several lines of experimental evidence, including our own, have shown that AB can rapidly dysregulate proteins mediating excitatory synaptic transmission (Selkoe, 2002; Coleman and Yao, 2003; Shankar and Walsh, 2009; Overk and Masliah, 2014; Tu et al., 2014; Olajide and Chapman, 2021; Olajide et al., 2021). These synaptic changes are thought to lead to cognitive deficits and memory failure early in AD (Arendt, 2009; Mucke and Selkoe, 2012; Marsh and Alifragis, 2018; Guo et al., 2020), but the molecular mechanisms through which Aß induces excitotoxicity and neurodegeneration are still poorly understood.

Synaptic transmission relies heavily on mitochondria that generate energy through ATP and nicotinamide adenine dinucleotide (NAD +), maintain calcium homeostasis and buffering, and regulate cell signaling (Calkins et al., 2011; Akhter et al., 2017; Pickett et al., 2018). Perturbation of mitochondrial functions may contribute directly to impaired synaptic transmission in early AD (Cavallucci et al., 2013; Akhter et al., 2017; Olajide et al., 2017a; Xiao et al., 2017; Pickett et al., 2018; Khosravi and Harner, 2020), and A\beta can disrupt mitochondrial energy production and lead to increases in the synthesis of reactive oxygen species (ROS) and oxidative damage (Calkins et al., 2011; Hampel et al., 2021; Olajide et al., 2021; Ashleigh et al., 2022). Cellular ROS are natural by-products of mitochondrial aerobic respiration that result predominantly from leakage of electrons at complexes I and III of the electron transport chain. While ROS serve essential cellular functions, excess ROS can induce oxidative damage and AD-related neuropathology (Murphy, 2009; Patten et al., 2010; Wang et al., 2020). Under physiological conditions, oxidative damage by the major ROS, superoxide (O^2-) , is prevented by mitochondrial superoxide dismutase (SOD2), an antioxidant enzyme that catalyzes the dismutation of O^2 - to hydrogen peroxide (Murphy, 2009). However, AB can induce mitochondrial dysfunction both by increasing O^2 – production and by depleting the cellular antioxidant defense system, which causes degeneration of synaptic elements and alters neurotransmission processes in neurons (Tönnies and Trushina, 2017; Wang et al., 2020; Ionescu-Tucker and Cotman, 2021; Misrani et al., 2021). The overexpression of SOD2 in AD mouse models reportedly reduces AB deposition and prevents memory deficits (Dumont et al., 2009; Massaad et al., 2009), whereas mutant AD mice with depleted SOD2 expression show increased Aß levels and accelerated synaptic dysfunction and cognitive decline (Li et al., 2004; Esposito et al., 2006).

The entorhinal cortex (EC) is important for cognitive functions including memory and is among the first cortical regions to be affected by AD pathology (van Hoesen et al., 1991; Velayudhan et al., 2013; Khan et al., 2014; Zhou et al., 2016; Grubman et al., 2019; Olajide et al., 2021). Subpopulations of neurons are differentially vulnerable to the toxicity of oxidative stress (Wang and Michaelis, 2010), and the EC is one of the earliest temporal lobe structures to show both oxidative impairment and AB accumulation (Nunomura et al., 2001; Terni et al., 2010; Olajide et al., 2021). Armand-Ugon et al. (2017) reported reduced expression of mitochondrial complexes I, II, IV, and V in the EC, but not in the frontal cortex, during the initial stages of AD, and this mitochondrial dysfunction and oxidative impairment may drive early synaptic failure in AD (Tönnies and Trushina, 2017). We have shown recently that human $A\beta_{1-42}$ (h $A\beta_{1-42}$) rapidly alters elements of glutamatergic synapses in the EC through activation of both GluN2A and GluN2B subunitcontaining NMDA receptors (Olajide and Chapman, 2021). This may drive the selective vulnerability of the EC to ADtype neurodegeneration (Olajide et al., 2021), but whether increased ROS and mitochondrial dysfunction contribute to the susceptibility of EC to early AD-related synaptic degeneration has not been explored.

Here, we investigated how mitochondrial proteins, respiratory function, and key synaptic proteins are affected in the EC of brain slices from wild-type rats following exposure to soluble $hA\beta_{1-42}$. We then assessed the ability of two novel and specific pharmacological ROS scavengers and inhibitors, the mitochondria-targeted antioxidant mitoquinone mesylate (MitoQ) and the SOD and catalase mimetic EUK134, to block the degradation of synaptic proteins induced by $hA\beta_{1-42}$ in the EC. Our results support the idea that mitochondrial alterations induced by $hA\beta_{1-42}$ are a central factor in early synaptic degeneration in the EC, and contribute to a more detailed understanding of the molecular mechanisms driving neurodegeneration in the EC during early AD.

Materials and methods

Animals and tissue preparation

Experiments were conducted according to the guidelines of the Canadian Council on Animal Care, and experimental procedures were approved by the Concordia University Animal Research Ethics Committee (Permit Number: 30000253). Six to ten-week-old male Long-Evans rats (Charles River) were housed and placed on a reverse 12 h: 12 h light-dark cycle, with free access to rat chow and water. Acute brain slices were obtained following isoflurane anesthesia and decapitation as previously described (Glovaci and Chapman, 2019). Brains were rapidly removed and cooled (4°C) in high-sucrose ACSF containing (in mM) 250 sucrose, 2 KCl, 1.25 NaH₂PO₄, 7 MgCl₂, 26 NaHCO₃, 0.5 CaCl₂ and 10 dextrose, saturated with 95% O₂ and 5% CO₂. Horizontal slices (400 µm thick) were obtained throughout the ventral to the dorsal extent of the brain in cooled highsucrose ACSF using a vibratome (Leica, VT1200). The EC was carefully excised from each slice, using a flat blade in contact with the vibratome blade (Paxinos and Watson, 1997; Olajide and Chapman, 2021). Obtained tissue was placed in normal ACSF consisting (in mM) of 124 NaCl, 5 KCl, 1.25 NaH₂PO₄, 2 MgSO₄, 2 CaCl₂, 26 NaHCO₃, and 10 dextrose saturated with 95% O_2 and 5% CO_2 at 32°C for 30 min. Assignment of EC slices taken from the right or left hemispheres were alternated between treatment groups at each consecutive level, so that tissue in the control and treated conditions was obtained from both hemispheres across the dorso-ventral extent of the EC.

Drug preparation and treatments

Drugs were prepared as stock solutions and diluted to final concentrations just before use. Tissue was incubated for 3 h in submersion chambers containing ACSF at 22–24°C, saturated with 95% O₂ and 5% CO₂. EC slices were incubated in 1 μ M hA β_{1-42} (MW 4514.08; Abcam, AB120301) with a final

concentration of DMSO of 0.1%, while control EC slices were exposed to 0.1% DSMO in ACSF. Solubilization and preparation of hA β_{1-42} were done as previously described (Olajide and Chapman, 2021). hA β_{1-42} was first solubilized in DMSO at 500 μ M, sonicated for 15 min at room temperature, and then centrifuged at 15,000 × g at 4°C for 20 min. The supernatant was stored at -80°C in 10 μ L aliquots and diluted in ACSF just before use. This method is known to result in low molecular weight β -oligomers including monomers to tetramers (Bitan et al., 2003; Stine et al., 2003) which are the most neurotoxic (Masters and Selkoe, 2012; Guo et al., 2020). It is possible for some protofibrils to develop following this preparation, but significant fibrillary aggregation requires longer incubation times and higher concentrations (>10 μ M; O'Nuallain et al., 2004; Wogulis et al., 2005).

The role of mitochondrial ROS and oxidative stress was assessed by applying the mitochondria-targeted antioxidant drug mitoquinone mesylate (MitoQ, 500 nM; Toronto Research Chemicals; M372215), and SOD/catalase mimetic drug EUK134 (250 nM; Cayman Chemical; 10006329) during incubation of slices in hA β_{1-42} or DMSO. Each treatment group included slices from at least 6 animals.

Mitochondrial respiration measurements

A sequential substrate addition protocol was conducted to assess mitochondrial coupled and uncoupled oxygen consumption, LEAK respiration, and membrane integrity using a two-chamber polarographic sensor (Oxygraph-2k; Oroboros Instruments, Innsbruck, Austria). $hA\beta_{1-42}$ -incubated and control EC samples were assessed simultaneously in both chambers under similar reaction conditions (n = 5 or 6). The measurements of oxygen consumption were performed in MiR05 at 37°C. MiR05 contains (in mM) 0.5 EGTA, 3.0 MgCl₂·6H₂O, 60 K-lactobionate, 20 taurine, 10 KH₂PO₄, 20 HEPES, 110 sucrose, and 1 g/L BSA (pH 7.1). The oxygen flux was registered and analyzed by the DatLab 7.0 software. Between 2 and 3 mg (wet weight) of treated entorhinal tissue was placed in the Oxygraph and oxygen levels were increased to approximately 480 pmol. Saponin (50 µg/mL) was added to the chambers to permeabilize the tissue before the experiment was begun. In the protocol, non-phosphorylating LEAK-respiration was induced by adding the Complex I-linked substrates malate (2 mM), pyruvate (5 mM), and glutamate (5 mM). Subsequently, the OXPHOS-capacity of Complex I-linked activity was measured after the addition of a saturating concentration of ADP (5 mM). Cytochrome c (10 µM) was then added to assess mitochondrial membrane damage. OXPHOScapacity with combined Complex I and II-linked substrates was assessed by the addition of succinate (10 mM). This was followed by FCCP (carbonylcyanide-4 (trifluoromethoxy)

phenylhydrazone, 1 $\mu M)$ to test for uncoupling. The acceptor control ratio (ACR), which measures the degree of coupling between oxidation and phosphorylation, was calculated as the rate of oxygen consumption during ADP phosphorylation divided by the rate of non-phosphorylating LEAK-respiration induced by glutamate administration.

Protein preparation and Western blotting

To perform Western immunoblotting, treated EC tissue was collected into microfuge tubes and snap-frozen. Tissues were disrupted with a tissue sonicator (QSonica: Q55) in homogenization buffer (10 mM Tris, pH 7.4, 1 mM EGTA, 1 mM EDTA, 0.5 DTT, 0.1 µM okadaic acid, 1 mM Na₃VO₄, 1 mM PMSF and 10 µg/mL leupeptin). Mitochondrial protein purification was done following extraction in mitochondrial homogenization buffer (10 mM Tris-HCl, pH 6.7, 10 mM KCl, 0.15 mM MgCl₂, 1 mM PMSF, 1 mM DTT) followed by resuspension in mitochondrial suspension buffer (10 mM Tris HCl-pH 6.7, 0.15 mM MgCl₂, 0.25 mM sucrose, 1 mM PMSF, 1 mM DTT). The quantity of protein in each sample was determined using BCA Protein Assay (Thermo Fisher: 23,227) and an ELISA Fluorostar Analysis System Plate Reader. Bovine serum albumin (BSA) was used as the standard for protein quantification. Protein samples (20-30 µg) were resolved on Tris-glycine 8-12% SDS-PAGE gels. The resolved proteins were transferred from gels to nitrocellulose membrane (Bio-Rad: 1620112) and blocked for 1-2 h in either 5% milk or 5% BSA (as indicated below for each specific antibody) in Tris-buffered saline (TBS) containing 0.2% Tween-20 (TBST). All antibodies used were solubilized (0.5-2.0 mg/mL) by manufacturers in buffered saline or culture supernatant containing 0.02-0.1% sodium azide (pH 7.2-7.6). Primary antibodies were diluted for overnight incubation of membranes at 4°C, and included total oxidative phosphorylation (OXPHOS) rodent antibody cocktail (1:2,000, 5% milk in TBST, MitoSciences, MS604), rabbit anti-SOD2 antibody (1:4,000, 5% milk in TBST; Proteintech, 24127-1-AP), rabbit anti-Cytochrome c antibody (1:2,000, 5% milk in TBST; Abcam, AB133504), mouse anti-VDAC-1/Porin antibody (1:2,000, 5% BSA in TBST, Abcam, AB14734), mouse anti-synaptophysin (1:3,000, 5% BSA in TBST; Sigma-Aldrich, MAB5258), rabbit anti-PSD95 (1:3,000, 5% milk in TBST; Abcam, AB18258), rabbit anti-choline acetylcholinesterase (1:1,000, 5% milk in TBST; Abcam, AB183591), rabbit antivesicular acetylcholine transporter (1:2,000, 5% milk in TBST; Abcam, AB235201), mouse anti-vinculin (1:4,000, 5% milk in TBST; Abcam, AB130007), and mouse anti-β-Actin (1:5,000, 5% BSA in TBST; Abcam, AB8226). Membranes were then washed 3 times for 5 min each in TBST and incubated at room temperature with either peroxidase-conjugated goat antimouse secondary antibody (1:4,000; Millipore Sigma, AP124P) or peroxidase-conjugated goat anti-rabbit secondary antibody (1:5,000; Millipore Sigma, AP132P) for 1–2 h. Immunoreactivity was detected using ECL Western blotting substrate (Thermo Fisher Scientific, 32106) and visualized using a CDP-STAR chemiluminescence system (Amersham hyperfilm ECL). All antibody signals were normalized against loading control (β -Actin or vinculin) immunoreactivity. Western blot data were compiled from six animals, and bands were quantified by densitometric analysis using Image-J software (version 1.41).

Data analysis

Results obtained from mitochondrial respiratory measurements were tested statistically using a two-tailed Student's *t*-test. Data from Western blot densitometry were analyzed using either two-tailed Student's *t*-tests or one-way analyses of variance (ANOVA) with Sidak's multiple comparisons tests. All data analyses were done using GraphPad Prism software version 8.0.1 with significance set at p < 0.05. Bar graphs indicate the mean and standard deviation, normalized to the largest control value in percentage (for protein analysis), and include plots showing values obtained from individual animals.

Results

Effects of $hA\beta_{1-42}$ on entorhinal mitochondrial function and integrity

During the pathogenesis of AD, the EC selectively accumulates neurotoxic $A\beta$ and shows the earliest signs of oxidative impairment and neurodegeneration (Terni et al., 2010; Olajide et al., 2021), but the mechanisms involved are still unclear. We therefore examined the rapid effects of $hA\beta_{1-42}$ on mitochondrial functions in entorhinal tissue using highresolution respirometry. Utilization of respiration substrates in entorhinal samples was significantly reduced by exposing brain slices to 1 μ M hA β_{1-42} for a period of 3 h (Figure 1A1). Relative to control, $hA\beta_{1-42}$ markedly reduced respiration in EC slices (in pmol/s/mg; n = 5-6) following the addition of glutamate (2.9 \pm 1.2 vs. 5.6 \pm 1.0 in controls; p = 0.0211), ADP (3.8 \pm 3.3 vs. 20.2 \pm 2.4; p = 0.0022), cytochrome c $(4.2 \pm 3.7 \text{ vs. } 19.4 \pm 2.7; p = 0.0036)$, succinate $(13.7 \pm 5.9 \text{ vs.})$ 36.0 ± 4.7 ; p = 0.0045), and FCCP (11.7 ± 5.0 vs. 27.0 ± 4.0 ; p = 0.0070). However, incubation of EC slices with hA β_{1-42} did not significantly alter the consumption of malate (0.7 \pm 0.2 vs. 1.1 \pm 0.5 in controls; *p* = 0.2427), and pyruvate (1.8 \pm 1.2 vs. 3.5 \pm 0.6; *p* = 0.0570). hA β_{1-42} also markedly reduced the ratio between oxidation and phosphorylation which measures the efficiency of the mitochondria in generating ATP per given amount of oxygen, and is shown by assessments of the ACR $(1.1 \pm 0.6 \text{ vs. } 3.6 \pm 0.4 \text{ in controls}; p < 0.0001; Figure 1A2).$



with vinculin serving as a loading control. (C2) Graphs showing normalized data for all five mitochondrial subunits in hA β_{1-42} -treated slices vs. the largest expression in the control group (n = 6) (*p < 0.05; **p < 0.01; ****p < 0.001).

We next assessed the impact of $hA\beta_{1-42}$ on immunoexpression of markers for the integrity of the mitochondrial membrane and for the five mitochondrial

protein complexes that make up the electron transport chain. Compared to control, $hA\beta_{1-42}$ treatment did not alter immunoblot expression of the mitochondrial membrane

and gatekeeper protein voltage-dependent anion channel 1 (VDAC1; 91.9 \pm 7.3 vs. 94.3 \pm 7.8 in controls; p = 0.6125) (Figure 1B). Similarly, immunoblotting of complexes I to V with total OXPHOS rodent antibody cocktail (Figure 1C) revealed no significant changes (n = 6) in protein expression between control and $hA\beta_{1-42}$ -treated entorhinal slices. When compared to control, $hA\beta_{1-42}$ (1 μM) treatment for 3 h did not alter the normalized protein expression of Complex V (93.2 \pm 4.6 and 94.0 \pm 5.0; p = 0.785), Complex III (90.5 \pm 7.4 and 91.4 \pm 7.0; *p* = 0.833), Complex IV (94.2 \pm 4.6 and 94.1 \pm 5.3; *p* = 0.961), Complex II $(95.1 \pm 3.6 \text{ and } 93.4 \pm 3.5; p = 0.560)$, and Complex I (91.1 \pm 9.3 and 94.0 \pm 4.3; *p* = 0.589) in entorhinal lysates. Therefore, although $hA\beta_{1-42}$ treatment reduces mitochondrial function as reflected in respirometry analysis, it did not significantly affect the expression of protein markers for mitochondrial integrity and for respiratory complexes.

$hA\beta_{1-42}$ reduces immunoexpression of mitochondrial enzymes, synaptophysin, postsynaptic density protein-95 and vesicular acetylcholine transporter

Our finding that $hA\beta_{1-42}$ rapidly impairs mitochondrial respiration in the EC suggests that a resulting oxidative redox imbalance, due to increased leakage of mitochondrial electrons and ROS formation, could have deleterious effects on the mitochondrial antioxidant system. We therefore characterized the expression of the mitochondrial oxidative scavenger enzyme SOD2, and both mitochondrial and cytosolic cytochrome c protein. Representative immunoblots and quantification data show that, compared to control, $hA\beta_{1-42}$ markedly reduced the relative expression of SOD2 (53.3 \pm 14.0 vs. 90.4 \pm 8.1 in controls; p = 0.0002) (Figure 2A). Mitochondrial cytochrome c expression was also reduced by application of $hA\beta_{1-42}$ $(33.6 \pm 17.2 \text{ vs. } 86.0 \pm 12.3 \text{ in controls}; p = 0.0001)$, and this effect was specific to mitochondria because immunoexpression of cytosolic cytochrome c was not significantly affected $(81.7 \pm 13.6 \text{ vs. } 89.6 \pm 11.3 \text{ } p = 0.298).$

A reduction in the efficiency and energy production capacity of the mitochondrial electron transport chain, increased mitochondrial ROS, and oxidative stress could impact both pre- and post-synaptic elements, and we therefore evaluated the effects of $hA\beta_{1-42}$ on the presynaptic marker synaptophysin, and the postsynaptic density protein 95 (PSD95). We found that $hA\beta_{1-42}$ -induced mitochondrial dysregulation was accompanied by a marked reduction in relative immunoexpression of both PSD95 (91.9 ± 6.6 and 26.4 ± 11.3; p < 0.0001) and synaptophysin (87.5 ± 12.2 and 31.3 ± 21.8; p = 0.0003) (Figure 2B).

Reductions in cholinergic function in the EC is one of the earliest manifestations of AD (Hampel et al., 2018), and we therefore assessed the effects of $hA\beta_{1-42}$ on the expression of the vesicular acetylcholine transporter (VAChT) and the degradative enzyme acetylcholinesterase (AChE). Although $hA\beta_{1-42}$ did not cause a significant change in AChE immunoexpression when compared to control (97.4 ± 18.2 vs. 91.5 ± 8.7 in controls; p = 0.5433), $hA\beta_{1-42}$ markedly reduced the expression of VAChT in entorhinal lysates vs. control (60.0 ± 16.6 vs. 87.7 ± 12.7; p = 0.0312) (Figure 2C), suggesting a disruption in the function of cholinergic terminals in the EC.

Mitochondria-targeted antioxidant mitoquinone mesylate and EUK134 reduce $hA\beta_{1-42}$ -induced impairments in synaptic proteins

To determine if increased ROS associated with mitochondrial dysregulation is related to the reductions in mitochondrial enzymes and synaptic proteins induced by $hA\beta_{1-42}$, we tested the ability of two novel, structurally distinct ROS scavengers (MitoQ and EUK134) to block protein reductions induced by $hA\beta_{1-42}$. The mitochondria-targeted antioxidant MitoQ administered alone had no significant effect on the expression of mitochondrial enzymes, but was found to block changes in both SOD2 and mitochondrial cytochrome c induced by $hA\beta_{1-42}$ (compare Figure 3A and Figure 2A). There was no significant main effect of treatment on the relative expression of SOD2 between control slices (92.2 \pm 6.4) and slices incubated with MitoQ alone (90.4 \pm 15.6) or MitoQ with $hA\beta_{1-42}$ (88.7 ± 8.9) [$F_{(2, 15)} = 0.15$, p = 0.8646]. Similarly, the expression of mitochondrial cytochrome c protein was not significantly different between control slices (90.7.0 \pm 8.3) and slices treated with either MitoQ (94.5 \pm 9.8) or MitoQ and $hA\beta_{1-42}$ (88.0 ± 11.4) [$F_{(2, 15)} = 0.65$, p = 0.5355] (Figure 3A).

We then assessed if blocking excess ROS induced by $hA\beta_{1-42}$ using MitoQ would also block reductions in the immunoexpression of PSD95, synaptophysin, and the vesicular acetylcholine transporter (VChAT). The application of MitoQ had no effect alone, but it prevented reductions in all three proteins induced by $hA\beta_{1-42}$ (compare **Figure 3B** and **Figures 2B,C**). There were no significant main effects of treatment (n = 6) on the expression of PSD95 [$F_{(2, 15)} = 0.14$, p = 0.8742], synaptophysin [$F_{(2, 15)} = 0.31$, p = 0.7386], and VChAT [$F_{(2, 15)} = 1.66$, p = 0.2226]. Preventing increases in ROS induced by $hA\beta_{1-42}$ can therefore prevent degradation of key synaptic proteins.

The idea that increases in mitochondrial ROS induced by $hA\beta_{1-42}$ mediate synaptic degeneration in entorhinal slices was further tested by determining if the synthetic SOD and catalase



cholinergic markers acetylcholinesterase (AChE), vesicular acetylcholine transporter (VAChT), and β -actin (loading control). (C) Bar graphs showing normalized expression of AChE and VAChT in hA β_{1-42} -treated slices and control (n = 6) (**p < 0.01; ***p < 0.005; ****p < 0.0001).

mimetic EUK134 could block the reductions in synaptic and mitochondrial proteins induced by $hA\beta_{1-42}$. Application of EUK134 blocked reductions in both SOD2 and mitochondrial

cytochrome c induced by $hA\beta_{1-42}$ (compare Figure 4A and Figure 2A), and there were no significant main effects of treatment with EUK134, or with EUK134 and $hA\beta_{1-42}$, (*n* = 6),



on the relative expression of SOD2 $[F_{(2, 15)} = 0.95, p = 0.4084]$ or mitochondrial cytochrome c $[F_{(2, 15)} = 0.61, p = 0.5578]$.

We also found that EUK134 rescued $hA\beta_{1-42}$ -induced reductions in immunoexpression of PSD95 and VChAT, and partially rescued $hA\beta_{1-42}$ -induced reductions in the presynaptic protein synaptophysin (compare Figure 4B and Figures 2B,C). There was no significant effect of treatment with EUK134 or EUK134 and $hA\beta_{1-42}$ on the relative expression of PSD95 $[F_{(2, 15)} = 0.04, p = 0.9650]$ and VChAT $[F_{(2, 15)} = 2.83, p = 0.0905]$. There was, however, a significant main effect of treatment on synaptophysin immunoexpression $[F_{(2, 15)} = 4.44, p = 0.0305]$ in which the expression of synaptophysin was significantly reduced in slices treated with EUK134 + hA β_{1-42} relative to control slices (69.8 \pm 21.6 vs. 92.7 \pm 5.7 in controls; p = 0.0480), but not relative to slices treated with EUK134 alone (90.4 \pm 12.1; p = 0.0811). The mean expression of synaptophysin following treatment with $hA\beta_{1-42}$ in the presence of EUK134 (69.8 \pm 21.6) was greater than that following treatment with $hA\beta_{1-42}$ alone $(31.3\pm21.8;$ Figure 2B), however, suggesting that EUK134 was partially protective in preventing reductions in synaptophysin. These findings suggest that $hA\beta_{1-42}$ -induced rapid synaptic impairments in entorhinal tissue can be significantly rescued by inhibiting mitochondrial ROS levels.

Discussion

A previous report from our lab has shown that $hA\beta_{1-42}$ causes rapid degeneration of presynaptic and postsynaptic elements in the EC through activation of GluN2A- and GluN2B- containing NMDA glutamate receptors (Olajide and Chapman, 2021). The selective vulnerability of the EC to early AD-related synaptic damage and neurodegeneration (Du et al., 2004; Velayudhan et al., 2013; Zhou et al., 2016), may reflect the dysregulation of neural mechanisms including NMDA glutamate receptors that contribute to learning and memory (Olajide et al., 2021). A β can accumulate in mitochondria where



it impairs mitochondrial dynamics and upregulates oxidative stress by impairing mitochondrial respiratory function and the production of adenosine triphosphate (ATP) (Wang et al., 2014, 2020; Ashleigh et al., 2022). Further, activation of NMDA receptors by $hA\beta_{1-42}$ can lead to increases in calcium influx, the rapid facilitation of AMPA glutamate receptor responses, and hyperexcitability that increases metabolic demands on mitochondria, ultimately culminating in oxidative stress and synaptic dysfunction (Findley et al., 2019; Guo et al., 2020; Wang et al., 2020). We have found here that exposure to 1 μ M $hA\beta_{1-42}$ for a period of 3 h markedly reduces mitochondrial respiratory function and integrity in the EC. Further, the resulting increase in ROS appears to be a major driver of the rapid degeneration of both pre- and post-synaptic proteins in the EC because the ROS scavengers Mito-Q and EUK134 strongly inhibited reductions in synaptophysin and PSD-95. The reductions in synaptic proteins observed here imply a reduction in number of synapses, but it is not clear if this would be expressed following this relatively brief application of $hA\beta_{1-42}$. However, these results point to a major role of mitochondrial dysregulation in the synaptic degeneration induced by $hA\beta_{1-42}$, which may contribute to the early progression of AD in the EC (Olajide et al., 2021). The present results may reflect mitochondrial dysfunction in glia as well as neurons, and mitochondrial dysfunction in glia is also thought to be a major contributor to the progression of AD (Mulica et al., 2021).

In our experimental model we have applied moderate concentrations of $A\beta$ to wildtype slices for 3 h, but the progression of AD involves circulation of lower concentrations of $A\beta$ over greatly extended periods of time. Although our model may reflect degenerative mechanisms that are expressed early in the progression of AD, and reflect the early selective vulnerability of the EC to $A\beta$ toxicity (Olajide et al., 2021), the changes in mitochondrial function and synaptic proteins that we have observed might also occur in later stages of AD after prolonged exposure to lower concentrations of $A\beta$.

We have found here that $hA\beta_{1-42}$ markedly reduces mitochondrial-coupled oxygen consumption, and functional membrane integrity as reflected by oxygen use in the presence of cytochrome c. High-resolution respirometry showed no significant changes in non-phosphorylating LEAK respiration during the addition of the Complex I-linked substrates malate, pyruvate or glutamate, but the OXPHOS capacity of Complex I-linked activity was markedly reduced by $hA\beta_{1-42}$, as shown by the approximately fivefold reduction in oxygen utilization during the addition of ADP. Exposure to $hA\beta_{1-42}$ also impaired the function of Complex II, because the addition of succinate, which reflects the combined OXPHOS capacity of Complex I and Complex II, was also significantly reduced. The rapid defects in mitochondrial-related metabolic pathways provide direct evidence that impaired bioenergetic machinery in mitochondria is an early driver of AD-related pathology in the EC. Similarly, OXPHOS and associated pathways are significantly downregulated within the hippocampus during AD, and reductions in Complex I, III, and IV of OXPHOS have been reported in both early and definite AD brains (Manczak et al., 2004; Brooks et al., 2007).

Further, the indication that $hA\beta_{1-42}$ -induced impairments in entorhinal electron transport chain complexes can dysregulate bioenergetics is corroborated by our findings that the efficiency of the link between oxidation and the production of ATP by phosphorylation of ADP was reduced by $hA\beta_{1-42}$, as reflected by both reductions in the uncoupling capacity of the mitochondria during the addition of FCCP, and by a reduction in the ACR. The ACR expresses the rate of oxygen consumption during ADP phosphorylation relative to the rate of non-phosphorylating LEAK-respiration induced by glutamate. Our data corresponds with several reports indicating a large and consistent decline in mitochondrial substrate utilization and energy production in the neocortex during the prodromal stages of AD (Croteau et al., 2018; Wang et al., 2020), suggesting that mitochondrial bioenergetics dysfunction plays an early role in the pathogenesis of AD in the EC.

Mitochondrial cytochrome c transfers electrons to complex IV in the electron transport chain which is the primary site of cellular oxygen consumption (Timón-Gómez et al., 2018). We found that $hA\beta_{1-42}$ caused about a 2.5-fold reduction in mitochondrial cytochrome c function, suggesting alterations to the mitochondrial membrane in entorhinal tissue. We further found that, although total cytoplasmic cytochrome c protein expression was unaffected, the expression of mitochondrial cytochrome c was greatly reduced by treatment with $hA\beta_{1-42}$. An increase in cytosolic cytochrome c could be expected from previous work showing that A β can activate intrinsic apoptotic

pathways and the release of mitochondrial cytochrome c into the cytosol (Kim et al., 2014), but the lack of an increase in cytosolic cytochrome c in the present study may be due to the relatively brief duration of exposure to A β . Deficiency of cytochrome c oxidase (Complex IV) is the most common defect in the mitochondrial electron transport chain in AD, leading to an increase in ROS production, a decrease in energy stores, and a disruption of energy metabolism (Rak et al., 2016). We found a marked reduction in the expression of SOD2, a key antioxidant enzyme that scavenges superoxide in mitochondria, and this is consistent with a rapid increase in mitochondrial ROS production induced by hA β_{1-42} that depletes the energy metabolism machinery in the EC.

Although $hA\beta_{1-42}$ significantly reduced respiration and bioenergetics, it did not alter the expression of protein markers for the five mitochondrial Complexes or the expression of the mitochondrial voltage-dependent anion channel 1 (VDAC1). VDAC1 is the most abundant protein on the outer membrane of mitochondria, and it serves as a gatekeeper for the passage of metabolites, and is crucial for the metabolic functions of mitochondria (Camara et al., 2017). It is well established that the expression of mitochondrial proteins and genes are among the most prominent changes in the cortex during AD (Cottrell et al., 2001; Valla et al., 2001; Minjarez et al., 2016; Adav et al., 2019), and the lack of a significant effect of $hA\beta_{1-42}$ on the expression of mitochondrial complex proteins observed here may reflect the resistance of these proteins to rapid degradation over the relativity short incubation time.

Preventing increases in reactive oxygen species inhibits hAβ₁₋₄₂-induced synaptic changes

Reductions in synaptic proteins induced by $hA\beta_{1-42}$ are likely to have resulted from increased ROS and oxidative stress in entorhinal neurons. Incubation in $hA\beta_{1-42}$ resulted in a marked reduction in SOD2, which is the primary antioxidant enzyme that scavenges superoxide in mitochondria, suggesting that SOD2 may have been depleted due to increases in ROS and overwhelming activity-dependent utilization of SOD2. The reduction in SOD2 may have further impaired ROS scavenging in entorhinal neurons thereby exacerbating oxidative damage and a loss of mitochondrial cytochrome c function. Excessive ROS and oxidative stress have been extensively linked to synaptic loss associated with aging and AD (Patten et al., 2010; Tönnies and Trushina, 2017; Guo et al., 2020), and increases in ROS associated with mitochondrial dysfunction that exceed the scavenging ability of entorhinal neurons is likely to have contributed to the observed loss of synaptic proteins. Consistent with our previous findings in the EC (Olajide and Chapman, 2021), both the presynaptic marker

synaptophysin and the postsynaptic marker PSD-95, were reduced by $hA\beta_{1-42}.$

The degeneration of cholinergic terminals in the EC is one of the earliest neuropathologies in AD (Francis et al., 1999; Gil-Bea et al., 2005; Hamam et al., 2007; Schaeffer and Gattaz, 2008). Although we found that the degradative enzyme acetylcholinesterase was not affected by $hA\beta_{1-42}$, we did observe a reduction in the vesicular acetylcholine transporter (VAChT), suggesting a reduced function in cholinergic terminals. Several other reports have also shown marked reductions in the expression of VAChT following A β infusion in rats and in AD models (Ikeda et al., 2000; Pákáski and Kálmán, 2008; Schliebs and Arendt, 2011; Hampel et al., 2018).

ROS-mediated oxidative stress is thought to have a major role in synaptic dysregulation and neurodegeneration in AD, and various antioxidants, and the genetic overexpression of SOD2, have been explored as therapeutic strategies for AD (Dumont et al., 2009; Massaad et al., 2009; Bonda et al., 2010; Olajide et al., 2017a,b, 2018; Tönnies and Trushina, 2017; Misrani et al., 2021). We co-incubated slices with $hA\beta_{1-42}$ and the ROS inhibitors MitoQ or EUK134 to determine if reducing oxidative stress could prevent the degeneration of synaptic proteins in the EC. Both MitoQ and EUK134 inhibited $hA\beta_{1-42}$ -induced reductions in both SOD2 and mitochondrial cytochrome c, likely by maintaining a sufficient ROS scavenging capacity in entorhinal neurons and preventing overwhelming demands on SOD2. In addition, co-incubation with MitoQ blocked hA_{β1-42}-induced decreases in synaptophysin, PSD-95, and VAChT, indicating that oxidative stress arising from insufficient mitochondrial capacity can drive degeneration of these synaptic proteins. Similarly, the ROS inhibitor EUK134 prevented reductions in PSD-95 and VAChT, and greatly attenuated reductions in synaptophysin induced by $hA\beta_{1-42}$. Others have also found that decreasing ROS using MitoQ and EUK134 can prevent synaptic degeneration induced by oxidative stress in cortical and hippocampal neurons (Ma et al., 2011; Mcmanus et al., 2011; Reddy et al., 2012; Sanmartín et al., 2017; Yu et al., 2018; Cenini and Voos, 2019). These studies suggest that the protective effect of ROS inhibition that we have observed following acute application of $hA\beta_{1-42}$ to wildtype slices might be similarly protective for the advancement of neurodegeneration in more chronic models of AD.

Conclusion

Mitochondrial dysfunction resulting in increased ROS production and oxidative stress precedes the formation of neuritic plaques and neurofibrillary tangles, and is thought to contribute substantially to the earliest stages of AD and the onset of cognitive decline and memory loss (Uttara et al., 2009; Wang et al., 2014; Tönnies and Trushina, 2017; Ashleigh et al., 2022).

We have shown here that short-term exposure of wild-type EC slices to $hA\beta_{1-42}$ results in deficits in mitochondrial respiration, and have provided evidence that oxidative stress associated with increased ROS is a major factor in the rapid degeneration of key pre- and post-synaptic proteins in the EC. Further, these mechanisms may be a major contributor to the reduction in cholinergic transmission in the EC, which is an early pathology thought to contribute to cognitive decline.

Data availability statement

The raw data supporting the conclusions of this article will be made available by the authors, without undue reservation.

Ethics statement

The animal study was reviewed and approved by Animal Research Ethics Committee, Concordia University.

Author contributions

OO, AB, and CC contributed to conception and design of the study. OO, AB, and CL acquired and analyzed the data. OO wrote the first draft of the manuscript. All authors contributed to manuscript revision and approved the submitted version.

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Conflict of interest

The authors declare that the research was conducted in the absence of any commercial or financial relationships that could be construed as a potential conflict of interest.

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