## Mitochondrial biogenesis mediated by melatonin in an APPswe/PS1dE9 transgenic mice model

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Alzheimer's disease (AD) is a chronic progressive neurodegenerative disease, but the pathogenesis is unclear. Damaged mitochondrial biogenesis has been observed in AD. Increasing evidence suggests that mitochondrial biogenesis is involved in the pathogenesis of AD, but the exact mechanism is unclear. In this study, we used the amyloid precursor protein Swedish mutations K594N/M595L (APPswe)/presenilin 1 with the exon-9 deletion (PS1dE9) transgenic mouse model of AD, which was successfully established by the expression of amyloid  $\beta$  precursor protein and presenilin 1 (PS1). Then, we compared APPswe/PS1dE9 transgenic mice with and without melatonin (MT) in drinking water for 4 months (estimated 0.5 mg/day) and control C57BL/6J mice without MT for expression of mitochondrial biogenesis factors (mitochondrial transcription factor A, nuclear respiratory factor 1 and 2, peroxisome proliferator-activated receptor  $\gamma$ coactivator 1-α), mitochondrial structure, mitochondrial DNA to nuclear DNA ratio, behavioral changes, and amyloid  $\beta$  $(A\beta)$  deposition and soluble A $\beta$  levels in the cerebral cortex and hippocampus. Compared with controls, APPswe/

#### Introduction

Alzheimer's disease (AD) is a chronic, progressive, irreversible neurodegenerative disorder pathologically characterized by amyloid  $\beta$  (A $\beta$ ) plaques, neurofibrillary tangles, and neuronal cell loss [1,2]. The most common early symptom is difficulty in remembering recent events (short-term memory loss) [3]. As the disease advances, symptoms can include problems with language, disorientation (including easily getting lost), mood swings, loss of motivation, and behavioral issues [4]. According to the Alzheimer's Disease International annual report, about 46.8 million individuals worldwide had dementia in 2015. In 2050, the estimated number of cases will be 131.5 million, 50–75% of them with AD.

Mitochondria are considered the energy provider in eukaryotic cell activity. Growing evidence indicates that mitochondrial dysfunction is the important cause of diseases such as neurodegenerative disease. Damage to mitochondrial PS1dE9 mice with long-term MT intake showed increased levels of mitochondrial biogenesis factors, alleviated mitochondrial impairment, enhanced mitochondrial DNA copy number, improved spatial learning and memory deficits, and reduced A $\beta$  deposition and soluble A $\beta$  levels. Defective mitochondrial biogenesis may contribute toward the damaged mitochondrial structure and function in AD. MT may alleviate AD by promoting mitochondrial biogenesis. *NeuroReport* 29:1517–1524 Copyright © 2018 The Author(s). Published by Wolters Kluwer Health, Inc.

NeuroReport 2018, 29:1517-1524

Keywords: Alzheimer's disease, APPswe/PS1dE9 transgenic mice, melatonin, mitochondrial biogenesis

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Received 31 July 2018 accepted 16 September 2018

biogenesis plays an important role in mitochondrial dysfunction. Mitochondrial biogenesis is regulated by the peroxisome proliferator-activated receptor  $\gamma$  coactivator 1- $\alpha$ nuclear respiratory factor 1 and 2-mitochondrial transcription factor A pathway (PGC-1 $\alpha$ -NRF-TFAM). Recent research showed decreased expression of PGC-1 $\alpha$ , NRF1, NRF2, and TFAM in the brains of AD patients and cells stably expressing amyloid  $\beta$  precursor protein (APP) [5]. In addition, the number of mitochondria was significantly reduced [6], which suggested that mitochondrial biogenesis was blocked in AD. An important cause of mitochondrial dysfunction in the process was impaired mitochondrial biogenesis [7].

Melatonin (MT) is a tryptophan metabolite synthesized mainly by the pineal gland and also synthesized by the mitochondria. It has many biological functions, such as regulating circadian rhythm, scavenging free radicals, improving immunity, and antioxidation [8]. MT was found to be significantly reduced in the brain of AD patients, and was found to be involved in sleep disorders with AD. In addition, MT plays an important role in the pathogenesis of AD [9]. Recent studies have shown that MT is synthesized in the pineal gland and also in other

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cells. The concentration of MT is higher in the mitochondria than in the plasma, cerebrospinal fluid, and peripheral blood [10]. MT can be transferred by peptide transporters 1 and 2. It is essential for maintaining the stability of mitochondrial morphology and function [10]. Finally, the pathological changes and cognitive function of APP/presenilin 1 (PS1) mice were found to be alleviated with oral MT treatment [11].

Increasing evidence suggests that mitochondrial biogenesis is involved in the pathogenesis of AD, but the exact mechanism is unclear. In this study, we treated APP/PS1 mice with MT to observe the expression of mitochondrial biogenesis factor, mitochondrial structure, mitochondrial DNA (mtDNA) to nuclear DNA (nDNA) ratio, behavioral changes, and A $\beta$  deposition and soluble A $\beta$  levels in the cerebral cortex and hippocampus. We aimed to analyze the role of mitochondrial biogenesis in the AD mouse model and the effect of MT on defective mitochondrial biogenesis and its underlying mechanisms.

#### Materials and methods Animals and tissues

Male mice were used in this study to avoid the possible influence of sex on A $\beta$  plaque formation in AD [12]. All mice were purchased from Beijing HFK Bio-Technology Co., Beijing, China. Mice were housed at 25±2°C with a relative humidity of 70±5°C in controlled rooms on a 12/12 h light, dark cycle. Treatment of mice followed the guidelines of the ethical committee for animal experiments of Shandong University (Jinan, China).

The APP/PS1 mouse is an AD mice model harboring mutant human transgenes of APP and presenilin 1 (PS1) [13]. A $\beta$  deposition appears in this mouse model early in life. Some research has shown that A $\beta$  plaques were present at 4 months of age and increased with age [14]. We used C57 mice as a control. Mice were divided into three groups (n = 15/group): C57 mice, APP/PS1 mice, and APP/PS1 mice that received MT in 4% PBS (100 mg/ l) from 4 to 8 months of age, when they were killed. Mice drink an average of 5 ml water/day; thus, their daily MT intake was estimated at 0.5 mg. Controls received PBS in water. Finally, there was 4% PBS in the drinking water.

#### Morris water maze

Behavior of mice was detected by the Morris water maze (MWM) after a 4-month treatment with MT. The aim of the MWM test was to test the spatial learning and memory ability of mice [15]. The maze consisted of a circular pool filled with water. Within the pool, there were four quadrants. A platform was placed in the center of a quadrant and submerged 1 cm below water. A camera was placed on the top of the MWM to track the swimming path of mice. In the place navigation test, mice had to find the platform within 60 s after they were released into water and the time until they found the platform was recorded. If they could not find the platform within 60 s,

escape latency was recorded as 60 s and mice were guided to the platform. Mice underwent four trials per day. The place navigation test was performed for 5 consecutive days. At 24 h after the end of testing, the hidden platform was removed and the spatial probe test was performed. Mice were placed in the water for 60 s to find the platform. The number of times the platform was crossed and the time spent in the target quadrant were recorded. All of the results were analyzed using the software of SPSS (SPSS Inc., Chicago, Illinois, USA).

#### Western blot analysis

After the MWM test, mice were deeply euthanized by 10% chloral hydrate, followed by cervical dislocation. The cerebral cortex and hippocampus from one hemisphere were isolated from three mice of each group. Tissues were homogenized in ice-cold RIPA lysing buffer and then centrifuged at 12 000g for 15 min at 4°C. The protein level of samples was tested using the bicinchoninic acid method. Protein samples were electrophoresed by SDS-PAGE (140 V) with a 10% separation gel and a 6% stacking gel, and transferred by constant current (180 mA, 1.5 h) to a polyvinylidene difluoride membrane that was blocked with milk powder at room temperature for 1 h and incubated overnight at 4°C with the following primary antibodies: rabbit anti-APP (1:1000), rabbit anti-PS1 (1:1000), rabbit anti-TFAM (1:5000; Abcam, Cambridge, Massachusetts, USA), rabbit anti-NRF1 (1:3000; Abcam), rabbit anti-NRF2 (1:3000), rabbit anti-PGC-1a (1:1000; Abcam); or mouse  $\beta$ -actin (1:1000; Golden Bridge International, Beijing, China). After washing with TBST three times, membranes were incubated at room temperature for 1.5 h with secondary antibodies: goat anti-rabbit IgG (1:5000) and goat anti-mouse IgG (1:5000; Golden Bridge International). Immunoreactive bands were visualized using electrochemiluminescence. The density of bands was analyzed by Image J (National Institutes of Health, Bethesda, Maryland, USA).

#### Transmission electron microscopy

The mice were deeply euthanized by 10% chloral hydrate, followed by cervical dislocation after the MWM test. We used three mice of each group for TEM. According to the conventional TEM sample preparation method [16], the brains were rapidly removed after mice were killed and cut into 1-mm-thick blocks, placed in glutaraldehyde, and embedded using Epon 812 (Sigma-Aldrich, St. Louis, Missouri, USA). Tissues were collected on formvar-coated, single-slot grids, and viewed by transmission electron microscopy.

## Real-time PCR to detect mitochondrial DNA copy number

Mice were deeply euthanized by 10% chloral hydrate, followed by cervical dislocation after the MWM test. Brains were rapidly removed after mice were killed and cut into 1-mm-thick blocks for TEM. The rest of the brains were used for Real-time PCR (three mice per group). MtDNA copy number was measured by real-time PCR using the Step one plus real-time PCR System (Applied Biosystems, Carlsbad, California, USA) with the SYBR Green detection method. Total intracellular DNA in the cortex and hippocampus CA1 neuronal cells was extracted using the QIAamp DNA mini kit (Qiagen, Germantown, Maryland, USA) following the manufacturer's instruction. Each real-time PCR reaction (20 ul total volume) contained  $3 \mu l$  template DNA (50 lg),  $10 \mu l$ of 2× SYBR Green Real-time PCR Master (Applied Biosystems), 1 µl each forward and reverse primers, and 5 µl ultrapure water. The primers for a subunit of the human electron transport chain used for mtDNA amplification were as follows: forward, 5'-CAAAC-CTACGCCAAAATCCA-3' and reverse, 5'-GAAATGA-ATGAGCCTACAGA-3' [17]. The mtDNA primers (human complex II, fragment length 164 bp) were designed to minimize amplification of the mtDNA pseudogenes embedded in the nuclear DNA. Primers for human nuclear 18s, as an internal control, were as follows: forward, 5'-ACGGACCAGAGCGAAAGCA-3' and reverse, 5'-GACATCTAAGGGCATCACAGAC-3'. Relative amounts of mtDNA and nDNA copy numbers were compared. The  $\Delta\Delta C_t$  (mtDNA to 18 s) represents the mtDNA copy number in a cell [18].

#### **Thioflavin S**

After the MWM test, the mice were anesthetized with chloral hydrate, and then immediately cardiac perfused with a 0.9% saline solution, followed by 4% paraformaldehyde in 0.1 M PBS (pH 7.4). After the perfusion, the brains were excised and postfixed overnight at 4°C. The brain tissue was then incubated in 30% sucrose at 4°C until equilibration (six mice per group). Brain sections were removed and cut in a sagittal plane at 30 µm thickness on a freezing microtome and mounted on slides. Slides were stained with 1% thioflavin-S for 5 min, and rinsed with 70% ethanol and distilled water. Images of plaques were captured with a camera connected to a fluorescence microscopy (1×71S1F; Olympus Tokyo, Japan). Plaque burden was calculated as percentage of the thioflavin-S staining area to the total area. Image Pro Plus 6 Media Cybernetics (Silver Spring, Maryland, USA) was used to analyze images.

#### ELISA

Mice were deeply euthanized by 10% chloral hydrate, followed by cervical dislocation after the MWM test. Brain samples were isolated from the mice (three mice per group). Hemibrains were flash frozen and stored at  $-80^{\circ}$ C until homogenization. The secretion of A $\beta_{1-40}$ and A $\beta_{1-42}$  in brain tissue was assessed by ELISA. Brain tissues were lysed in guanidine-Tris buffer and centrifuged at 12 000 rpm/min for 30 min; 50 µl sample was added to each well of plates, then incubated with antibodies for A $\beta_{1-40}$  and A $\beta_{1-42}$ , and quantified by ELISA kits (Invitrogen, Carlsbad, California, USA) following the manufacturer's instruction. The wavelength of absorbance was set at 450 nm.

#### Statistical analysis

All data were expressed as mean  $\pm$  standard error. Student's *t*-test was used to compare two groups. In twovariable experiments, two-way repeated-measures analysis of variance was used to evaluate the significance of differences between group means. *P* value less than 0.05 was considered statistically significant. All experiments were repeated independently at least three times. Data were analyzed using SPSS 17.0 software (SPSS Inc.).

#### Results

# Effect of melatonin on protein expression of transcription factor A pathway, nuclear respiratory factor 1, nuclear respiratory factor 2, and proliferator-activated receptor $\gamma$ coactivator 1- $\alpha$

We verified the AD mouse model by measuring the protein level of APP and PS1. The expression of APP and PS1 was greater in APPswe/PS1dE9 transgenic (APP/PS1) than C57 mice (Fig. 1a). The protein expression of TFAM, NRF1, NRF2, and PGC-1 $\alpha$  in the cortex and the hippocampus was lower in APP/PS1 than C57 mice (P < 0.05); MT restores the expression of TFAM, NRF1, NRF2, and PGC-1 $\alpha$  in APP/PS1 mice (Fig. 1b and c). Thus, MT promotes mitochondrial biogenesis in AD in mice.

#### Effect of melatonin on the mitochondrial structure in Alzheimer's disease mice

The mitochondrial structure was severely damaged in cortex and hippocampus CA1 neuronal cells of APP/PS1 mice compared with C57 mice (Fig. 2). The structures of mitochondrial membrane and mitochondrial crista were obviously destroyed. Arrows showed altered structures: (a) mitochondria split rings formed; (b) mitochondria swelling with broken cristae; And (c) mitochondria split rings significantly increased. MT could reverse the damaged mitochondrial structure.

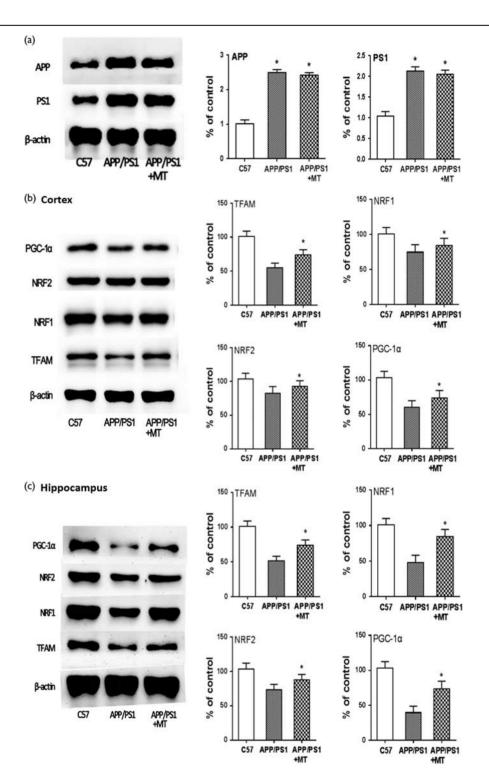
#### Effect of melatonin on mitochondrial DNA copy number

The number of mitochondria was significantly reduced in neurons of mice with AD (Fig. 3). The mtDNA/nDNA ratio in the cortex and hippocampus was lower in APP/ PS1 than C57 mice. However, with MT treatment for 4 months, the mtDNA/nDNA ratio was increased significantly in APP/PS1 mice.

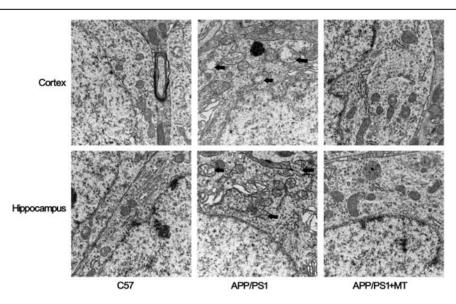
### Effect of melatonin treatment on Morris water maze behavior

The escape latency was longer for APP/PS1 than C57 mice, and MT treatment could ameliorate the latency in APP/PS1 mice (Fig. 4a). The effect of MT was similar for the number of platform crosses and the time spent in the target quadrant (Fig. 4b and c). The swimming route to

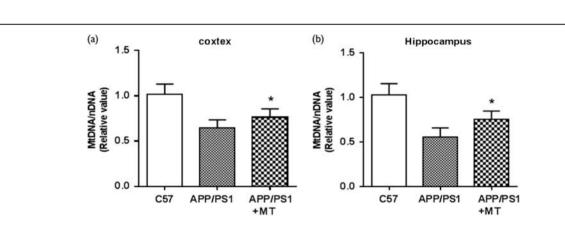
Fig. 1



Verification of AD mouse model and expression of mitochondrial biogenesis factors. Western blot analysis of protein levels of APP and PS1 (a) and TFAM, NRF1, NRF2, and PGC-1 $\alpha$  in the cortex (b) and hippocampus (c) of APP/PS1 and C57 mice. The scanned image of western blot was analyzed using the software Image J. Data are mean ± SE; \**P* < 0.05 versus APP/PS1 alone. AD, Alzheimer's disease; APP, amyloid precursor protein; MT, melatonin; NRF, nuclear respiratory factor 1; PGC-1 $\alpha$ , proliferator-activated receptor  $\gamma$  coactivator 1- $\alpha$ ; PS1, presenilin 1; TFAM, transcription factor A pathway.



Melatonin alleviated the damaged mitochondrial structure in the AD mouse model. Arrows show the altered structure of mitochondria. Transmission electron microscopy of the mitochondrial structure of cortex and hippocampus CA1 neuronal cells in APP/PS1 mice with and without melatonin and in C57 mice. AD, Alzheimer's disease; APP, amyloid precursor protein; MT, melatonin; PS1, presenilin 1.



Effect of melatonin on the mtDNA/nDNA ratio in cortex and hippocampus CA1 neuronal cells of mice with AD. Relative amounts of mtDNA and nDNA copy numbers were measured by real-time PCR. Data are mean±SE; \*P<0.05 versus APP/PS1 alone. AD, Alzheimer's disease; APP, amyloid precursor protein; mtDNA, mitochondrial DNA; MT, melatonin; nDNA, nuclear DNA; PS1, presenilin 1.

the previous goal location was longer for APP/PS1 than C57 mice (Fig. 4e) and was shorter with MT treatment. However, the swimming speed of the three groups did not differ (Fig. 4d). Hence, MT could alleviate the spatial learning and memory deficits in APP/PS1 mice and similar results have been shown previously by O'Neal-Moffitt *et al.* [11].

## Effect of melatonin on amyloid $\beta$ plaques and levels of $A\beta_{1-40}$ and $A\beta_{1-42}$

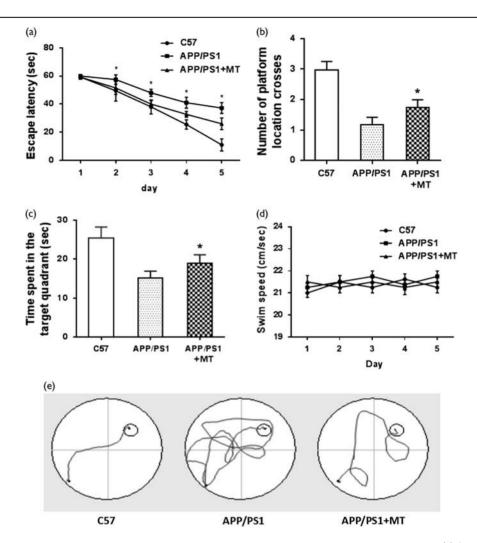
We used thioflavin-S staining to examine A $\beta$  plaques. The number of plaques in the cortex and hippocampus was significantly greater in APP/PS1 than C57 mice (Fig. 5a and b). MT reduced the number of plaques in APP/PS1 mice. We used ELISA to examine the levels of  $A\beta_{1-40}$  and  $A\beta_{1-42}$ . The levels of  $A\beta_{1-40}$  and  $A\beta_{1-42}$  were higher in APP/PS1 than C57 mice (P < 0.05); MT could reverse the levels in APP/PS1 mice (P < 0.05; Fig. 5c and d). Similar results have been reported previously by O'Neal-Moffitt *et al.* [11].

#### Discussion

In this study, we observed the effect of MT on the regulators of mitochondrial biogenesis in the model. We also studied the relation between mitochondrial biogenesis and the AD model. Compared with the controls, APP/PS1 mice with long-term oral MT intake showed increased levels of mitochondrial biogenesis factors, alleviated mitochondrial

Fig. 3





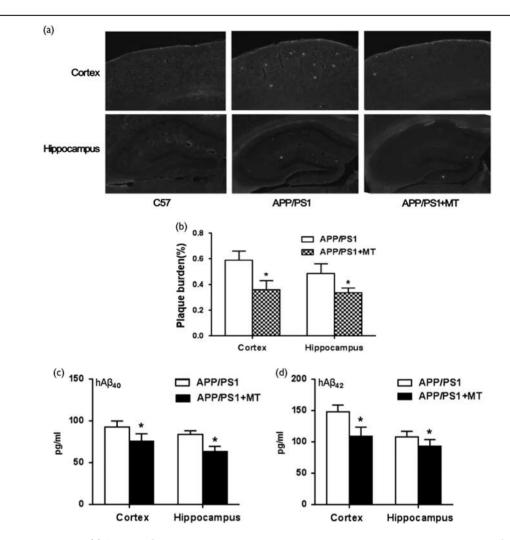
Melatonin improved spatial learning and alleviated memory impairments in mice with AD by the Morris water maze test. (a) Spatial learning measured as escape latency per day (n = 15 mice/group).(b, c). Spatial memory evaluated by the number of platform location crossings. (b)Time spent in the target quadrant (c). (d) Swimming speed. (e) Representative swimming route to previous goal location after 5 days of training. Data are mean ± SE; \*P < 0.05 versus APP/PS1 alone. AD, Alzheimer's disease; APP, amyloid precursor protein; MT, melatonin; PS1, presenilin 1.

impairment, enhanced mtDNA copy number, improved spatial learning and memory deficits, and reduced A $\beta$  plaque deposition and soluble A $\beta$  levels.

NRF1 and NRF2, which control the nuclear genes to encode mitochondrial protein, and TFAM, which drives transcription and replication of mtDNA, were significantly increased in APP/PS1 mice with long-term oral MT compared with controls as shown in Fig. 1. We believed that MT alleviated mitochondrial impairment, and enhanced mtDNA copy number through improving the expression levels of TFAM, NRF1, and NRF2. The expressions of NRF1, NRF2, and TFAM are regulated by PGC-1α [19]. Therefore, we speculate that MT can improve the structure and function of the mitochondria by the PGC-1α-NRF-TFAM pathway.

In Fig. 5, MT can reduce  $A\beta$  plaque deposition and soluble  $A\beta$  levels. In addition, MT can improve spatial

learning and memory of APP/PS1 mice as shown in Fig. 4. Similar results have been reported previously by O'Neal-Moffitt *et al.* [11]. Their findings showed that prophylactic MT significantly reduced AD neuropathology and associated cognitive deficits in a manner that was independent of antioxidant pathways. However, they did not reveal the exact mechanism of MT's action. According to Sheng et al. [7], they reported that damaged mitochondrial biogenesis was associated closely with mitochondrial dysfunction and played an important role in the pathogenesis of AD. In our study, MT can increase the expression levels of TFAM, NRF1, NRF2, and PGC-1 $\alpha$ . Therefore, we speculate that the PGC-1 $\alpha$ -NRF-TFAM pathway plays an important role in the pathogenesis of AD and the increase in spatial learning and memory and the decline in A $\beta$  plaque deposition and soluble  $A\beta$  levels may have a potential connection with



Melatonin reduced A $\beta$  deposition. (a) Thioflavin-S staining for senile plaques of the cerebral cortex and hippocampus in mice. Scale bar, 200 µm. (b) Quantification of thioflavin-S staining (*n* = 6 mice/group). The plaque burden was calculated as the percentage of thioflavin-S staining area over the total area (*n* = 6 mice/group). Image Pro Plus 6 (Media Cybernetics) was used to analyze the images. Melatonin reduced soluble A $\beta$  levels in AD mice. ELISA was used to detect A $\beta_{1-40}$  (c) and A $\beta_{1-42}$  (d) levels in the cortex and hippocampus of mice. Data are mean ± SE; \**P* < 0.05 versus APP/PS1 alone. MT, melatonin. A $\beta$ , amyloid  $\beta$ ; AD, Alzheimer's disease; APP, amyloid precursor protein; PS1, presenilin 1.

the PGC-1 $\alpha$ -NRF-TFAM pathway. We infer that MT improves mitochondrial dysfunction by promoting mitochondrial biogenesis, resulting in an increase in spatial learning and memory and a decline in A $\beta$  plaque deposition and soluble A $\beta$  levels.

Overall, this study showed reduced mitochondrial biogenesis signaling in the AD mouse model. Defective mitochondrial biogenesis may contribute to mitochondrial dysfunction in AD, and the use of MT, which can enhance mitochondrial biogenesis, may be a potential approach for the treatment of AD.

#### Acknowledgements

This work was supported by the National Natural Science Foundation of China (81571052, 81401052), the Special fund for basic research and business funds of Chinese Centre Universities (2018BTS01), the Key Research and Development Program of Shandong Province (2017GSF218036, 2017GSF218046), the Youth Talent Found of the 2nd Hospital of Shandong University (2018YT09), and the Fundamental Research Funds of Shandong University (2016JC022).

#### **Conflicts of interest**

There are no conflicts of interest.

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