

EXPERIMENTAL STUDY

Mechanism of protein phosphatase-2A regulating phosphorylation of amyloid precursor proteosome and A β generation

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Abstract: *Objective:* To discuss whether PP-2A influences the phosphorylation level at APP threonine 668 locus thus regulating A β secretion.

Method: In the experiment, 24 hours after N2a cells of stably transfected human APP (N2a/APP) were treated with okadaic acid (OA) or DES (C6-ceramide) (N2a/APP), an injection of OA cerebral stereotactic was administered to a SD rat in the hippocampal region, or PP-2A overexpressed plasmids was transfected transiently, the aggregate levels of APP phosphorylated APP, and APP-CTF were detected through immunoblotting and the activity of PP-2A and secretase was also detected using a reagent kit.

Result: The phosphorylation level of APP was significantly increased after the PP-2A activity was inhibited by OA. DES activated PP-2A or over-expressed PP-2A was able to reduce the phosphorylation level of APP. Either can inhibit PP and reduce the phosphorylation level of APP. The level of phosphorylated APP was increased significantly after the SD rat was injected with OA through the hippocampal region. The activity of β - and γ -secretases in N2a/APP cells significantly increased after OA treatment whereas the α -secretase activity had no significant changes; the A β level increased.

Conclusion: We discovered that PP-2A was capable of regulating the A β level by regulating APP phosphorylation level and β and γ -secretase activity (Fig. 5, Ref. 30). Text in PDF www.elis.sk.

Keywords: PP-2A, alzheimer disease, phosphorylation, A β , protein.

Introduction

Alzheimer's disease (AD) is a progressive neurodegenerative disease and its histopathological changes mainly include formation of a large number of extracellular senile plaques and entanglement of intracellular neurofibrils. The main component of the senile plaque is β -amyloid protein (A β) and malnutrition-related neuritis is generated in its vicinity (1). A β is generated in such a way that the amyloid precursor proteosome is severed continuously by two protein secretases, e.g. β -secretase (BACE-1) and γ -secretase (presenilin, PS-1/PS-2) (2). In amyloid pathologic changes, products resulted from β -secretase severing, e.g. N-terminal soluble segment (sAPP- β) and C-terminal segment can continue to be severed by γ -secretase and produce A β polypeptide finally (3). The toxic actions of A β are extensive and capable of causing cell death. The mechanisms mainly include: peroxidation damage, influencing calcium balance, activating aspartase proteinase, stimulating protein phosphorylation, mitochondria abnormality etc. (4). In addition, A β fibril can also specifically lead to neuron malnutrition. In the rat cortical neuron undergoing primary culture, over-expressed APP may lead to cell apoptosis, whereas γ -secretase inhibitor is

capable of blocking such apoptosis (5). In the transgenic rat model, A β aggregation can lead to malnutrition-related neuroinflammation, pathologic changes in tau protein, neuronal death, and DNA damage (6). When APP is over expressed or severed unusually, A β will form a toxic oligomer, deposit and form senile plaque, and cause damage to the learning memory (7). It has been reported that gene mutation of APP and PS-1 is the major cause of increased generated A β in the genetic AD disease. However, the cause of a significant rise of A β in the majority of sporadic AD patients still remains unclear (8).

In the research, we have discovered that overexpression or medication activated PP-2A in N2a/APP (stably transfected human APP695) cells could lower the phosphorylation level of 668 locus. The phosphorylation level of 668 locus could be lowered and generated A β increased after the PP-2A activity was inhibited. Meanwhile, we have also discovered that PP-2A was able to influence the activity of APP-related secretases and PP-2A could regulate secretion of A β by influencing phosphorylation and secretase activity of the threonine 668 locus.

Material and methods

Material

DMEM, OPTI-MEM, foetal bovine serum (FBS), and neomycin (G418) were purchased from Beijing Ruixiang Biotechnology Co., Ltd. The BCA protein determination reagent kit was purchased from U.S. SantaCruz Company, and the plasmid midi

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preparation kit was purchased from U.S. Amresco Company. The tetramethylbenzidine color developing agent was purchased from Beijing Biocoen Biotechnology CO., Ltd. The DNA molecular weight marker was purchased from Beijing Bole Bioscience Development Co. Ltd. The reverse transcription reagent kit and Taq enzyme were purchased from Beijing CoWin Bioscience Co., Ltd. The OA was purchased from U.S. Amresco Company. The Lipofectamine TM2000 transfection reagent was purchased from U.S. Amresco Company.

The cryogenic refrigerator was purchased from LEAD-TECH (Shanghai) Scientific Instrument Co., Ltd. The incubator-co2 was purchased from Zhengzhou Nanbei Instrument Equipment Co., Ltd. The inverted microscope was purchased from U.S. MEDICA Company. The gel electrophoresis analytic system was purchased from U.S. Labnet Company. The multi-function microplate reader was purchased from Beijing Jingli Centrifuge Co. Ltd.

Treatment of animal samples

The SD rats were purchased from Shanghai Laboratory Animal Centre of The Chinese Academy of Sciences, Conformity Certificate No. SCXK (Shanghai) 2013-0042 and all animals were in accordance with *Law of the People's Republic of China on Inspection and Quarantine of Animals and Plants*.

All animals were given sufficient water and food. The rats were sacrificed after intraperitoneal injection anaesthesia. The rats were decollated immediately after death and placed on the ice. The skull was scissored from the occipital bone hole of the centre of the head with scissors to remove the bones. The brain was raised along the edge of the cranial cavity with tweezers and placed on a piece of filter paper for blunt dissection of the hippocampus. The hippocampal part of the rat brain was taken out quickly, numbered as per EP tube, weighted respectively, and stored in a refrigerator at -80°C for homogenate in the future (9).

It was boiled for 10 minutes after homogeneous mixing of the tissue block, centrifuged at a room temperature at 8000g for 10 minutes. The insoluble components were taken out with a pinhead and the supernatant was stored in a refrigerator at -80°C for use in subsequent experiments.

Stereotactic injection to rat brain

20 SD rats weighing 200–220g were divided into the two groups with 10 for each group. The SD rats were anesthetized by 0.3% chloral hydrate, fixed onto the brain stereotactic instrument, cut in the centre of the top of the head to expose the bregma after the fur was removed and skin was sterilized. A microsyringe was inserted vertically by 2.5mm from the brain surface at 3.0mm behind the bregma and 2.5mm right of the centre line in accordance with the rat brain orthostatic map. The bilateral hippocampal regions were gently injected with 5 μl for 3min. The syringe needle was retracted gently after retaining for 5 minutes. All operation procedures were performed under an aseptic condition. Penicillin was applied to the skin incision site for antibiosis and the wound was sutured. The rats of the sham-operated group were injected with normal saline of an equivalent volume.

Cell culture and transfection administration

Mouse neuroblastoma cells (N2a) were purchased from Shanghai Boyan Biotechnology Co., Ltd. N2a cells of stably transfected human APP (N2a/APP). The cells were cultured in the incubator containing 5% CO_2 at 37°C and the culture medium was replaced at an interval of 3 days. The cells could be used for passaging or further experiment when the density reached approximately 90%.

LipofectAMINETM 2000 reagent was used for plasmid transfection in strict accordance with the instructions. The cells were transferred to a 96-well micro-plate and transfection was performed until the cell density reached 80–90%. The culture medium containing no blood serum neither antibiotic was used, replaced after 6 hours, and corresponding cell treatment and detection within 48 hours.

Extraction and concentration determination of protein samples

The tissue sample was stored in a cryogenic refrigerator following homogenization. Partial homogenate was taken and 2 \times SDS sample buffer of an equivalent volume was added, PMSF, protease inhibitor, subject to lysis on ice for 30 minutes. The lysis solution may freeze and mixing was performed at an appropriate interval. Then it was water bathed for 10 minutes and subjected to 20000Hz ultrasound 3s X15 times at the amplitude of approximately 20–30%. Excessive violent operation should be avoided to prevent the histone bound DNA from being broken. It was centrifuged at 8000g for 3 minutes. The supernatant was collected. The protein concentration was tested with the BCA method (10).

Western blotting

The separation gel was prepared in accordance with the molecular weight of the antigen. 3ml separation gel was infused into the gel trough and sealed after being full of double distilled water, 2ml spacer gel prepared was added to the gel trough after solidification. A broach was inserted and the electrophoresis channel was marked. The electrophoresis channel was washed with double distilled water and filled with electrophoretic buffer solution. Specimen of the identical quality was taken for polyacrylamide gel electrophoresis and 20 μg , 30 μg and 40 μg specimen were added on the basis of contents of different antigens in the cells. 5% skim milk powder was diluted with TBS and shaken slowly in a shaking table for 1 hour or stored in a refrigerator at 4°C overnight after addition of properly diluted primary antibody. It was washed with 1 \times TBS buffer solution, 10 min for each time. The Odyssey two colour fluorescence second antibody was incubated for 1h after diluted as per 1:1000. It was washed with 1 \times TBS buffer solution for 3 times, 10 minutes for each time; slightly washed with double distilled water and developed by Odyssey scanning.

ELISA adsorption method

G2-10 stock solution was diluted with 100 mM NaHCO_3 , (pH 9.6), 1 μg /well, G2-11 stock solution was diluted with 100 mM Tris-Cl, (pH 8.4), 1 μg /well. The coated antibody was added to a high affinity 96-well ELISA plate with 100 μl /well. The ELISA plate was placed in a wet box at 4°C overnight. It could be incubated for two nights if the final developing signal was excessively low

during Aβ42 detection. The coated antibody was thrown away. The wells were filled with TBS and rinsed on the shaking table for 3 times and 3 minutes. Each well of the ELISA plate was filled with TBS containing 5% BSA, blocked at room temperature for 2 hours. The ELISA plate was rinsed with TBS (TTBS) containing 0.1% Tween-20 for 3 times and 3 minutes. 10% BSA was diluted as per 1:100 and biotin-labelled 4G8, or diluted as per 1:500 and biotin-labelled WO2, 20μl/well. Then, 10μl (Aβ40) or 40μl (Aβ42) animal homogenate supernatant or cell culture media supernatant was added and normal saline or culture media was to supplement the insufficient part of the total volume of 100μl per well. It was stored at 4°C overnight. The sample and antibody were thrown away. The culture plate was rinsed with TTBS for 3 minutes. The avidin-labelled horse radish peroxidase HRP-NeuroAvidin (1mg/ml) was diluted as 1:1000 with 2% BSA. 100μlvidin-labelled horse radish peroxidase was added to each well and placed on the shaking table for incubation at room temperature for 1 hour. The culture plate was rinsed for 5 times with TTBS and once with TBS for 3 minutes after the HRP was thrown away. 100μl TMB:H₂O₂(1:1) solution was added to each well to observe changes in colour. 50μl of 1M H₂SO₄ was added to each well immediately after the solution became blue. At the time, the solution colour would become yellow. The absorbance value was read at a wave length of 450nm on the microplate reader and the content was calculated as per the standard curve.

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the centre of the top of the head to expose the bregma after the fur was removed and skin was sterilized. A microsyringe was inserted vertically by 2.5mm from the brain surface at 3.0mm behind the bregma and 2.5mm right of the center line in accordance with the rat brain orthostatic map. The bilateral hippocampal regions were gently injected with 5μl for 3min. The syringe needle was retracted gently after retaining for 5 minutes. All operation procedures were performed under an aseptic condition. Penicillin was applied to the skin incision site for antibiosis and the wound was sutured. The rats of the sham-operated group were injected with normal saline of an equivalent volume.

Data statistics

The gray values of the immunoblotting stripes were calculated with the Odyssey software for semiquantitative analysis. The immunofluorescence gray values were calculated with the Image-Pro-Plus 6.0. The data was expressed with the mean number ± standard deviation (mean±SD). The one-way analysis of variance or T test was used for comparison among groups. The statistical software, EXCEL, was used to analyse and process the data.

Results

Inhibition of PP1 in reducing phosphorylation level of APP

As a protein phosphatase inhibitor, OA can inhibit activity of PP-2A and PP1 respectively within different dosage ranges. The inhibition ranges are: 0.1–10 nM specificity inhibition PP-2A. The OA above 10nM is able to inhibit the activity of both PP-2A and PP1. As a threonine/serine esterase, PP1 may also influence the

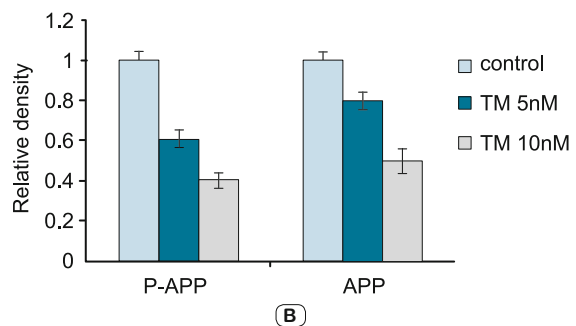
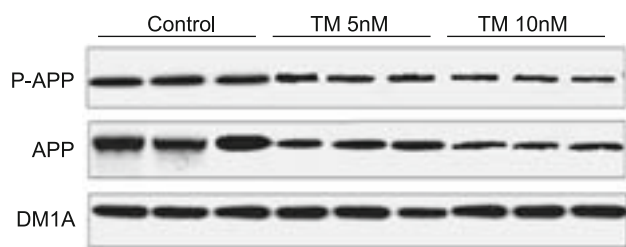


Fig. 1. Inhibit PP1 by Decreasing PAPP Activity in N2a/APP Cells via TM.

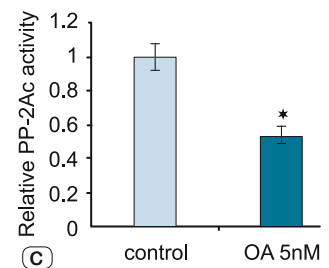
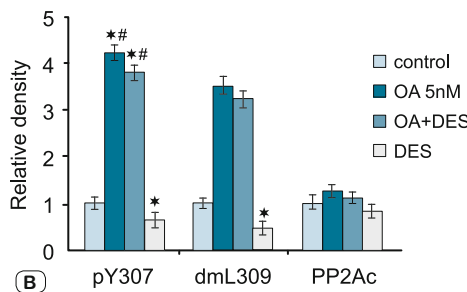
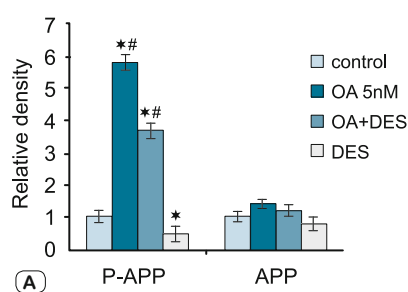


Fig. 2. Effect of Low-concentration OA and DES Specifically Regulating PP-2A Activity on APP Phosphorylation Level. Compared with the control group, * p <0.05; compared with the DES group, # p <0.05.

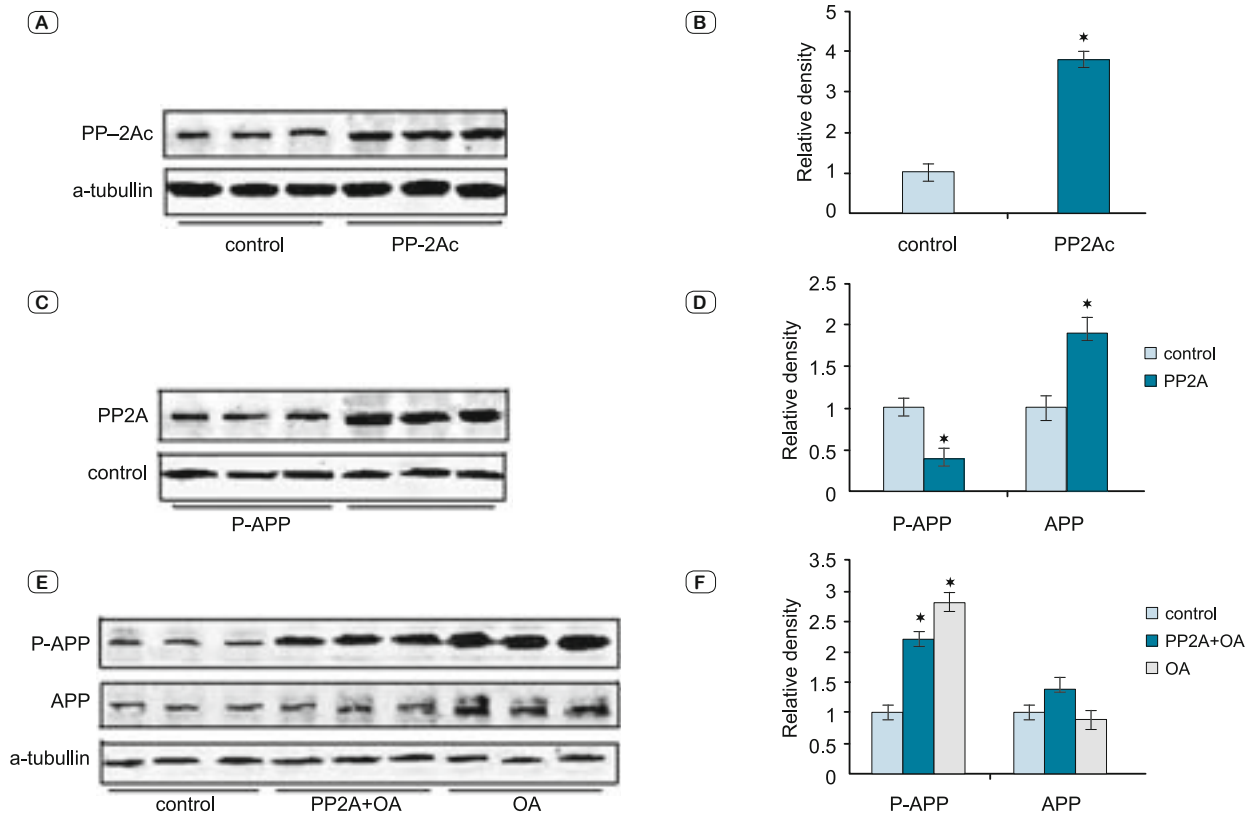


Fig. 3. Over-expressed PP-2A Reducing Phosphorylation Level of APP. The comparison with the control group indicates that * $p < 0.05$.

phosphorylation level of the APP668 locus. We have also detected the effect of solely regulating PP1 activity on phosphorylated APP. In the experiment, TM (tautomycin) with concentrations of 5nM and 25nM was administered to N2a/APP cells respectively. TM could serve as a specific protein esterase inhibitor and specifically inhibit PP1 activity within the above concentration range. Compared to the control group, the APP phosphorylation level in cells with concentrations of 5nM and 25nM exhibited a negative correlation with the dosage of administration indicating that when the PP1 activity was inhibited, the phosphorylation level of the APP668 locus decreased rather than increased (Fig. 1A and 1B). The result excluded the effect of PP1 in rising APP phosphorylation level induced by OA.

Low-concentration OA and DES specifically regulating PP-2A activity cause corresponding changes in APP phosphorylation

The OA effect dose was 0.1–10 nM and only had inhibitional effect on PP-2A. Therefore, we used 5nM as the administration concentration to study the role of PP-2A in APP phosphorylation level in the subsequent study. It was observed that the phosphorylation level of APP increased significantly compared to the control group during treating N2a/APP cells with OA for 24 h. PP-2A specificity agonist DES was administered and the phosphorylation level of APP decreased. DES was capable of reversing the rise in APP phosphorylation caused by OA after combined administration

(Fig. 2A). The result has shown that the phosphorylation level of the PP-2AY307 locus of the OA group increased and the demethylation level of L309 decreased after detection of active sites Y307 and L309 of PP-2A. This is exactly converse in the DES group indicating that OA inhibited PP-2A activity while DES activated PP-2A (Fig. 2B). Meanwhile, we have detected the activity of PP-2A and found that the activity of PP-2A in cells stimulated by 5nM OA decreased significantly compared to that of the control group (Fig. 2C). The above result has sufficiently demonstrated that PP-2A was able to regulate the phosphorylation level of APP.

Over-expressed PP-2A can reverse the rise in APP phosphorylation level caused by OA

In the previous experiment, we have demonstrated that low-concentration OA was capable of inhibiting the activity of PP-2A and increasing the phosphorylation level of APP whereas the rise in APP phosphorylation level induced by OA could be reversed after PP-2A was activated by DES. To further verify whether PP-2A participated in regulation of the phosphorylation level of APP, we have first over-expressed the PP-2A plasmid in N2a/APP cells (Figs 3A and 3B) and meanwhile tested the expression level of phosphorylated APP. The result has shown that compared to the control group, over-expressed PP-2A significantly decreased the protein level of APP while the total level of APP had no significant changes (Figs 3C and 3D). In addition, we have discovered

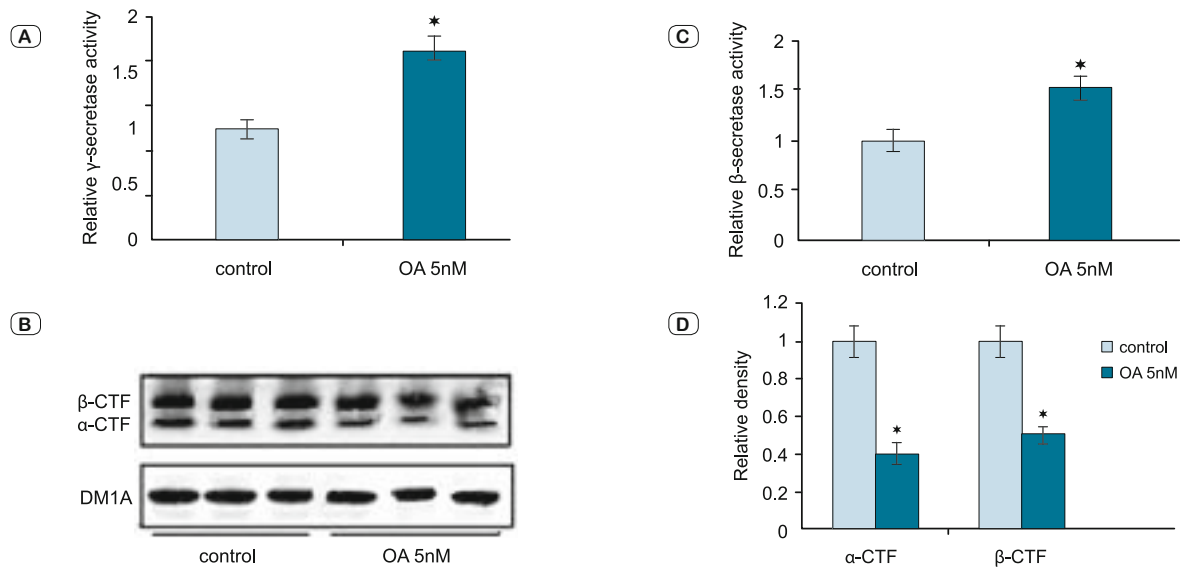


Fig. 4. low-dose OA activating the activities of β -secretase and γ -secretase and reducing the contents of α -CTF and β -CTF. The comparison with the control group indicates that * $p < 0.05$.

that the over-expressed PP-2A was able to reverse the rise in phosphorylation level of APP induced by OA (Figs 3E and 3F).

Inhibition of PP2A can up-regulate the activity of the β and γ secretases in N2a/APP cells and reduce output of α and β -CTF

The protein was extracted with the enzyme activity kit for corresponding enzyme activity detection 24 hours after 5 nM OA stimulation was administered to N2a/APP cells. The result has shown that compared to the control group, the activity of β -secretase in cells treated with OA increased, the activity of γ -secretase also increased (Figs 4A and 4B) whereas, the activity of α -secretase had no significant changes. The result indicated that the level of APP in cells severed by β -secretase and γ -secretase increased accordingly when the phosphorylation level of APP increased. $A\beta$ and β -CTF form under the action of β -secretase and γ -secretase. We have also detected the protein levels of α -CTF and β -CTF and the result has shown that the content of α -CTF and β -CTF decreased compared to that of the control group (Figs 4C and 4D).

Inhibition of PP2A causes rise in $A\beta$ generated

5 nM low-concentration OA stimulation was administered to the N2a/APP cells. The sample was collected after 24 hours and subjected to western blotting test by gradient tricine gel electrophoresis. The result has shown that compared to the control group $A\beta$ of OA-treated cells significantly increased indicating that decrease of the PP-2A activity can increase the quantity of $A\beta$ generated by increasing the phosphorylation level of the APP668 site. Meanwhile, we have also observed the protein level of β -CTF. The protein level of β -CTF in cells treated by OA significantly decreased compared to that of the control group, which was consistent with the above result (Figure 5A and B).

Discussion

For Alzheimer’s disease, senile plaques caused by amyloid proteinosis in the patient’s brain are one of the major pathologic changes. The major constituent is $A\beta$ polypeptide formed from the

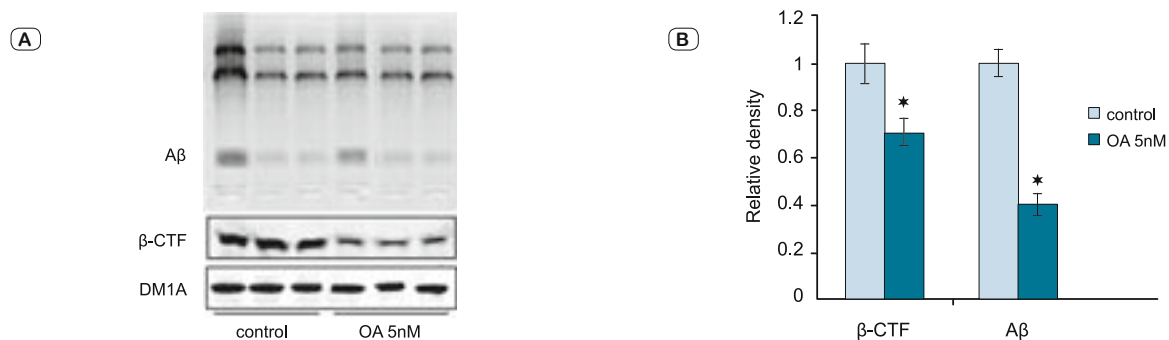


Fig. 5. Low-Dose OA Reducing the Quantity of $A\beta$ Generated. The comparison with the control group indicates that * $p < 0.05$.

amyloid precursor proteasome APP severed by secretase (11–12). APP is a transmembrane protein. Its intracellular domain contains multiple phosphorylation sites and two functional domains that can be bound with other proteins (13). Previous studies have shown that phosphorylation of APP threonine 668 site could influence the generation of intracellular A β and high phosphorylation of APP threonine 668 site could strengthen A β generation (14). PP-2A is a protein phosphatase widely present and conservative in cells, which can catalyze in vivo dephosphorylation of the phosphorylated site of the substrate (15). A number of reports have demonstrated that the activity of PP-2A in AD patients was decreasing (16).

APP is a membrane protein, which has a very long N-terminal extracellular fragment (NTF) and a very short C-terminal intracellular fragment (CTF). We have known very well that C-terminal intracellular fragment could be phosphorylated by multiple phosphokinases and the sites (17). It is considered that two elements in the APP intracellular domain could specifically regulate APP, could be severed by secretases, engulfed by cells or A β generation. (18). APP phosphorylation can regulate synapse extension in the differentiated PC12 cells (19). The NMR research have indicated that threonine 668 phosphorylation could significantly change APP structurally and might influence binding of APP and other proteins. It is thus considered that threonine 668 phosphorylation could regulate APP intracellular transfer and metabolism (20). Rise in phosphorylation of the threonine 668 site could strengthen the role of secretase in severing amyloid protein generated. Some reports have demonstrated that colocalization existed between APP (P-APP) and BACE1 of the phosphorylated threonine 668 within the hippocampal neurons and primarily cultured cortical neurons of AD patients and it has been clinically indicated that APP phosphorylation level increased significantly in AD patients' brains (21).

Protein phosphatase 2A is a ubiquitous and highly conserved serine/threonine protein esterase, which can be extensively bound with different substrates (22). In many reports, it was stated that imbalance of protein kinase and esterase effects was one of the major factors for AD onset (23). Research has found that the activity of PP-2A was increased after specifically down-regulating PP-2A specificity inhibitory proteins I2PP-2A in lentivirus infected hAPP transgenic mice and the A β level in the brains of the transgenic mice after down-regulating the APP phosphorylation level (24). These research results have indicated that PP-2A might regulate the phosphorylation level of APP threonine 668 site as an important threonine substrate (25).

PP-2A can reduce the phosphorylation level of APP and it was reported that the rise of the phosphorylation level of APP could increase the A β generated (26). Up to now, people still have not been well aware of the pathogenesis of AD but the majority of the scholars have stated that A β deposition was the common approach for all factors causing AD (27). An experiment has shown that A β has neurotoxicity, which can cause neuronal necrosis, apoptosis and amyloidosis (28). It has been generally accepted that the neurotoxicity of A β was the critical factor for AD formation and progression (29). A β accumulation is an early certain event during the AD onset. As for the onset sequence, A β occurs earlier than pathological changes and clinical symptoms such as nerve fibre

entanglement, neuraxon lacking nutrition etc. An experiment has demonstrated that A β was toxic to the cultured nerve cells and also the trophism and toxic effect of A β depended on the aggregation state transforming from solubility to insolubility, i.e. A β in the aggregation state has a toxic effect (30).

To discuss the effect of changes in the APP phosphorylation level on A β generation in our experimental system we detected the changes in quantity of A β generated by OA treatment in N2a/APP cells. Small molecular weight protein electrophoresis immunoblotting was used. We have discovered that the A β polypeptide content increased significantly in cells treated by OA. We have detected the activities of α -, β - and γ -cand severing products β -CTF and α -CTF due to the fact that the major approach for A β generation was secretase severing. The result has shown that the activities of β - and γ - secretases increased somewhat whereas the activity of α -secretase had no significant change compared to that of the control group. Contents of β -CTF and α -CTF decreased significantly. These results have indicated that the decreased PP-2A activity caused by OA could be increased by raising the APP phosphorylation level, inducing the rise of the activities of β - and γ - secretases, thus strengthening APP severing. We have discovered that the activities of γ -secretase and β -secretase increase, A β generated increases and β -CTF content decreases. The reason may be that the activity of γ -secretase increasing and acting upon the severed product β -CTF of β -secretase produced more A β polypeptide and decreased the contents of α -CTF and β -CTF.

Based on the above experimental results, in the research, we have discovered that PP-2A had a direct effect on the APP threonine 668 site in the N2a/APP (stably transfected human APP695) cells. Over-expressed or activating PP-2A by medication decreased the phosphorylation level of the APP668 site and A β generated. The phosphorylation level of APP668 site could be raised and the generated A β increased after the PP-2A activity was inhibited by medication. We have also obtained a consistent result in the animals. Meanwhile, we have also discovered that PP-2A was able to influence the activity of APP severing-related secretases and thus PP-2A could regulate secretion of A β by influencing activities of phosphorylated β - and γ -secretases at the threonine 668 site.

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