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# Salidroside attenuates hypoxia-induced abnormal processing of amyloid precursor protein by decreasing BACE1 expression in SH-SY5Y cells

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### ABSTRACT

Hypoxia which is mainly mediated by hypoxia-inducible factor 1 (HIF-1), can greatly contribute to the occurrence of Alzheimer's disease (AD) by increasing  $\beta$ -site APP cleaving enzyme (BACE1) gene expression, protein level and  $\beta$ -secretase activity, resulting in a significant generation of amyloid-beta (A $\beta$ ). Salidroside has been reported to have great neuroprotective effects. The aim of this study was to investigate the effects of salidroside on hypoxia-induced abnormal processing of the amyloid precursor protein (APP) in SH-SY5Y cells and its possible mechanism. Western blot analysis showed that 200  $\mu$ M of salidroside pretreatment significantly decreased BACE1 protein level and promoted the secretion of sAPP $\alpha$  in hypoxic condition. Salidroside had no effect on the level of APP, ADAM10 and ADAM17. ELISA analysis revealed that salidroside was able to inhibit the increase of  $\beta$ -secretase activity and A $\beta$  generation induced by hypoxia, with no effect on  $\gamma$ -secretase activity. Notably, under hypoxia condition, mRNA of *BACE1* and protein level of HIF-1 $\alpha$  were decreased by salidroside pretreatment. These results demonstrated for the first time that salidroside was able to attenuate abnormal processing of amyloid precursor protein induced by hypoxia in SH-SY5Y cells, providing a new insight into prevention and treatment of Alzheimer's disease.

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Alzheimer's disease (AD) is the most common type of dementia among older people. The aggregation and accumulation of extracellular A $\beta$  in the brain are important pathologic features of AD [7]. A $\beta$  is generated by sequential proteolysis of the amyloid precursor protein (APP) by  $\beta$ -secretase and then by  $\gamma$ -secretase [7,12]. BACE1 is a transmembrane aspartyl protease and is the principal  $\beta$ secretase responsible for A $\beta$  generation *in vivo* [1,4]. APP can also be cleaved by  $\alpha$ -secretase (ADAM10, ADAM17) within the A $\beta$  domain to generate soluble neurotrophic sAPP $\alpha$  [7,12].

Many studies have revealed that hypoxic/ischemic injury can greatly contribute to the occurrence of AD [13,15,17,26]. Hypoxia

significantly increases *BACE1* mRNA and protein levels [10,19,20]. Recent studies have shown that *BACE1* promoter contains a physiological functional hypoxia response element (HRE), and hypoxia can facilitate AD pathogenesis by increasing HIF-1 $\alpha$  binding to the HRE, thereby up-regulate *BACE1* gene expression, protein level and  $\beta$ -secretase activity, and eventually leading to a significantly generation of A $\beta$  both *in vivo* and *in vitro* [8,16,24,25].

Salidroside(2-(4-hydroxyphenyl)ethyl  $\beta$ -D-glucopyranoside) is extracted from the root of Rhodiola rosea L., which has been used in traditional Tibetan medicine for a long history. Recently, salidroside has been found to have a protective effect against amyloid-beta induced cytotoxicity in PC12 cells [9], and against H<sub>2</sub>O<sub>2</sub>-induced cell apoptosis in SH-SY5Y human neuroblastoma cells, primary rat hippocampal neurons and PC12 cells [2,3,22]. Our previous studies revealed that salidroside conferred neuroprotection against MPTPinduced nigrostriatal dopaminergic damage in mice (unpublished data). Another study reported the protective effects of salidroside on hypoxia/hypoglycemia-induced injury in cultured neurons [23]. Taken together, these studies suggest that salidroside exerts great neuroprotective effects and potential pharmaceutical applications for neurodegenerative disorders. However, whether salidroside has protective effects against hypoxia facilitated Alzheimer's disease remains unknown. Therefore, the aim of present study was to test

Abbreviations: A $\beta$ , amyloid-beta; AD, Alzheimer's disease; ADAM, a disintegrin and metalloproteinase; APP, amyloid precursor protein; BACE,  $\beta$ -site APP cleaving enzyme; sAPP $\alpha$ , soluble amyloid precursor protein  $\alpha$ ; HIF-1 $\alpha$ , hypoxia-inducible factor 1 $\alpha$ ; HRE, hypoxia response element; PS, presenilin; PEN-2, presenilin enhancer 2; APH-1, anterior pharynx-defective 1.

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the effect of salidroside on hypoxia-induced abnormal processing of APP and the potential mechanism.

Human neuroblastoma SH-SY5Y cells stably transfected with human wild type APP695 cDNA (SH-SY5Y<sup>APP695</sup>, kindly provided by Prof. Dai Zhang, Institute of Mental Health, Peiking University, Beijing) were cultured in DMEM/F12 (GIBCO, Carlsbad, CA) containing 1% non-essential amino acids, 1% glutamine, 300  $\mu$ g/ml G418 (Sigma–Aldrich Inc.) and 10% heat-inactivated FBS (Hyclone, Rockford, IL) at 37 °C with 5% CO<sub>2</sub>. Twelve to sixteen hours before drug treatment, the media was replaced with FBS-free DMEM/F12 media. Salidroside (Purity (HPLC) 99.74%. Shanghai Pharm Valley Corp., Shanghai, China) was dissolved in PBS and sterilized through a 0.2  $\mu$ m filter prior to use. For drug and hypoxia treatment, SH-SY5Y<sup>APP695</sup> cells were pretreated with indicated concentration of salidroside for 1 h then incubated in a 37 °C chamber containing 1% O<sub>2</sub> and 5% CO<sub>2</sub> as indicated time.

As for ELISA assay, the supernatants of SH-SY5Y<sup>APP695</sup> cells were collected and the concentrations of secreted Aβ40 and Aβ42 were determined using commercial ELISA kits (Shanghai ExCell Biology, China), following manufacturer's protocols. β-Secretase activity and γ-secretase activity of cultured SH-SY5Y<sup>APP695</sup> cells were detected using β-Secretase Activity Assay Kit (BioVision, CA, US) and γ-Secretase Activity Assay Kit (R&D Systems, Inc. Minneapolis, US), according to manufacturer's instructions.

LDH (lactate dehydrogenase) activity was determined using the Cytotoxicity Detection Kit (LDH) (Roche Applied Science, Mannheim, Germany), following manufacturer's protocols.

*Western blot analysis*: The culture media of SH-SY5Y<sup>APP695</sup> cells were collected, concentrated with Centricon YM-30 filter units (Millipore, Bedford, MA) and frozen at -80 °C until further analysis. Treated cells were lysed in ice-cold RIPA (50 mM Tris–HCl, pH 7.4, 150 mM sodium chloride, 1% Triton X-100, 1% sodium deoxycholate, 0.1% sodium dodecyl sulfate). Protein concentrations were determined using the Pierce BCA Protein Assay Kit (Thermo Scientific). Equal amounts of lysates were separated by SDS-PAGE, then transferred into PVDF membrane (Millipore). The membranes were blocked with 5% skim milk in 1× Tris-buffered saline containing 0.8‰ Tween-20 for 1 h. Then the membranes were incubated overnight at 4°C with the following antibody: HIF-1 $\alpha$ , Novus Biologicals, Littleton, US; BACE1, Alzheimer precursor protein A4 monoclonal antibody, 22C11, ADAM 10 and ADAM17, Millipore; Human  $\beta$  Amyloid (1-17), clone 6E10, Aph-1aL,  $\beta$ -actin and  $\alpha$ -tubulin, Sigma–Aldrich Inc. The membranes were then incubated with goat anti-mouse or goat anti-rabbit horseradish (HRP) secondary antibody (Jakson ImmunoResearch Laboratories) at room temperature for 1 h. The signals were developed with Immobilon Western Chemiluminescent HRP Substrate (Millipore, Billerica, MA).

Quantitative real-time polymerase chain reaction (RT-PCR): Total cellular RNA was extracted by TRIzol reagent (Invitrogen). RT was performed with a cDNA synthesis kit according to manufacturer's instructions (Takara, Dalian, China). For quantitative SYBR Green (2×SYBR Green Real-time PCR Master Mix, TOYOBO, Osaka, Japan) real-time PCR, 50 ng of cDNA was used per reaction. The following specific oligonucleotide primers were used for BACE1 (sense, 5'-CCGGCGGGAGTGGTATTATG-3', antisense, 5'-GCAAACGAAGGTTGGTGGT-3'), HIF-1 $\alpha$  (sense, 5'-ACTAGCCGA-GGAAGAACTATGAA-3', antisense, 5'-TACCCACACTGAGGTTGGTTA-3'), with  $\beta$ -actin as an internal control (sense, 5'-CATGTACGTT-GCTATCCAGGC-3', antisense, 5'-CTCCTTAATGTCACGCACGAT-3'). Real-time RT-PCR was performed with ABI 7900HT Sequence Detection System (Applied Biosystems, Foster City, CA) and data were analyzed using  $\Delta\Delta$ CT method.

All data in bar graphs were expressed as mean  $\pm$  SEM ( $n \ge 3$ ). Statistical analysis was performed by Student's *t*-test or ANOVA method using Prism software (GraphPad, San Diego, CA). A value of p < 0.05 was considered to be statistically significant.

We first evaluated whether hypoxia could increase the expression of BACE1 in human neuroblastoma SH-SY5Y<sup>APP695</sup> cells. For this, we treated SH-SY5Y<sup>APP695</sup> cells with hypoxic condition for 0, 3, 6, 12 or 24 h. As shown in Fig. 1A and B, hypoxia markedly increased the protein level of BACE1, with a significantly 2.6-fold increase at 12 h. Quantitative real-time PCR analysis showed 1.4-fold increased expression of *BACE1* mRNA (Fig. 1C). Concomitantly, the HIF-1 $\alpha$  protein level elevated obviously at 6 h (Fig. 1A). We also observed a significant increase of total BACE1 activity (Fig. 1D) and significant higher level of secreted Aβ40 (Fig. 1E) and Aβ42



**Fig. 1.** Hypoxia increased BACE1 expression,  $\beta$ -secretase activity and  $A\beta$  generation in SH-SY5Y<sup>APP695</sup> cells. SH-SY5Y<sup>APP695</sup> cells were treated with 1% O<sub>2</sub> for indicated time. (A) BACE1 and HIF-1 $\alpha$  were detected by Western blot with  $\beta$ -actin as an internal control. (B) Quantification of BACE1 protein levels. (C) *BACE1* mRNA level was detected by real-time quantitative RT-PCR. (D) Lysates of hypoxia-treated (1% O<sub>2</sub>, 12 h) and untreated SH-SY5Y<sup>APP695</sup> cells were assayed for  $\beta$ -secretase activity using  $\beta$ -Secretase Activity Assay Kit.  $\beta$ -secretase inhibitor was used as negative control. RFU, the relative fluorescence units per  $\mu$ g of protein sample. "p < 0.01, "p < 0.001 by ANOVA. (E and F) ELISA was performed to measure A $\beta$ 40(E) and A $\beta$ 42(F) in conditioned media from hypoxia-treated (1% O<sub>2</sub>, 12 h) and untreated SH-SY5Y<sup>APP695</sup> cells. "p < 0.01 by Student's *t*-test.



**Fig. 2.** Salidroside inhibited hypoxia-induced increase of BACE1 protein level,  $\beta$ -secretase activity and  $A\beta$  generation in SH-SY5Y<sup>APP695</sup> cells. After pretreatment for 1 h in the presence or absence of indicated concentration of salidroside, SH-SY5Y<sup>APP695</sup> cells were treated with normoxic or 1% O<sub>2</sub> for 12 h. (A) BACE1 and APP protein were detected by Western blot with  $\beta$ -actin as a loading control. (B) Densitometric analysis of BACE1 protein levels. (C) Lysates of SH-SY5Y<sup>APP695</sup> cells were applied to ELISA assay for  $\beta$ -secretase activity using  $\beta$ -Secretase Activity Assay Kit. RFU, the relative fluorescence units per  $\mu$ g of protein sample.  $\beta$ -Secretase inhibitor was used as negative control. (D and E) Conditioned cell culture media were collected, ELISA was performed to measure A $\beta$ 40(D) and A $\beta$ 42(E). (F) LDH was assayed using the Cytotoxicity Detection Kit.  ${}^{*}p < 0.05$ ,  ${}^{*}p < 0.01$  by ANOVA.

(Fig. 1F). These observations strongly indicated that hypoxia could increase BACE1 expression,  $\beta$ -secretase activity and A $\beta$  level in SH-SY5Y<sup>APP695</sup> cells.

To investigate the effect of salidroside on BACE1, SH-SY5Y<sup>APP695</sup> cells were preincubated with salidroside (0, 50, 100, 150 or 200  $\mu$ M) for 1 h, then incubated in hypoxic condition for 12 h. As shown in Fig. 2A and B, 100–200  $\mu$ M of salidroside pretreatment significantly decreased BACE1 protein level in a dose-dependent manner. In the presence of 150 or 200  $\mu$ M of salidroside, there were only 1.412- or 1.332-fold BACE1 protein as compared with only hypoxia-treated cells (2.635-fold).

We further tested the effect of salidroside on  $\beta$ -secretase activity in hypoxia condition by ELISA analysis. As shown in Fig. 2C, 200  $\mu$ M of salidroside pretreatment significantly attenuated hypoxia-induced up-regulation of  $\beta$ -secretase activity. Based on the results of BACE1 protein level and  $\beta$ -secretase activity, we expected that salidroside might inhibit the generation of A $\beta$ . Therefore, we examined extracellular A $\beta$  level in cultured medium from SH-SY5Y<sup>APP695</sup> cells by ELISA analysis. The results indicated that 200  $\mu$ M of salidroside preincubation significantly reduced hypoxia-induced excessively release of A $\beta$ 40 (Fig. 2D) and A $\beta$ 42 (Fig. 2E) to normal level. Further, we did not find obviously change in APP protein level either by salidroside or by hypoxia treatment (Fig. 2A), this implied that the reduced A $\beta$  generation was not the result of decreased APP, but may due to down-regulation of BACE1 by salidroside.

LDH analysis showed that 200  $\mu M$  of salidroside preincubation had no significant effect on LDH release into the culture medium under normoxia or hypoxia condition (Fig. 2F). This suggested that the decreased BACE1 protein level,  $\beta$ -secretase activity and A\beta generation after salidroside pretreatment were not induced by toxicity of salidroside to cells.

To investigate the effect of salidroside on  $\alpha$ -secretase and sAPP $\alpha$ . we performed Western blot analysis to detect sAPP $\alpha$  release in the supernatants of SH-SY5Y<sup>APP695</sup> cells using the 6E10 antibody, which recognized amino acid resides 1-17 of the human amyloid-beta. Compared with normoxia, hypoxia had no significant effect on the level of sAPP $\alpha$  (Fig. 3A and B). Notably, 200  $\mu$ M of salidroside pretreatment significantly increased sAPP $\alpha$  level (about 2-fold) either in hypoxia-treated or in untreated SH-SY5Y<sup>APP695</sup> cells (Fig. 3A and B). Consistent with some studies [18,25], hypoxic treatment did not significantly influence the levels of ADAM10 and ADAM17 (Fig. 3A). However, 200 µM of salidroside pretreatment failed to alter the expression of APP, ADAM10 and ADAM17 in hypoxic condition (Fig. 2A and Fig. 3A). It was thus implied that the promotion of sAPP $\alpha$  secretion was not the result of an increase in cellular APP, ADAM10 and ADAM17 expression, but was possibly due to decrease in  $\beta$ -secretase activity and other pharmacological properties.

As documented [10,18], hypoxia dramatically enhanced APH-1a protein expression. Here, we also observed a significantly increase (1.6-fold) in the APH-1 $\alpha$  protein level by 12 h of hypoxic treatment and 200  $\mu$ M of salidroside pretreatment reduced APH-1 $\alpha$  levels to 1.446-fold (Fig. 3C and D). Hypoxia treatment had no effect on the levels of PEN-2, PS1 and Nicastrin, and their protein levels remained unchanged by 200 µM of salidroside pretreatment (data not shown). We next investigated the effect of salidroside on  $\gamma$ -secretase activity. As can be seen in Fig. 3E, 12 h of hypoxic treatment with or without 200 µM of salidroside preincubation failed to significantly affect  $\gamma$ -secretase activity in SH-SY5Y<sup>APP695</sup> cells. This could be attributed to the unchanged other three components of  $\gamma$ -secretase (PEN-2, PS1 and Nicastrin) and their coordinate regulation. This indicated that only the increase of APH-1 $\alpha$  was insufficient to alter the activity of  $\gamma$ -secretase complex.





**Fig. 3.** Effects of salidroside on sAPPα release and γ-secretase activity. After pretreatment for 1 h in the presence or absence of 200 μM of salidroside, SH-SY5Y<sup>APP695</sup> cells were treated with normoxic or 1% O<sub>2</sub> for 12 h. (A) sAPPα level in the supernatants of SH-SY5Y<sup>APP695</sup> cells was detected. The cell lysates were analyzed and immunoblotted with antibodies against ADAM10, ADAM17, with β-actin as an internal control. (B) Quantification of sAPPα protein level. (C) APH-1α was detected by Western blot with β-actin as an internal control. (B) Lysates of SH-SY5Y<sup>APP695</sup> cells were applied to ELISA assay for γ-secretase activity using γ-Secretase Activity Assay Kit. RFU, the relative fluorescence units per μg of protein sample. "" p < 0.001, "p < 0.05 by ANOVA.

To understand the potential mechanism underlying the reduced expression of BACE1 protein after salidroside pretreatment, *BACE1* mRNA was measured in salidroside and/or hypoxia-treated SH-SY5Y<sup>APP695</sup> cells by real-time quantitative RT-PCR. Under hypoxic condition, endogenous *BACE1* mRNA levels showed significantly 1.4-fold increase compared with normoxia, and 200  $\mu$ M of salidroside pretreatment could reduce *BACE1* mRNA to 1.051-fold (Fig. 4A). These results suggested that salidroside could inhibit hypoxia-induced up-regulation of *BACE1* transcription. Since hypoxia-induced *BACE1* mRNA and protein up-expression was principally mediated by activation of HIF-1 $\alpha$  [8,16,25], we

continued to analyze whether HIF-1 $\alpha$  protein level was decreased by salidroside pretreatment under hypoxic condition. As shown in Fig. 4B and C, 200  $\mu$ M of salidroside pretreatment significantly reduced HIF-1 $\alpha$  protein level (0.55  $\pm$  0.06229) compared with only hypoxia-treated SH-SY5Y<sup>APP695</sup> cells, with no influence on *HIF*-1 $\alpha$  mRNA level (Fig. 4D). These results indicated that salidroside attenuated hypoxia-induced increase of BACE1 possibly through down-regulating HIF-1 $\alpha$  protein level.

Our study validated the findings that hypoxia could increase BACE1 expression,  $\beta$ -secretase activity and A $\beta$  generation [10,16,19,25] in human neuroblastoma SH-SY5Y cells. Here we



**Fig. 4.** Effects of salidroside on mRNA level of *BACE1* and protein expression of HIF-1 $\alpha$  in hypoxia-treated SH-SY5Y<sup>APP695</sup> cells. SH-SY5Y<sup>APP695</sup> cells were treated with or without 150 or 200  $\mu$ M of salidroside for 1 h then incubated in normoxic or 1% O<sub>2</sub> for 12 h. (A) *BACE1* mRNA level was detected by real-time quantitative RT-PCR. <sup>\*\*</sup> p < 0.01 by ANOVA. (B) HIF-1 $\alpha$  was detected by Western blot with  $\alpha$ -tubulin as an internal control. (C) Quantification of HIF-1 $\alpha$  protein levels. <sup>\*\*\*</sup> p < 0.001 by Student's *t*-test. (D) *HIF-1\alpha* mRNA level was detected by real-time quantitative RT-PCR.

found that salidroside pretreatment markedly attenuated hypoxiainduced increase in BACE1 protein levels and β-secretase activity. We also examined extracellular  $A\beta$  level by ELISA analysis and detected that 200 µM of salidroside preincubation significantly reduced hypoxia-induced excessively release of Aβ. LDH analysis confirmed that pretreatment with salidroside had no neurotoxicity to SH-SY5Y<sup>APP695</sup> cells. Thus, the reduced BACE1 protein level, βsecretase activity and AB generation by salidroside pretreatment were not induced by neurotoxicity of salidroside to SH-SY5YAPP695 cells. We also found that 200 µM of salidroside pretreatment significantly promoted sAPP $\alpha$  secretion. However, hypoxia and salidroside had no significant effects on the levels of ADAM10, ADAM17, APP and  $\gamma$ -secretase activities, this indicated that the elevated release of sAPP $\alpha$  was due to reduction in BACE1 protein level,  $\beta$ -secretase activity and other pharmacological properties of salidroside, not appear to be mediated by the promotion of APP, ADAM10 and ADAM17 expression.

Multiple studies have shown significantly elevated BACE1 protein level and activity in sporadic AD patients, suggesting that abnormal increased BACE1 may constitute a determining factor in the pathogenesis of sporadic AD [5,6,21]. Some studies demonstrated that BACE1 inhibition dramatically reduced cerebral Aβ levels, and BACE1 deficiency rescued memory deficits in AD mouse model without profound side effects [1,11,14]. These findings indicate that BACE1 is a prime therapeutic target for prevention and therapy of AD. Several studies have revealed that BACE1 gene promoter contains a physiological functional HIF-1 $\alpha$  binding element, and HIF-1 $\alpha$  can modulate BACE1 mRNA and protein level by binding to the BACE1 promoter [8,16,24,25]. Overexpression of HIF-1 $\alpha$  with HIF-1 $\alpha$  cDNA increased BACE1 mRNA and protein level, while down-regulation of HIF-1 $\alpha$  by siRNA reduced the level of BACE1 in both hypoxia-treated and untreated cells [8,16,25]. Furthermore, in hippocampus and cortex of HIF-1α conditional knock-out mice, BACE1 protein level was significantly reduced [25]. Thus, we tested whether the inhibition of BACE1 by salidroside under hypoxia condition was mediated by regulation of HIF-1 $\alpha$ . As we observed, salidroside significantly reduced HIF-1a protein and BACE1 mRNA level. This indicated that salidroside may inhibit binding of HIF-1 $\alpha$  to BACE1 promoter by reducing HIF-1 $\alpha$  protein level, thereby down-regulated BACE1 mRNA and protein level, modulating process of APP.

In conclusion, this work showed for the first time that salidroside could inhibit hypoxia-induced increase of BACE1 expression,  $\beta$ -secretase activity,  $A\beta$  generation, and promote the secretion of sAPP $\alpha$ . These findings suggest a new potential application value of salidroside in the treatment of AD.

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