Original Article
E3 ubiquitin ligase Siah-1 downregulates synaptophysin expression under high glucose and hypoxia

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Abstract: Background: Diabetes is proved to be one of the independent risk factors for cognitive dysfunction. The pathophysiologic changes caused by diabetes including hyperglycemia and tissue hypoxia may contribute greatly to cognitive decline. In the present study, we demonstrate E3 Ubiquitin Ligase Siah-1 downregulates the key synaptic protein Synaptophysin expression under high glucose and hypoxia condition which may be the underlying factor leading to cognitive dysfunction in diabetic patients. Methods: In this study, hypoxia (2% oxygen) and high glucose (50 mM) were used to treat primary neuronal culture. By using quantitative PCR and western blotting we determined the influence of hypoxia and high glucose on the expression of synaptophysin and Siah-1 and the phosphorylated forms of extracellular signal-regulated kinase (ERK). Knockdown of Siah-1, inhibitors for proteasome, lysosome and ERK kinase was employed to evaluate the role of Siah-1 and ERK activity on the expression of synaptophysin. By immunoprecipitation we also examined the role of Siah-1 in the ubiquitination of synaptophysin under hypoxic and hyperglycemic condition. Results: We demonstrated that hypoxia and high glucose together but not hypoxia or high glucose alone mediated posttranscriptional reduction of synaptophysin with increased ERK phosphorylation and Siah-1 expression. The downregulation of synaptophysin was reversed by inhibition of ERK and Siah-1 knockdown. Overexpression of Siah-1 accelerated the degradation of synaptophysin under hypoxia and high glucose conditions and promoted the ubiquitination of synaptophysin. Conclusions: The present results demonstrate that Siah-1 is the key factor that contributes to hypoxia and high glucose mediated synaptophysin degradation.

Keywords: Siah-1, diabetes, synaptophysin, ERK, cognitive dysfunction

Introduction
Vascular risk factors such as hypertension, stroke, diabetes, have been demonstrated in close relationship with cognitive decline in the elderly. Among them, diabetes has been affirmed to cause cognitive dysfunction in clinical research [1-5].

A multi-center European prospective study (LADIS) found that diabetes is one of the independent risk factors for cognitive dysfunction [5]. There is also evidence showing that blood glucose control could significantly improve cognitive function in elderly patients [6-8]. Animal experiments showed that diabetes can cause significantly reduced hippocampal mossy fiber endings and the number of synaptic vesicles, leading to cognitive decline in animals [9, 10]. Studies have indicated that people with diabetes may also have obviously reduced oxygen supply to the tissue [11-14]. Considering hypoxia as another important risk factor for cognitive decline, diabetes related cognitive dysfunction may be caused by chronic hyperglycemia and tissue hypoxia.

Synaptophysin (Syp) is a key synaptic vesicle protein which is involved in synaptic vesicle exocytosis and synaptic vesicle biogenesis. Syp was demonstrated to relate closely with cognitive function [15-17]. In neurons, Syp is mainly degraded through the ubiquitin-proteasome system. Studies have shown that E3 ubiquitin ligase Siah (Mammalian Homologues of Seven in Absentia) is the main protease that degrades Syp [18]. The expression of Siah was elevated significantly by both hypoxia and...
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hyperglycemia. We postulated that in diabetes, both hyperglycemia and hypoxia might cause great synaptic changes, resulting in cognitive dysfunction. Thus, in this study, we examined the influence of hypoxia together with higher glucose level on the expression of Syp in cultured neurons, and evaluated the role of Siah in this pathological change.

Methods

Primary hippocampus neuron culture

Sprague-Dawley rats were purchased from the Shanghai Institute of the Chinese Academy of Science. All experimental procedures were carried out in accordance with the experimental standards of Tongji University, as well as international guidelines on the ethical treatment of experimental animals. Neurons were isolated from embryonic day (E) 14.5 embryos [19]. Briefly, the embryos were removed and placed into ice-cold Dulbecco's minimal essential medium. Using a dissecting microscope, the hippocampus was dissected away from the brainstem, the meninges were removed and the telencephalon was placed into ice-cold HBSS. The tissue was digested in trypsin-EDTA (0.1%) for 20 min, washed three times in Dulbecco's minimal essential medium, and dispersed in neurobasal medium containing B27, L-glutamine and glutamic acid. The cells were plated on poly-L-lysine coated-glass coverslips, 96-well plates or 100 mm dishes at a density of 3-10×10^5 cells/ml and maintained at 37°C in a humidified 5% CO_2 incubator. Under these conditions, neuronal purity was approximately 90%, as estimated by positive immunocytochemical staining for neurofilament proteins and the negative immunocytochemical staining for glial fibrillary acidic protein (data not shown). The cultured neurons were used for studies on in vitro days 8-10 (DIV 8-10).

Cell treatment

Optimal survival rate and neurite growth of hippocampal neurons require 25 mM basal glucose [20], reflecting the fact that neurons have high metabolic rates. Neurobasal medium containing 25 mM glucose (normal cultured, NC) meets these metabolic requirements. After 7 days in culture, cells were incubated with 50 mM of glucose (high glucose, HG) or with 25 mM glucose + 25 mM mannitol (osmotic control, OC), Then maintained for further 3 days under hypoxia (2% oxygen) or Normoxia (21% oxygen) condition. The proteasome inhibitor MG132 or lactacystin (20 μM for both; Calbiochem, San Diego, CA), the lysosome inhibitor E64 (50 μM; Sigma, St.Louis, MO, USA), the extracellular signal-regulated kinase (ERK) kinase inhibitor U0126 (10 μM; Cell Signaling Technology, Beverly, MA) or PD98059 (10 μM; Calbiochem) or vehicle were added before the hypoxia procedure.

Cellular injury assays-LDH measurement

Cellular injury was determined by measuring concentration of lactate dehydrogenase (LDH) released into the medium [21]. Briefly, the media was removed and LDH in the medium and total cellular LDH were determined using the CytoTox 96® Non-Radioactive Cytotoxicity Assay (Promega, Madison, WI, USA). The maximal LDH release was obtained in each well following repeated cycles of freezing and thawing. Each experimental condition was repeated in triplicate. Results were expressed as a percentage of maximal LDH release, after the subtraction of background levels was determined from the medium alone.

Cell viability assay

The viability of cells was examined by 3-(4,5-dimethylthiazole-2-yl)-2,5-diphenyltetrazolium bromide (MTT) assay. After hypoxia and high glucose exposure, MTT was added to a final concentration of 0.5 mg/mL for 4 h. The supernatant was removed and 150 uL dimethylsulfoxide was added for 20 min. The MTT optical density values were measured on a microplate reader at 570 nm and 630 nm wavelengths. Each experimental condition was repeated in triplicate.

Quantitative PCR

The knockdown efficiency of siah-1 siRNA was confirmed by measuring siah-1 expression using quantitative real-time PCR. And the expression level of synaptophysin was also been detected. Briefly, total RNA was purified using RNA easy columns (Qiagen, Valencia, CA, USA) and cDNA synthesis was performed using SuperScript III (Invitrogen, Carlsbad, CA, USA). Primers were designed to amplify a segment to rat siah-1 and synaptophysin. Rat β-actin prim-
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ers were used in the same reactions to control for the amount of starting template. All samples were measured in duplicate and were compared to standard curves of known concentrations of different genes; all mRNA expression data is expressed relative to β-actin.

**Immuno blotting**

The cultured primary neurons cells were placed into RIPA buffer. The whole cell lysate (40 μg) were separated using SDS-PAGE and transferred to nitrocellulose membranes (Millipore, Bedford, MA, USA). After nonspecific binding was blocked with 4% skim milk, the membranes were incubated at 4°C overnight with a rabbit polyclonal antibody against phosphorylated forms of ERK or total ERK (1:1,000, 1:1,000, respectively; Cell Signaling Technology), a mouse monoclonal antibody against synaptophysin (1:500; Sigma), a goat polyclonal antibody against Siah-1 (1:2000; Abcam, Cambridge, MA, USA) or mouse monoclonal antibody against tubulin (1:2,000; Sigma). The membranes were then incubated with a horseradish peroxidase-conjugated goat antibody against rabbit or mouse immunoglobulins or with a biotinylated secondary antibody followed by avidin-biotin horseradish peroxidase complexes (Vectastain Elite ABC Kit; Vector, Burlingame, CA). The signals were visualized with chemiluminescence (ECL Blotting Analysis System; Amersham, Arlington Heights, IL), measured by ImageJ software (National Institutes of Health, Bethesda, MD) and normalized to -tubulin.

**siRNA mediated siah-1 knockdown**

For siah-1 knockdown, hippocampus neurons were transfected with siRNA against rat siah-1 (5’-GAAAUCCGACAACAACUUUU-3’; Dharmacon, Lafayette, CO, USA) or a scrambled control siRNA with no significant homology to any known gene sequences (ID#4611; Ambion, Carlsbad, CA, USA) using Lipofectamine™ 2000 (Invitrogen), according to the manufacturer’s instructions. After 5 h incubation, the medium was replaced with regular culture medium, and the cells were cultured for an additional 48 h. The efficiency of siah-1 knockdown was confirmed by real time RT-PCR and Immuno blot analysis.

**Transient transfection of constructs**

Cultured HEK 293 cells were transfected with Myc-synaptophysin, Flag-Siah-1 or HA-ubiquitin plasmid (1 ug per 1×10^5 cells) using Lipofectamine™ LTX (Invitrogen) according to company instructions. Six hours later, medium was replaced with DMEM containing 10% fetal

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**Figure 1.** Neuronal cell death was induced by both hypoxia and high glucose exposure. A. Primary cultured hippocampus neurons were treated by 50 mM glucose (HG) under hypoxia (2% oxygen) or Normoxia (21% oxygen) condition for 3 days. Cells treated by 25 mM glucose + 25 mM mannitol were used as osmotic control (OC). LDH release was measured. Data are mean ± SD from six independent experiments. B. Primary cultured hippocampus neurons treated either by 50mM glucose or 25 mM glucose + 25 mM mannitol were subjected to 3 days hypoxia or normoxia exposure and cell viability was assessed with MTT assay. Cells under normal cultured medium were used as normal control (NC). Data are mean ± SD from six independent experiments. *, P<0.05 vs. NC or OC group. NC, normal control; OC, osmotic control; HG, high glucose.
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bovine containing either 50 mM glucose or 25 mM glucose + 25 mM mannitol. Transfected cells were detected by immunoblotting with antibodies against Myc (1:1000, Millipore), Flag (1:1000, Sigma) or HA (1:1000, Roche).

**Ubiquitination assays**

For immunoprecipitation (IP), HEK293 cells were grown in 100 mm dishes and transiently transfected with 1 μg of each plasmid containing HA-ubiquitin, Myc-synaptophysin in the presence or absence of Flag-Siah-1. Cells were then collected and lysed as described [27] in IP buffer [1X PBS, 0.5% Triton-X-100, Complete Mini protease inhibitor (Roche)] and rotated for 1 h at 4°C. Supernatant was collected for IP after centrifugation at 17,500 g for 15 min at 4°C. Primary monoclonal anti-Myc antibody (2 μg, Millipore) or 2 μg mouse IgG (Santa Cruz) was bound to Dynabeads Protein G (Invitrogen) at room temperature for 30 min by rotation. The beads-Ab complexes were then washed in PBS containing 0.01% Tween 20 prior to being incubated with the soluble cell lysates (500 μg) overnight at 4°C with rotation. The beads-Ab-Ag complexes were then washed with citrate-phosphate buffer (pH 5.0). After washes of nonspecific binding to beads, the bound protein was solubilized in sample buffer, loaded on an SDS-PAGE, and processed for immunoblotting.

**Statistical analysis**

All values are expressed as mean ± S.E. Differences were analyzed using either one-way or two-way ANOVA followed by Newman-Keuls post hoc testing for pairwise comparisons using SigmaStat Version 3.5. The null hypothesis was rejected when the p value < 0.05.

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**Figure 2.** Posttranscriptional reduction of synaptophysin was detected in primary cultured hippocampus neurons after hypoxia and high glucose exposure. A. Primary cultured hippocampus neurons treated either by 50 mM glucose or 25 mM glucose + 25 mM mannitol were subjected to 3 days hypoxia or normoxia exposure. The levels of synaptophysin were analyzed by immunoblot. Shown are representative blots from six independent experiments with similar results. B. Neurons were treated as described above; the mRNA expression levels of synaptophysin are shown. There was no significant difference between different groups. Data are mean ± SD from six independent experiments. C. Densitometric quantification of synaptophysin in hippocampus neuronal cells treated with different culture medium in response to hypoxia or normoxia exposure. Data are mean ± SD from six independent experiments. *, P<0.05 vs. NC or OC group. NC, normal control; OC, osmotic control; HG, high glucose.
Results

The cell viability was decreased when the primary cultured neurons were exposed to both hypoxia and high glucose

To evaluate the effect of both hypoxia and high glucose on the neuronal cell viability, we cultured primary hippocampus neurons and treated the cells with either 50 mM glucose or 25 mM glucose + 25 mM mannitol as osmotic control for 3 days under hypoxia (2% oxygen) or Normoxia (21% oxygen) condition. Our data demonstrated that neither hypoxia nor high glucose exposure alone could elicit decrease in cell viability and increase in LDH release (Figure 1). However, when the cultured neurons were exposed to hypoxia and high glucose together, the cell viability were dramatically decreased. This evidence demonstrates the neuronal damage caused by diabetes may relate to both tissue chronic hypoxia and high glucose environment in pathological conditions.

Hypoxia and high glucose mediated posttranscriptional reduction of synaptophysin was regulated by the ubiquitin-proteasome system in neuronal cells

To elucidate the molecular mechanisms involved in the diabetes related cognitive decline. We used the in vitro culture system with primary hippocampus neurons stimulated with high glucose or isomotic medium and exposure to hypoxia for 3 days. As we showed in Figure 2A and 2B, the exposure of hippocampus neurons to elevated glucose or hypoxia did not affect the mRNA or protein content of synaptophysin, but in vitro application with high glucose and hypoxia together significantly reduced synaptophysin protein, but not mRNA. Because synaptic vesicle proteins including synaptophysin have recently proven to be physiologically degraded by the ubiquitin-proteasome system (UPS) for the maintenance of synaptic plasticity [18, 22, 23], we examined the involvement of UPS with hypoxia and high glucose posttranscriptional decrease in synaptophysin protein. As we showed in Figure 3, application with the proteasome inhibitor MG132 or lactacystin, but not the lysosome inhibitor E64, led to significant suppression of hypoxia and...
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high glucose induced degradation of synaptophysin. In contrast, the baseline levels of synaptophysin in neurons were unaffected with these inhibitors.

Hypoxia and high glucose mediated ERK activation was required for synaptophysin degradation in neuronal cells

Synaptophysin protein may be degraded by the mammalian homolog of Drosophila seven in absentia (sina), an E3-ligase selective for synaptophysin named seven in absentia homologue (Siah). Since Drosophila sina is regulated by ERK signaling [24, 25], the Siah may also be regulated by ERK activation. We further examined the involvement of ERK activation with hypoxia and high glucose mediated degradation of synaptophysin in the neuronal cells. Consistent with synaptophysin degradation, ERK phosphorylation was significantly enhanced after hypoxia and high glucose exposure. Hypoxia or high glucose exposure alone could induce the phosphorylation of ERK, but not as dramatic as in Hypoxia + HG group (Figure 4A and 4B). Inhibition of ERK activation with U0126 or PD98059 led to a significant suppression of hypoxia and high glucose induced degradation of synaptophysin (Figure 5A and 5D). In addition to ERK signaling, Siah-1 expression levels were also been detected. In the neuronal cells, consistent with phosphorylation of ERK and synaptophysin degradation, Siah-1 was also dramatically upregulated and this elevation was totally reversed by ERK phosphorylation inhibitor (Figures 4A, 4C, 5A and 5C). These results suggested that ERK activation is responsible for hypoxia and high glucose induced UPS degradation of synaptophysin through the regulation of E3-ligase Siah-1.

Posttranscriptional reduction of synaptophysin in hypoxia and high glucose exposed neuronal cells were reversed by Siah-1 gene knockdown

To further determine whether Siah-1 protein regulates the degradation of synaptophysin, we examined the effect of knockdown Siah-1 on synaptophysin protein expression levels in hypoxia and high glucose exposed primary cultured hippocampus neurons. Here, we generated a hypoxia and high glucose model, and manipulated Siah-1 by knocking down endogenous Siah-1 expression by siRNA. Siah-siRNA significantly reduced siah-1 mRNA and protein levels over 80% in the primary neurons (Figure 6A-C). Silencing of Siah-1 significantly reversed hypoxia and high glucose induced reduction of synaptophysin (Figure 7A-C). Taken together, these results indicate that Siah-1 proteins have the ability to regulate synaptophysin expression under hypoxia and high glucose conditions.
Siah-1 downregulates synaptophysin expression via the ubiquitin-proteasome pathway during hypoxia and high glucose conditions

Next, we sought to determine whether the enhanced degradation of synaptophysin by Siah-1 is mediated by the proteasome pathway. HEK293 cells exposed to hypoxia and high glucose that have been co-transfected with myc-synaptophysin and Flag-Siah-1 were treated by various inhibitors of proteolytic pathways, and the synaptophysin levels were then analyzed by Western blotting. As shown in Figure 8, the enhanced degradation of synaptophysin by Siah-1 overexpression is blocked by MG132, a potent inhibitor of proteasome function [26]. A similar effect was also observed when cells were treated with lactacystin, an irreversible inhibitor of the proteasome pathway (data not shown). In contrast, E64, an inhibitor of lysosomal cysteine proteases, had no effect.

Because proteasome-dependent proteolysis involves the ubiquitination of target proteins, we investigated whether Siah-1 accelerates the degradation of synaptophysin under hypoxia and high glucose conditions by promoting the ubiquitination of synaptophysin. After hypoxia and high glucose exposure, HEK293 cells were co-expressed with Myc-tagged synaptophysin along with HA tagged ubiquitin in the absence or presence of exogenous Siah-1. Cell lysates were subjected to immunoprecipitation with an anti-myc antibody followed by immunoblotting.

Figure 5. Hypoxia and high glucose mediated ERK activation and Siah-1 production was required for synaptophysin degradation in neuronal cells. A. Inhibition of ERK activation with U0126 or PD98059 led to a significant suppression of Hypoxia and high glucose induced upregulation of Siah-1. Consistent with this result, inhibition of ERK activation also blocked the decline of synaptophysin under hypoxia and high glucose condition. B-D. Densitometric quantification of ERK Siah-1 and Synaptophysin in hippocampus neuronal cells exposed to high glucose under hypoxia condition. Data are mean ± SD from six independent experiments. *, P<0.05 vs. control group.
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with an anti-HA antibody to detect ubiquitin-conjugated synaptophysin (Figure 9). Compared to normoxia + OC group of cells, hypoxia and high glucose exposure dramatically increased the ubiquitination of synaptophysin. Under hypoxia and high glucose conditions, in the absence of exogenous Siah proteins, synaptophysin was ubiquitinated, and the ubiquitinated synaptophysin could be detected by the anti-HA antibody immunoblotting. Overexpression of Siah-1 enhanced the ubiquitination of synapto-

Figure 6. siRNA-mediated knockdown of Siah-1 in primary cultured hippocampus neurons. Primary culture hippocampus neurons were transfected with different concentrations of Siah-1-siRNA indicated for 48 hours. Cells without transfection or transfected with scramble siRNA (Scra-siRNA) were used as controls. A. Quantitative real-time RT-PCR was performed to quantify the extent to which endogenous Siah-1 was knocked down. Data are mean ± SD from three independent experiments. *P<0.05 compared to scramble siRNA. B. Representative immunoblots of Siah-1 showing knockdown of Siah-1 by the specific Siah-1 siRNA in a dose-dependent manner but not by the scramble siRNA. C. Densitometric quantification of Siah-1 immunoblots in hippocampus neurons transfected with scramble siRNA or Siah-1 specific siRNA. Data are mean ± SD from three independent experiments. *, P<0.05 compared with scramble siRNA.

Figure 7. Siah-1 mediates the hypoxia and high glucose induced-synaptophysin degradation in neuronal cells. A. Primary cultured hippocampus neuron were transfected with either siRNA targeting Siah-1 or non-targeting scrambled control. The cells were then treated by hypoxia and high glucose for 3 days. Cells incubated in isotonic medium under normoxia condition were used as control. Protein were isolated and immunoblotted for Siah-1 and synaptophysin. Representative blots showed silencing of siah-1 significantly reversed hypoxia and high glucose induced reduction of synaptophysin; B, C. Densitometric quantification of Siah-1 and Synaptophysin in neuronal cells. Data represent mean ± SD from 4 independent experiments and are expressed as either the ratio of Siah-1 or Synaptophysin to tubulin and normalized as percentage control to the scramble-siRNA cells under Normoxia + OC condition. *, P<0.05 compared with scramble siRNA.
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physin, because increased levels of HA-tagged ubiquitin were detected on synaptophysin. In the control IP using mouse IgG pulled down, no synaptophysin or HA-ubiquitin was detected from triple transfected cell lysates (Figure 9). In combination, these data indicate that Siah-1 targets synaptophysin for ubiquitin-mediated degradation under hypoxia and high glucose conditions.

Discussion

Studies have shown that diabetes is an independent factor for cognitive dysfunction [27-29]. The positive control of blood glucose is thought to improve cognitive function in the elderly [30-32]. Yet more than high glucose level maybe involved in diabetes related cognitive dysfunction. For example, there was evidence showing that more than 75% of diabetic patients suffered from obstructive sleep apnea (OSA) [11, 33, 34]. This indicated that chronic hypoxia may also participate in the pathophysiologic changes caused by diabetes. Hypoxia has been shown to be one of the most important risk factor for cognitive impairment [35, 36]. Hypoxia could cause animal short-term memory decline in animal models [37, 38]. In this context, we hypothesized that both high glucose level and hypoxia may contribute to cognitive dysfunction cause by diabetes. Synaptic transmission is important to cognitive function. And synaptophysin as the specific marker of synapses, has been proved to be involved in neurotransmitter release, synaptic plasticity, synaptic vesicle formation and reuse [17, 39, 40]. It is involved in learning, memory, etc. and abnormal synaptophysin expression has been observed in a variety of neurodegenerative disease [41-43].

In this study we employed high glucose and hypoxia in cultured hippocampal neurons to mimic the pathological changes in diabetes, and then detected synaptophysin expression. It is quite interesting that in the different conditions we used, only the coexist of high glucose and hypoxia could cause significant down regulation of synaptophysin protein level while its mRNA levels were not significantly changed. This makes us speculate that post translational processes which regulate the synaptophysin degradation may be involved in the down regulation of synaptophysin. By using ubiquitin inhibitors MG-132 and lactacystin or lysosomal inhibitor E64 we tested synaptophysin expres-

Figure 8. Siah-1 targets synaptophysin for degradation by the ubiquitin-proteasome pathway under hypoxia and high glucose condition. A. The effect of Siah-1 on synaptophysin degradation is blocked by proteasome inhibitors. HEK293 cells were co-transfected with Myc-synaptophysin and Flag-Siah-1. HEK293 cells were then treated by 50 mM glucose under hypoxia condition for 3 day and incubated with proteasome inhibitor MG132 or cysteine protease inhibitor E64. Cells were then lysed, and an equal amount of protein from each lysate was analyzed by immunoblotting for synaptophysin and tubulin. B. Densitometric quantification of Synaptophysin in HEK293 cells. Data represent mean ± SD from 4 independent experiments and are expressed as the ratio of Synaptophysin to tubulin. *, P<0.05 compared with Myc-syp transfection alone. †, P<0.05 compared with Myc-syp and Flag-Siah-1 co-transfected cells. Syp, synaptophysin.
Siah-1 downregulates synaptophysin expression in hippocampal neurons under different conditions. Although there was slight upregulation of synaptophysin expression in MG-132 treated neurons, under normal culture condition no significant changes were observed by these inhibitors. While in high glucose and hypoxia treated neurons both ubiquitin inhibitors can improve the expression of synaptophysin, yet the lysosomal inhibitor E64 could not change the expression of synaptophysin. These results suggest the ubiquitination process may be involved in the down regulation of synaptophysin by high glucose and hypoxia.

E3 ubiquitin ligase Siah (Mammalian Homologues of Seven in Absentia) is the key enzyme degradation of Synaptophysin [18]. Siah protein, the Drosophila SINA (seven in absentia) homologous, is an E3 ubiquitin ligase which ubiquitinates the substrate protein and induces its degradation by the proteasome pathway [44, 45]. Mammalian animal SIAH proteins are highly conservative in evolution. SIAH-1a, SIAH-1b and SIAH-2 are the main forms in rodents while in human SIAH-1 and SIAH-2 dominant [46-48]. SIAH has been proved to be involved in the regulation of a variety of substrate protein, those participate in different signal transduction pathway, including cell cycle, cell differentiation, apoptosis, and neurodegenerative diseases [49-51].

We have identified the ubiquitination inhibitor MG132 can inhibit high glucose and hypoxia induced downregulation of synaptophysin, this suggest ubiquitination is involved in the degradation of synaptophysin under high glucose and hypoxia condition. To demonstrate whether Siah-1 is involved in the ubiquitination of synaptophysin under this condition. We further demonstrated that high glucose and hypoxia can promote synaptophysin ubiquitination, and increased expression of SIAH-1 can promote an increase in synaptophysin ubiquitination, and reduced expression of synaptophysin. This suggests that Siah-1 is the key factor that result in the degradation of synaptophysin. The next question is what signaling pathway may control the expression of Siah-1 under this condition.

Because the ERK pathway is involved in regulating Drosophila Sina expression [52], we hypothesized that ERK pathway maybe activated under hypoxia and high glucose condition, thus influence Siah pathway. In this research we observed high glucose or hypoxia along can cause increase in the phosphorylation of ERK, while the co-effect of high glucose and hypoxia
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induced ERK phosphorylation more significantly. To determine whether the change of ERK phosphorylation affect Siah-1 expression, we use ERK activation inhibitor U0126 or PD98059 to inhibit ERK phosphorylation. As expected, there was obvious decrease of SIAH-1 expression concomitantly with an upregulation of synaptophysin level. We hypothesized that ERK signaling pathway regulates E3 ligase Siah-1 under our high glucose and hypoxia condition.

In summary, the co-effect of high glucose and hypoxia can cause the degradation of synaptophysin this may explain partially the diabetes related cognitive dysfunction. In high glucose under hypoxia environment synaptophysin degradation is mainly regulated by the E3 ligase Siah-1.

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Disclosure of conflict of interest

None to disclose.

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