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## Methylene blue-induced neuronal protective mechanism against hypoxiareoxygenation stress

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### Abstract

Brain ischemia and reperfusion (I/R) injury occurs in various pathological conditions, but there is no effective treatment currently available in clinical practice. Methylene blue (MB) is a century old drug with a newly discovered protective function in the ischemic stroke model. In the current investigation we studied the MB-induced neuroprotective mechanism focusing on stabilization and activation of hypoxia inducible factor-1 $\alpha$  (HIF-1 $\alpha$ ) in an *in vitro* oxygen and glucose deprivation (OGD)-reoxygenation model.

**Methods**—HT22 cells were exposed to OGD (0.1% O<sub>2</sub>, 6h) and reoxygenation (21% O<sub>2</sub>, 24h). Cell viability was determined with the calcein AM assay. The dynamic change of intracellular O<sub>2</sub> concentration was monitored by fluorescence lifetime imaging microscopy (FLTIM). Glucose uptake was quantified using the 2-[N-(7-Nitrobenz-2-Oxa-1,3-Diazol-4-yl)Amino]-2-Deoxy-D-Glucose (2-NBDG) assay. ATP concentration and glycolytic enzyme activity were examined by spectrophotometry. Protein content changes were measured by immunoblot: HIF-1 $\alpha$ , prolyl hydroxylase 2 (PHD2), erythropoietin (EPO), Akt, mTOR, and PIP5K. The contribution of HIF-1 $\alpha$  activation in the MB-induced neuroprotective mechanism was confirmed by blocking HIF-1 $\alpha$  activation with 2-methoxyestradiol-2 (2-MeOE2) and by transiently transfecting constitutively active HIF-1 $\alpha$ .

**Results**—MB increases cell viability by about 50% vs. OGD control. Compared to the corresponding control, MB increases intracellular O<sub>2</sub> concentration and glucose uptake as well as

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the activities of hexokinase and G-6-PDH, and ATP concentration. MB activates the EPO signaling pathway with a corresponding increase in HIF-1 $\alpha$ . Phosphorylation of Akt was significantly increased with MB treatment followed by activation of the mTOR pathway. Importantly, we observed, MB increased nuclear translocation of HIF-1 $\alpha$  vs. control (about 3 folds), which was shown by a ratio of nuclear:cytoplasmic HIF-1 $\alpha$  protein content.

**Conclusion**—We conclude that MB protects the hippocampus derived neuronal cells against OGD-reoxygenation injury by enhancing energy metabolism and increasing HIF-1 $\alpha$  protein content accompanied by an activation of the EPO signaling pathway.

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## Introduction

Ischemic stroke is the leading cause of severe adult disability and the 4<sup>th</sup> leading cause of death in the U.S. While the incidence of stroke in the Medicare population 65 years of age has dropped about 40% in the last two decades, possibly due to the preventive interventions such as anti-hypertension and diet control<sup>1</sup>, there is no currently available treatment for ischemic stroke. Indeed, recombinant tissue plasminogen activator (rtPA) remains the only FDA approved treatment for ischemic stroke<sup>2</sup>. Unfortunately, the therapeutic window of rtPA is less than 4.5 hours, thus, less than 4 % of stroke patients have received rtPA thrombolytic therapy<sup>3,4</sup>. Development of an alternative or combined therapy for ischemic stroke is urgently needed.

Hypoxia inducible factors (HIF) are the most pertinent transcription factors in the maintenance of cellular homeostasis under ischemic/hypoxic conditions. Three isoforms of HIF have been identified: HIF-1, HIF-2, and HIF-3<sup>5</sup>. Although physiological functions and regulatory mechanisms of HIF-1 and HIF-2 are relatively well documented, the role of HIF-3 is less known, except as a negative regulator of HIF-1 and HIF-2<sup>6</sup>. HIF-1 plays critical roles as a transcriptional activator regulating various proteins in energy metabolism and in maintenance of cellular homeostasis in low O<sub>2</sub> conditions<sup>5,7</sup>. The importance of HIF can be underscored with its primary functions in angiogenesis, hematopoiesis, energy metabolism, and anti-/pro-apoptosis<sup>5,8-10</sup>. All of the primary functions of HIF-1 $\alpha$  directly or indirectly contribute to the increase the O<sub>2</sub> content in ischemic organs. HIF has two subunits:  $\alpha$  and  $\beta$  (also known as aryl hydrocarbon nuclear translocator (ARNT)). Those proteins belong to the basic helix-loop-helix-per-ARNT (bHLH-PAS) protein family<sup>11</sup>. Although both subunits are continuously synthesized, the O<sub>2</sub>-regulated HIF- $\alpha$  subunit is recognized as a critical regulatory subunit because of short half-life (< 3 minutes) under normoxia. In the presence of O<sub>2</sub>, iron, and ascorbate, the HIF- $\alpha$  subunit is rapidly degraded via proline hydroxylation in the O<sub>2</sub>-dependent degradation domain, ubiquitination by von Hippel-Lindau protein (VHL), and the proteasomal degradation pathway. Interaction between HIF-1 $\alpha$  and PHD2, which is a key enzyme catalyzing proline hydroxylation, occurs in both nucleus and cytoplasm, but recent reports state that a significant portion of HIF-1 $\alpha$  proline hydroxylation occurs in the nucleus<sup>12</sup>. Many studies focusing on pharmacological inhibition of PHD2 to study the role of HIF-1 $\alpha$  have been conducted<sup>13,14,15</sup>.

A series of metabolic and pharmacological HIF-1 $\alpha$  stabilizers under normoxia are proposed, such as pyruvate<sup>16,17</sup>, moderate level of reactive oxygen species (ROS)<sup>18,19</sup>, and PHD

inhibitor<sup>15</sup>. Stabilization of HIF protects O<sub>2</sub>-sensitive organs, such as brain and heart, by mitigating inflammatory<sup>20</sup> and apoptotic<sup>21</sup> responses, but a HIF-1-induced neuronal protective mechanism is not clear. Empirically, outcomes of HIF-1 $\alpha$  stabilization and HIF-1 activation seem varied depending on the way HIF-1 $\alpha$  is stabilized or activated.

Methylene blue, the first synthetic drug, has been used in clinics for various diseases for more than a century<sup>22,23,24</sup>. Recently MB has been proposed as a potential treatment for cancer<sup>25</sup>, hepatopulmonary syndrome<sup>26</sup>, and septic shock<sup>27,28</sup>. Furthermore, the neuroprotective function of MB, which is able to traverse the BBB, has been reported<sup>29</sup>. Our previous studies have demonstrated that MB increases ATP synthesis and minimizes ROS generation<sup>30,31</sup>. The MB-induced antioxidant function is unique compared to the traditional ROS scavengers. MB is not capable of detoxifying the glucose oxidase-generated H<sub>2</sub>O<sub>2</sub><sup>30</sup>, rather it reduces newly formed ROS by minimizing electron leakage from the mitochondrial electron transport chain (ETC), shuttling electrons from complex I to cytochrome c bypassing complex II and complex III, a primary source for superoxide generation<sup>30</sup>. Since MB increases byproducts of glucose metabolism and reduces ROS generation, it could be reasonably speculated that MB stabilizes HIF-1 $\alpha$  under a normoxic environment.

Therefore, we hypothesized, based on previous reports including ours, that MB-induced neuroprotection is mediated by stabilizing HIF-1 $\alpha$  and MB activates HIF-1 by increasing glucose metabolism along with activating the EPO-mTOR pathway and enhancing nuclear translocation.

## Materials and Methods

### OGD and reoxygenation stress model

Murine hippocampal cell line, HT22 (< 20 passages), was maintained in DMEM supplemented with 10% fetal bovine serum (FBS) and penicillin (10,000 units/ml)-streptomycin (10,000  $\mu$ g/ml). At 18 h prior to OGD and reoxygenation, HT22 cells (5,000/well, 96-well plate) were seeded in high glucose DMEM supplemented with 10% FBS and 1% penicillin-streptomycin cocktail. OGD stress was introduced by replacing media with DMEM without FBS, glucose or pyruvate (Gibco, NY, USA) in which O<sub>2</sub> concentration is maintained at 0.1% with auto-controlled N<sub>2</sub> gas injection. After 6h OGD, reoxygenation was initiated by transferring the cells to normoxic 5% CO<sub>2</sub> cell culture incubator. At the beginning of reoxygenation, Dextrose (11mM) and pyruvate (1mM) were restored to simulate *in vivo* reperfusion.

### Primary Neuron preparation

Primary neurons were generated as described previously with modifications. Briefly, cortex tissue from postnatal day 0 C57BL/6 pups was dissociated by incubating in TrypLE Express (Invitrogen) at 37 °C for 20 minutes. Single cell suspension was made by passing tissue through fire-polished glass pipettes and a cell strainer. Cells were re-suspended in neuron culture medium (neurobasal medium with 2% B27 and 1% Glutamax) and seeded on poly-L-lysine coated coverslips. After incubation in cell culture incubator at 37 °C for 3–6 h, old

medium was removed and fresh warm neuron culture medium was added. Two days after plating, cytosine arabinoside (araC; 1- $\beta$ -D-arabinofuranosylcytosine) was added to the medium to a final concentration of 5  $\mu$ M to inhibit the proliferation of glia cells. After 24 hours, old medium containing araC was replaced with fresh warm neuron culture medium. Cells were cultured for ~10 days before experiment.

### Cell viability

After 24 h reoxygenation, cell viability was tested with the calcein AM assay. Cells were washed with PBS (pH 7.0) and incubated with calcein AM (1 $\mu$ M; Anaspec, Fremont, CA, USA) in PBS for 15 min at 37 °C. Fluorescence was measured using a Tecan Infinite F200 plate reader (Maennedorf, Switzerland) with 485/530 nm excitation/emission. Percent viability was calculated by comparing to the corresponding control. Treatment groups consisted of four independent trials, each in sextuplicate. After spectrophotometric analysis, green fluorescent images were obtained using a Zeiss Observer Z1 fluorescence microscope.

### Glucose uptake, glycolytic enzyme activity, pyruvate concentration and ATP assay

The effect of MB on glucose uptake was determined by using the 2-[N-(7- Nitrobenz-2-Oxa-1,3-Diazol-4-yl)Amino]- 2-Deoxy-D-Glucose (2-NBDG) assay as previously described<sup>32</sup>. HT22 cells were seeded in 96-well plates 24 h prior to the experiment, and subjected to the treatments. HT22 cells were incubated in glucose-free Krebs Ringer HEPES (KRH) buffer (NaCl (129 mM), NaHCO<sub>3</sub> (5 mM), KCl (4.8 mM), KH<sub>2</sub>PO<sub>4</sub> (1.2 mM), CaCl<sub>2</sub> (1 mM), MgCl<sub>2</sub> (1.2 mM), HEPES (10 mM); pH 7.4) for 30 min, and in glucose free KRH buffer containing 2-NBDG (100  $\mu$ M) and different concentrations of MB (1 and 10  $\mu$ M) for 5 min. Glucose uptake was determined and photographed by spectrophotometry at 465/540 nm of excitation/emission and Zeiss Observer Z1 microscope, separately. Activity of glucose-6-phosphate dehydrogenase (G-6-PDH) and hexokinase (HK), and pyruvate concentration were examined with spectrophotometry. Enzyme activities were measured with Flexstation (Molecular devices) at 37 °C as described earlier with minor modification<sup>33</sup>. Enzyme activities were expressed as units per milligram protein. ATP concentration was measured with commercial assay kit (Invitrogen, Eugene, OR, USA)<sup>34</sup>. ATP concentrations were normalized to the protein concentration of corresponding samples measured with a protein assay kit (Thermo Scientific, Rockford, IL, USA) at 660 nm, and presented as percentage of the mean ATP concentration of normoxic control (% normoxic control) experiments.

### Intracellular O<sub>2</sub> concentration

HT22 cells were seeded at a density of 20,000 cells/mL and cultured on cell culture dishes (35mm) with a cover glass in high glucose DMEM supplemented with pyruvate (1mM), L-glutamine (4 mM), and 10% FBS. For the fluorescence life time imaging microscopy (FLTIM), cells were incubated in tris (2, 2'-bipyridyl) dichlororuthenium (II) hexahydrate (120  $\mu$ M), an oxygen sensing dye, for 2 h. Washed Cells were incubated in Dextrose (10 mM)-supplemented sterile Dulbecco's phosphate buffered saline. MB (10  $\mu$ M) and glucose oxidase (GO), as a positive control, were added during microscopy. Time resolved images were obtained on a confocal MicroTime 200 system (PicoQuant GmbH, Berlin). Excitation was provided from 470 nm pulsed diode laser operating at a 320 kHz repetition rate and it was reflected off of a 490 nm dichroic plate into an Olympus IX71 inverted microscope. The

light passed through an Olympus 60X 1.2 NA objective, and the collected fluorescence was filtered by a 488 nm long wave pass, interference filter before passing through a 50 $\mu$ m confocal pinhole. The signal from the detector was routed into time correlated single photon counting module (PicoHarp 300). Fluorescence decay curves were analyzed by the software SymPhoTime, v. 5.3.2.

### Western blot analysis

A series of protein contents were analyzed with immunoblotting (n=5) of whole cell and nuclear lysates;  $\alpha$  subunit of HIF-1 $\alpha$ , Erythropoietin (EPO), Akt and phosphorylated Akt, mTOR and phosphorylated mTOR, phosphorylated P70S6K, and prolyl hydroxylase 2 (PHD2). Whole cell protein was extracted as following; PBS-washed HT22 cells were lysed in the cell lysis buffer, containing Tri (20mM), NaCl (100mM), and EDTA (1mM), mixed with inhibitors of protease and phosphatase for 15 min at the 4 °C on shaker and followed by centrifugation at 110,000g for 20 min at 4 °C. Supernatant containing whole cell extract was saved at -80 °C for later use. Nuclear protein was extracted with commercial kit (FIVEphoton Biochemicals, San Diego, CA). Cells were extracted with cytoplasmic isolation solution mixed with inhibitors of protease cocktail and phospholipase. Cells and cytoplasmic isolation solution mixture was centrifuged at 588g for 3 min at 4 °C to separate cytoplasmic fraction. The nuclear fraction was isolated with nuclear isolation solution plus protease inhibitor cocktail and phospholipase inhibitor. Equal nuclear protein loading was confirmed with histone deacetylase 1 (HDAC1). Protein concentration of lysates was measured with a protein assay kit (Thermo Scientific, Rockford, IL, USA) at 660 nm to ensure equal loading. Protein (15 $\mu$ g/lane) was separated with SDS-PAGE electrophoresis and transferred to nitrocellulose membrane. The following primary antibodies were used; HIF-1 $\alpha$ , EPO, and PHD2 (Novus Biologicals, Littleton, CO); Akt and p-Akt (Santa Cruz, Dallas, TX); and mTOR signaling pathway (Cell signaling technology, Danvers, MA, USA). Goat anti-mouse and rabbit secondary antibodies (Jackson ImmunoResearch, West Grove, PA, USA) were used at 1:4500 dilutions for 1 hour at the room temperature. Protein contents were quantified with protein densitometry (Ultraviolet Products, Upland, CA, USA) and normalized to actin band density.

### Nuclear Translocation of HIF-1 $\alpha$

Constitutively active HIF-1 $\alpha$  plasmids (gift from Dr. JA Garcia, UT Southwestern Medical Center<sup>35</sup>) encoding constitutively active human HIF-1 $\alpha$  (by alanine substitution of the conserved Pro or Asp residues normally modified in an oxygen-dependent manner), tagged with GFP, and cloned into pIRES-hrGFP (Stratagene, La Jolla, CA) was transiently transfected to the HT22 cells. DNA-liposome complex was formed with P1P2N HIF-1 $\alpha$  plasmid (200ng)<sup>36</sup> and transfection agent (1 $\mu$ L) (Lipofectamine 2000; Invitrogen, Carlsbad, CA). DNA-liposome complex was added to the 50% confluent HT22 cells in 48 well plates, incubated 48 h in normoxic CO<sub>2</sub> incubator at 37 °C. After transfection, before and after MB treatment, live cell images of HIF-1 $\alpha$  translocation was photographed with a temperature and time controlled (37°C) Zeiss Observer Z1 fluorescence microscope. Additionally, nuclear HIF-1 $\alpha$  protein content/Cytoplasmic HIF-1 $\alpha$  content ratio was obtained from the immunoblot analysis.

### Inhibition of HIF-1

2-MeOE2 (2-methoxyestradiol) inhibits transcriptional expression of target genes regulated by HIF-1 $\alpha$  without affecting HIF-1 $\alpha$  transcription<sup>37</sup>. 2-MeOE2 inactivates HIF-1 $\alpha$  by blocking nuclear translocation of HIF-1 $\alpha$ . HT22 cells were treated with 2-MeOE2 (1 $\mu$ M) during OGD and reoxygenation. The involvement of HIF-1 activation in the MB-induced neuronal protection was confirmed with calcein AM assay with 2-MeOE2 treatment.

### Inhibition of mTOR

in order to inhibit mTOR, KU-63794 (1  $\mu$ M, EMD Milipore) was pretreated 3 h prior to the OGD. KU-63794 is known to inhibit the phosphorylation of mTOR at the Ser2448/2481.<sup>38,39</sup> Inhibition of mTOR phosphorylation was confirmed with immunoblot and effects of mTOR inhibition on MB-induced protection against OGD-reoxygenation was monitored with Calcein AM assay.

### Statistical analysis

Data was expressed as mean  $\pm$  SEM. Multiple comparisons between OGD-reoxygenation exposed groups and normoxia groups was accomplished by one way analysis of variance combined with Turkey multiple comparison test to identify statistically meaningful differences. Probability values < 0.05 were taken to indicate statistically significant effects.

## Results

### MB protects HT22 cells from OGD and reoxygenation protection

MB-induced neuroprotection was studied in hippocampus derived HT22 cells subjected to OGD/reoxygenation. Six hours of OGD and 24 h of reoxygenation killed about 45% of HT22 cells in the control group (Figure 1). MB (100 nM) treatment at the time of reoxygenation was able to significantly rescue the cells from OGD/reoxygenation injury.

### MB enhances glucose utilization

MB increases ATP after 24h reoxygenation in a dose dependent manner (Figure 2A). Increased ATP concentration with MB treatment is associated with increased glucose uptake. MB (100 nM and 1  $\mu$ M) increases glucose uptake in a dose dependent manner (25% and 50%, respectively) which was manifested with 2-deoxy-2-[(7-nitro-2,1,3-benzoxadiazol-4-yl) amino]-D-glucose (2NBDG) (Figure 2B). Immunoblot assay shows that MB increased PIP5K which is known to function in glucose uptake by stimulating endocytosis of glucose (Figure 2C). The role of PIP5K in anti-apoptosis was also reported<sup>40</sup>. MB increases intracellular O<sub>2</sub> concentration, which provides a favorable condition for aerobic glucose metabolism. (Figure 2D) MB-enhanced ATP concentration was consistent with increased activities of hexokinase (Figure 3A-a,b) and glucose-6-phosphodehydrogenase (G6PDH). (Figure 3B-a,b) This increased enzyme activity resulted in an increased endogenous pyruvate concentration. (Figure 3C-a,b)

### MB-enhanced stability of HIF-1 $\alpha$ contributes to the MB-induced neuroprotection

The role of HIF in cell survival from I/R injury has been widely reported<sup>14,15,21,41</sup>. MB significantly increases HIF-1 $\alpha$  protein content compared to both normoxic and OGD control groups (Figure 4A). EPO is a protective cytokine that is mainly regulated by activation of HIF-1. EPO was significantly increased with MB treatment (Figure 4A,B). The binding of EPO to its membrane receptor triggers the Jak-STAT signaling pathway, and, in turn, activates anti-apoptotic protein kinase, such as Akt. Akt and mTOR pathways were activated by MB treatment (Figure 4C). In the current setting, the involvement of HIF activation in the MB-induced neuroprotective mechanism was examined by blocking HIF-1 $\alpha$  translocation to the nucleus with 2-methoxyestradiol (2-MeOE<sub>2</sub>) (Figure 4D). 2-MeOE<sub>2</sub> treatment significantly abolished MB's neuroprotection against OGD-reoxygenation (Figure 4D). Activated mTOR is newly proposed as an HIF-1 regulator at the messenger level<sup>42-45</sup>. The involvement of mTOR activation in MB-induced protection was tested by treatment with KU-63794, which inhibits phosphorylation of mTOR. KU-63794 dampened MB-induced protection against OGD-reoxygenation stress. (Figure 4E)

### MB enhances nuclear translocation of HIF-1 $\alpha$

To examine the role of MB on HIF-1 $\alpha$  nuclear translocation, HT22 cells were transiently transfected with constitutively active P1P2N HIF-1 $\alpha$  or empty vector. GFP-tagged HIF-1 $\alpha$  enables us to track the HIF-1 $\alpha$  translocation. Additionally, primary neuron cultures were treated with CoCl (100 $\mu$ M) to stabilize HIF-1 $\alpha$  under normoxia. HIF-1 $\alpha$  accumulation was observed in the nucleus of both HT22 cells (Figure 5A) and primary neurons (Figure 5D) within 10 minutes of treatment with 1 $\mu$ M and 10 $\mu$ M MB, respectively. Additionally, the nuclear/cytoplasmic HIF-1 $\alpha$  ratio was significantly increased in the MB treated group *vs.* both normoxic and OGD/reoxygenation controls. (Figure 5B) It has been reported that PHD2 interacts with HIF-1 $\alpha$  located in the nucleus as well as with cytoplasmic HIF-1 $\alpha$ , thus, even after HIF-1 $\alpha$  enter the nucleus, HIF-1 $\alpha$  degradation continuously occurs. MB lowered PHD2 content in the whole cell extract compared to the both normoxic and OGD control group. (Figure 5C)

## Discussion

Although MB's neuroprotective function failed to be proven in some cases<sup>46</sup>, several laboratories have reported the neuroprotective effects of MB in I/R models<sup>30,47</sup>. The role of ROS in I/R injury in various pathological cases, such as ischemic stroke<sup>48</sup>, myocardial infarction<sup>48</sup>, traumatic brain injury<sup>49</sup>, and organ transplantation<sup>50</sup> is well documented. Previously, we reported that MB minimizes electron leakage from the electron transport chain of the mitochondria, significantly reducing ROS generation<sup>30,31</sup>. Herein, we are proposing a possible mechanism for this MB protection from *in vitro* OGD-reoxygenation stress in neurons with special emphasis on the effect of MB on HIF-1. We demonstrated that MB-induced neuronal protection is mediated via activation of HIF-1, and proposed that MB stabilizes and increases HIF-1, a regulatory subunit of HIF-1 by 1) increasing glucose metabolism, culminating in increased endogenous pyruvate, which is consistent with increased ATP concentration, 2) activating EPO and mTOR signaling pathways, and 3) stimulating nuclear localization of HIF-1 $\alpha$ .

HIF-1 is an important transcription factor for cellular adaptation to intense hypoxic conditions. Activation of HIF-1 is involved in the regulation of numerous genes in glucose metabolism, anti- and pro-apoptosis, cellular proliferation, pH regulation, angiogenesis, erythropoiesis, matrix metabolism, and metal transport<sup>51,52</sup> in the ischemic condition. HIF-1 $\alpha$  is a well characterized oxygen-dependent transcription factor<sup>53</sup>, but recently also proposed as a mTOR-dependent transcription factor<sup>54,55</sup>. The  $\alpha$ -subunit of HIF-1 is continuously produced, but readily degraded by the activation of prolyl hydroxylases in the presence of oxygen and cofactors, such as Fe<sup>2+</sup> and ascorbate<sup>56</sup>. Although, HIF-1 $\alpha$  is rapidly degraded under normoxic conditions, in the presence of normoxic HIF-1 $\alpha$  stabilizers, HIF-1 $\alpha$  could become stable in a normoxic or even a hyperoxic environment<sup>16</sup>. Normoxic HIF-1 $\alpha$  stabilizers fall into the following categories: metabolites from glucose metabolism, growth factors, antioxidants, and inflammatory cytokines. For example, insulin increases HIF-1 $\alpha$  in both the messenger and the protein levels and non-pathological concentrations of ROS also stabilize HIF-1 $\alpha$  by inhibiting PHD<sup>57</sup>. The current study showed that MB significantly increased HIF-1 $\alpha$  even 24 h after 6 h OGD. This MB-induced stabilization of HIF-1 $\alpha$  is correlated with an increased effect of MB on glucose metabolism. In the *in vitro* OGD-reoxygenation model and also in the rodent Parkinson model, MB significantly increased ATP concentration, and reduced H<sub>2</sub>O<sub>2</sub> toxicity<sup>31,32</sup>. Conceivably, the increased ATP concentration with MB treatment is closely related to the increased glucose uptake, intracellular O<sub>2</sub> concentration, and content of phosphatidylinositol 4-phosphate 5-kinase (PIP5K) protein in the MB group. PIP5K generates phosphatidylinositol 4,5-bisphosphate (PIP<sub>2</sub>), which plays a key role in increased translocation of GLUT4<sup>58</sup>. In accordance with increased ATP concentration, the activities of HK and G-6-PDH were significantly increased with MB treatment. This culminated in an increased concentration of endogenous pyruvate, an intermediary metabolite and an efficient energy yielding fuel, entering into the TCA cycle. Previously, we reported that exogenous pyruvate perfused during reperfusion in rodent transient middle cerebral artery occlusion (MCAO) model significantly reduced brain lesion volume and DNA fragmentation<sup>17</sup>. Pyruvate-induced cerebral protection against I/R injury is mediated by activation of HIF-1 $\alpha$  and the endogenous EPO signaling pathway<sup>17</sup>. Indeed, knock down of HIF-1 $\alpha$  with small interfering RNA transfection and inhibition of the EPO signaling pathway with soluble EPO receptor abrogated pyruvate-induced cerebral protection<sup>17</sup>. Similarly, Inhibition of HIF-1 $\alpha$  nuclear translocation eliminated the favorable effect of MB on neuroprotection against OGD-reoxygenation. Therefore, MB-induced neuroprotection is mediated via HIF-1 $\alpha$  stabilization and a possible mechanism of action for this MB stabilization of HIF-1 $\alpha$  is the increase of glucose metabolism, so that increased endogenous pyruvate could contribute to the mitigation of HIF-1 $\alpha$  degradation.

HIF-1 was originally characterized as an O<sub>2</sub>-dependent transcription factor, and involvement of mTOR in HIF-1 activation was recently reported<sup>44,59</sup>. In addition to MB's function in the maintenance of HIF-1 $\alpha$  stability under normoxia by increasing glucose metabolism, MB could also increase HIF-1 $\alpha$  protein content by stimulating transcription of HIF-1 $\alpha$  by activating the mTOR pathway which could be a second mechanism of MB-induced HIF-1 $\alpha$  regulation. MB activates the EPO signaling pathway including the activation of anti-apoptotic kinase, Akt, followed by activation of the mTOR pathway. Indeed, combined



treatment of MB with mTOR inhibitor decreased HIF-1 $\alpha$  protein content and MB's neuroprotective effect was significantly moderated. This suggests that activation of mTOR is involved in the sustainably elevated HIF-1 $\alpha$  protein content with MB treatment. Therefore, transiently activating HIF-1 $\alpha$  by MB treatment might lead to a second phase elevation of HIF-1 $\alpha$  and possibly to long-term recovery.

Because of the short half-life of HIF-1 $\alpha$  in normoxia, its stabilization is a critical step. Nuclear translocation of HIF-1 $\alpha$  is also an essential step for HIF activation. In HIF-1 $\alpha$  degradation, the role of PHD is important. Of special significance is the reported HIF-1 $\alpha$  degradation by direct interaction between PHD2 and HIF-1 $\alpha$  in the nucleus<sup>12</sup>. MB significantly decreased PHD2 protein content and increased nuclear HIF-1 $\alpha$  protein content in this study. Thus, the rate of HIF-1 activation should be associated with both stability of HIF-1 $\alpha$  and inhibition of interaction between PHD2 and HIF-1 $\alpha$  in the nucleus, which results in increased expression of HIF-1 $\alpha$  in the nucleus.

In conclusion, MB protects hippocampus-derived neuronal HT22 cells by enhancing glucose metabolism and activating the HIF-1/EPO signaling pathway. MB also stabilizes the  $\alpha$  subunit of HIF-1 by increasing endogenous pyruvate, a known normoxic HIF-1 $\alpha$  stabilizer (Figure 6).

### Limitation

This study provided convincing data to support the proposed hypothesis that MB protects HT22 cells from OGD and reoxygenation stress by stabilizing HIF-1 $\alpha$  and activating HIF-1. Due to possible aberrant physiological characteristics of transformed cells, *i.e.* aberrant energy metabolism, confirmation of the current findings with primary neurons have been begun to demonstrate neuron-specific HIF-1 activation in OGD-reoxygenation. Furthermore, it is highly possible that in the *in vivo* environment various endogenous factors are interconnected and might create other outcomes. Therefore, it is necessary for the findings presented here to be confirmed in *in vivo* models.

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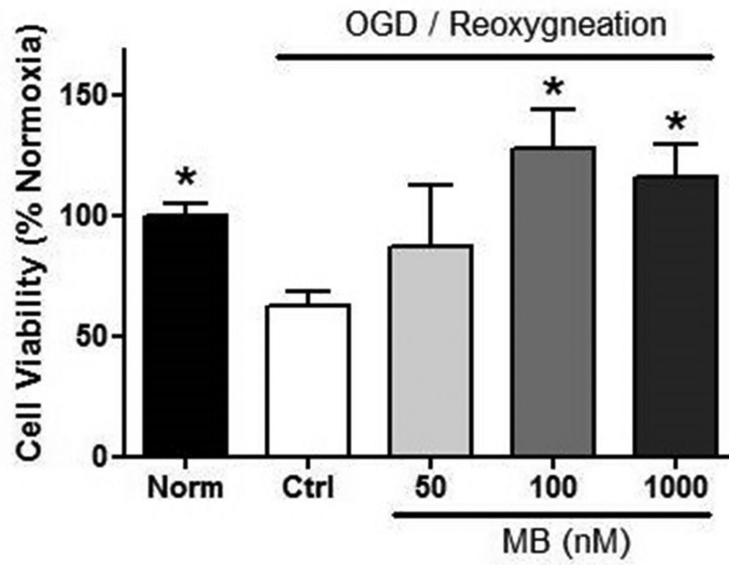
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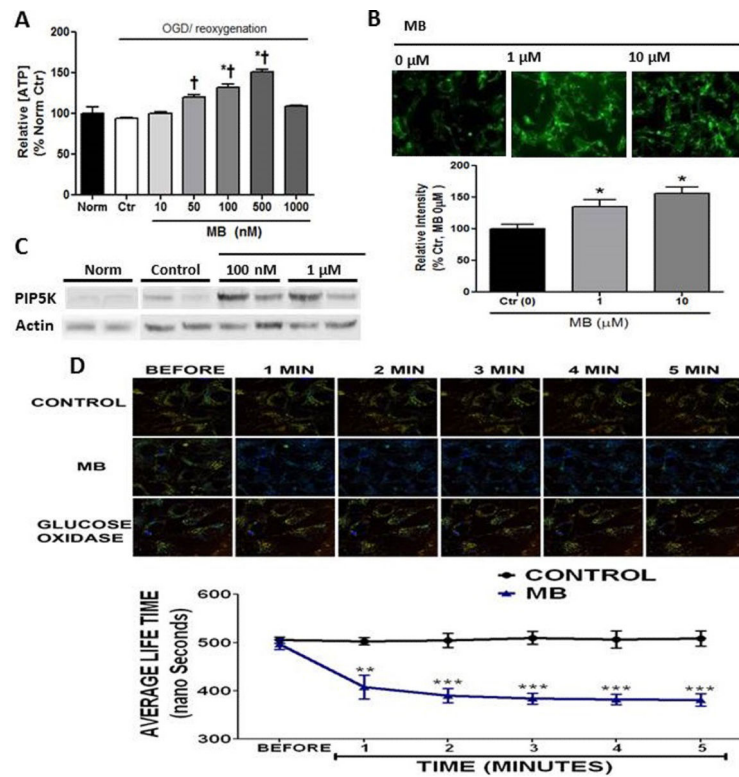
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**Highlights of submitted manuscript entitled “Methylene blue-induced Neuronal Protective Mechanism against Hypoxia-Reoxygenation Stress”**

- MB stabilizes hypoxia inducible factor-1 $\alpha$ .
- MB increased ATP production after OGD-reoxygenation stress.
- MB-induced HIF-1 $\alpha$  stabilization is associated with increased glucose metabolism.
- MB produces favorable intracellular environment for the aerobic glycolysis.

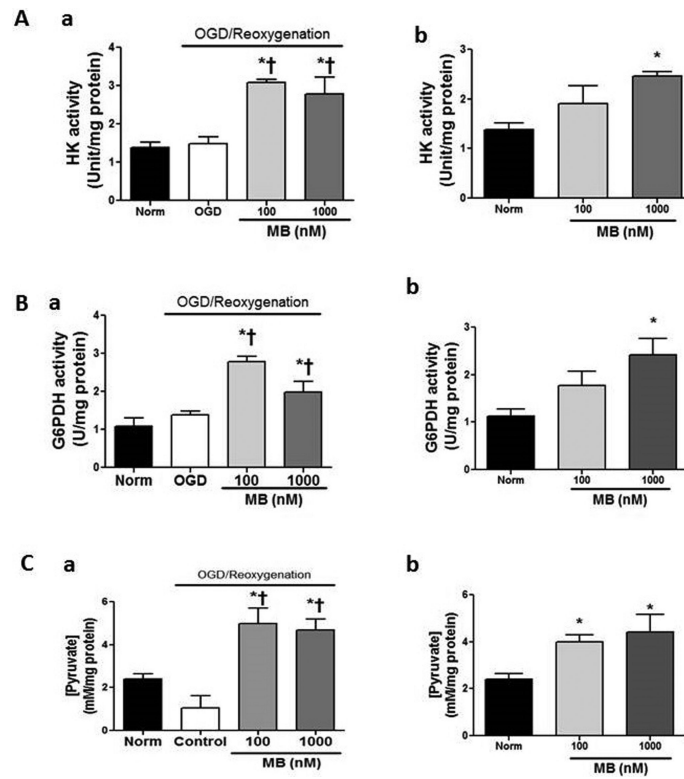


**Figure 1. MB protects HT22 cells against OGD-reoxygenation stress**  
MB treated at the reoxygenation protects HT22 cells in a dose dependent manner. †  $p < 0.05$  vs. Norm, \*  $p < 0.05$  vs. Ctr



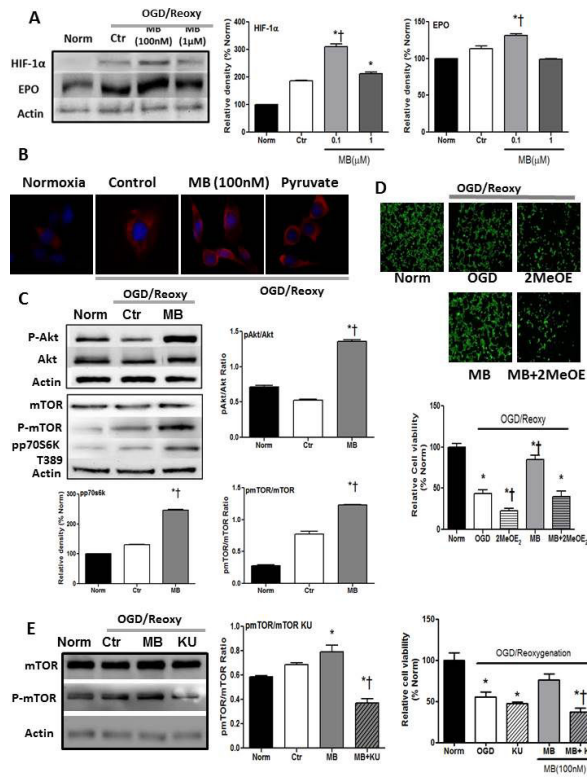
### Figure 2. MB increases energy production

**A)** ATP concentration was significantly increases in a dose dependent manner, but MB (1µM) increased toxicity. ATP value in control group:  $9.89 \pm 0.88 \mu\text{mol/g protein}$  (Mean  $\pm$  SE) **B)** MB increases glucose uptake, tested with 2-[N-(7-Nitrobenz-2-Oxa- 1,3-Diazol-4-yl)Amino]-2-Deoxy-D-Glucose (2-NBDG) assay, because of experiment limitation, relatively high doses of MB (1µM and 10µM) were used. **C)** Phosphatidylinositol 4-phosphate 5-kinase (PIP5K) protein content was increased with MB vs. normoxic and OGD controls **D)** Fluorescence life time imaging microscopy (FLTIM) showed that MB enhanced intracellular  $\text{O}_2$  concentration vs. Control. Glucose oxidase was used as a positive control.  $\dagger$   $p < 0.05$  vs. Norm, \*  $p < 0.05$  vs. Ctr

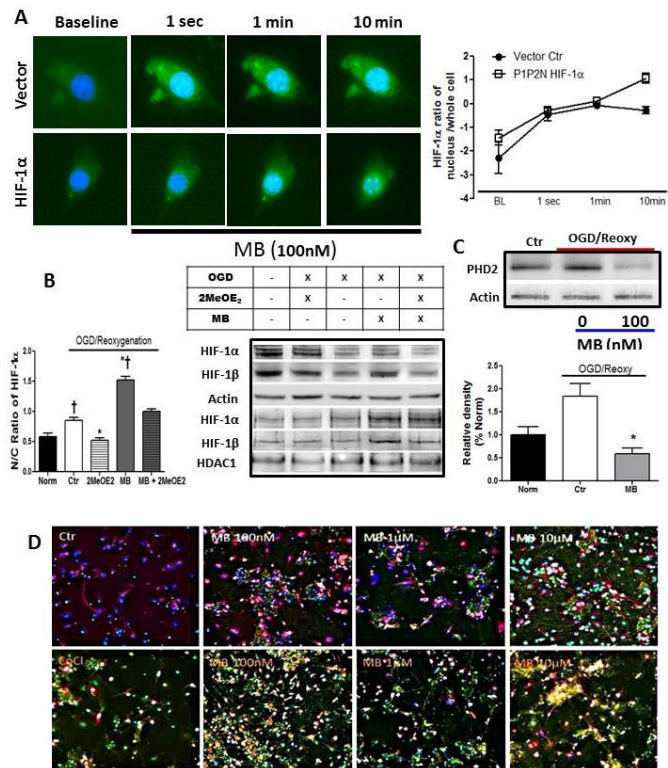


**Figure 3. MB enhances glycolytic enzyme activity and increases endogenous pyruvate**  
 Activities of hexokinase (**A**) and glucose-6-phosphate dehydrogenase (**B**) are significantly increased in both normoxia (**a**), and OGD (**b**) vs. Norm, OGD controls. **C**) Increased pyruvate concentration with MB treatment in both normoxia (**a**) and OGD (**b**) was consistent with enzyme activities. †  $p < 0.05$  vs. Norm, \*  $p < 0.05$  vs. Ctr



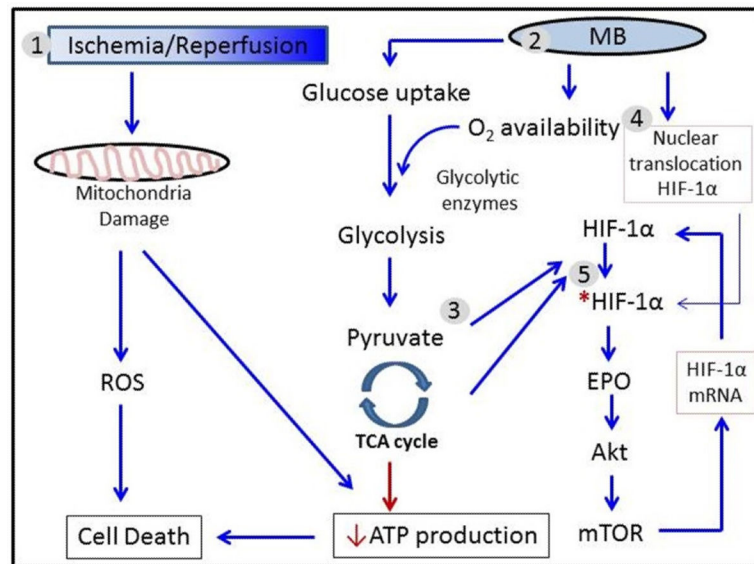


**Figure 4. MB-induced neuroprotection mediated via activation of HIF-1**  
 (A) HIF-1 $\alpha$  and EPO (A,B) protein contents are significantly increased in MB treated group.  
 C) Akt -mTOR in EPO signaling pathway are activated with MB treatment. D) Inhibition of HIF-1 activation with 2MeOE<sub>2</sub> dampened MB-induced protection against OGD-reoxygenation stress. E) Inhibition of mTOR with KU-63794 blocks protective effects of MB. n=5, \* p < 0.05 vs. Norm, † p < 0.05 vs. OGD



**Figure 5. MB stimulates nuclear translocation of HIF-1α**

**A)** GFP-tagged HIF-1α reveals the concentrated HIF-1α expression in the nucleus with MB treatment (green, P1P2N HIF-1α, blue: DAPI). **B)** The ratio of nuclear and cytosolic HIF-1α significantly increased in MB group vs. Norm and OGD groups. 2MeOE<sub>2</sub> inhibits nuclear translocation of HIF-1α. (n=4) **C)** MB significantly lowers prolyl hydroxylase 2 (PHD2) content vs. OGD control **D)** MB enhances nuclear HIF-1α expression in primary neurons which express constant HIF-1α expression with CoCl (100μM treatment. Red: Neurofilament, Green HIF-1α, Blue: DAPI. \* p < 0.05 vs. Ctr, † p < 0.05 vs. Norm



**Figure 6. Proposed mechanism of MB-induced neuronal protection against ischemia and reperfusion injury**

1) Ischemia and/or reperfusion damages cells and tissues by increased ROS and energy depletion which mediated by mitochondrial damage. 2) MB preserves mitochondria function, and increases intracellular O<sub>2</sub> concentration and glucose uptake which culminate in restoration of ATP. 3) Increased energy production increases endogenous pyruvate and other glycolytic metabolites, which stabilize HIF-1 $\alpha$ . Additionally, 4) MB enhances nuclear translocation of HIF-1 $\alpha$  followed by activation of HIF-1. 5) HIF-1 activation stimulates several protein synthesis, including EPO. Activated EPO in turn activate mTOR signaling pathway. Activated mTOR transcriptionally increases HIF-1 $\alpha$ . Therefore, MB could make HIF-1 being involved with protection of HT22 cells against ischemia-reperfusion injury. HIF-1 $\alpha$ : Hypoxia inducible factor-1 $\alpha$ ; EPO: Erythropoietin; mTOR: mammalian target of Rapamycin; ROS: reactive oxygen species;  $\downarrow$  Positive effect;  $\downarrow$  Negative effect; \* Activation.