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Preconditioning of bone marrow mesenchymal stem cells by prolyl hydroxylase inhibition enhances cell survival and angiogenesis in vitro and after transplantation into the ischemic heart of rats

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Abstract

Introduction: Poor cell survival and limited functional benefits have restricted the efficacy of bone marrow mesenchymal stem cells (BMSCs) in the treatment of myocardial infarction. We showed recently that hypoxia preconditioning of BMSCs and neural progenitor cells before transplantation can enhance the survival and therapeutic properties of these cells in the ischemic brain and heart. The present investigation explores a novel strategy of preconditioning BMSCs using the Hypoxia-inducible factor 1α (HIF-α) prolyl hydroxylase inhibitor dimethyloxalylglycine (DMOG) to enhance their survival and therapeutic efficacy after transplantation into infarcted myocardium.

Methods: BMSCs from green fluorescent protein transgenic rats were cultured with or without 1 mM DMOG for 24 hours in complete culture medium before transplantation. Survival and angiogenic factors were evaluated in vitro by trypan blue staining, Western blotting, and tube formation test. In an ischemic heart model of rats, BMSCs with and without DMOG preconditioning were intramyocardially transplanted into the peri-infarct region 30 minutes after permanent myocardial ischemia. Cell death was measured 24 hours after engraftment. Heart function, angiogenesis and infarct size were measured 4 weeks later.

Results: In DMOG preconditioned BMSCs (DMOG-BMSCs), the expression of survival and angiogenic factors including HIF-1α, vascular endothelial growth factor, glucose transporter 1 and phospho-Akt were significantly increased. In comparison with control cells, DMOG-BMSCs showed higher viability and enhanced angiogenesis in both in vitro and in vivo assays. Transplantation of DMOG-BMSCs reduced heart infarct size and promoted functional benefits of the cell therapy.

Conclusions: We suggest that DMOG preconditioning enhances the survival capability of BMSCs and paracrine effects with increased differentiation potential. Prolyl hydroxylase inhibition is an effective and feasible strategy to enhance therapeutic efficacy and efficiency of BMSC transplantation therapy after heart ischemia.
Introduction

Myocardial infarction (MI) is a leading cause of congestive heart failure that results from irreversible loss of cardiomyocytes, scar formation and ventricular remodeling [1]. A large amount of animal and clinical research has demonstrated that bone marrow mesenchymal stem cells (BMSCs) are capable of improving the post-MI recovery due to continued self-renewal, transdifferentiation and paracrine effects [2]. However, the morphological and functional benefits of BMSC therapy are limited primarily due to the low survival rate of transplanted cells in the ischemic host environment [3]. Strategies to improve BMSC tolerance to hypoxic/ischemic conditions as well as their functional benefits are therefore critically needed for clinical translation of the cell therapy.

Earlier work demonstrated that pretreatment of adult BMSCs with cardiogenic growth factors before transplantation improves the cells’ in vivo cardiac differentiation as well as functional recovery in a dog model of the chronically infarcted myocardium [4]. In a recent study, a recombinant cocktail consisting of transforming growth factor beta-1, bone morphogenetic protein-4, activin A, retinoic acid, insulin-like growth factor-1, fibroblast growth factor-2, α-thrombin, and interleukin-6 was formulated to engage human mesenchymal stem cells into cardiopoiesis. Compared with unguided counterparts, cardiopoietic mesenchymal stem cells delivered into infarcted myocardium achieved superior functional and structural benefits [5]. In order to enhance cell survival and repair potential, some studies have overexpressed anti-apoptotic and trophic/growth factors in stem cells or progenitor cells [6-9]. This molecular biological approach can effectively reduce cell death; however, it causes a practical concern for increased risk of tumor growth due to lasting and high expression of anti-apoptotic factors. Alternatively, we recently explored a new approach of hypoxic preconditioning by exposing stem cells and progenitor cells to sublethal hypoxia before transplantation [10]. This hypoxic preconditioning increased the expression of prosurvival and proangiogenic factors including hypoxia-inducible factor (HIF)-1α, angiopoietin-1, vascular endothelial growth factor (VEGF) and its receptor Flk-1, erythropoietin (EPO), Bcl-2, and Bcl-xL. Cell death and caspase-3 activation in these cells was significantly lower compared with that in normoxic cells both in vitro and after transplantation. Transplantation of hypoxic preconditioned BMSCs, neural progenitors or cardiac progenitors, resulted in greater cell survival, angiogenesis, better tissue repair and enhanced functional recovery after myocardial and cerebral ischemia [7,10-18].

It is suggested that a preconditioning procedure sets cells into a primed state before they encounter the harsh microenvironment of hypoxia/ischemia and elevated levels of injurious factors [19,20]. At present, the preconditioning strategy has been increasingly accepted and applied in cell-based transplantation therapy and shows multiple therapeutic benefits after ischemic disorders in the central nervous system and peripheral organs [21-23]. Besides hypoxic preconditioning, we and others have shown that stem cells/neural progenitors can be preconditioned by some endogenous factors and neural peptides such as growth factors, interleukin-6, apelin and EPO [4,17,18,24-26]. Similarly, pharmacological preconditioning has emerged as a means of priming transplanted cells. For example, pretreatment of skeletal myoblasts and mesenchymal stem cells with diazoxide achieved protective and functional benefits in the treatment of cardiomyocyte ischemia [27,28]. Diazoxide pretreatment also protects neurons from glutamate toxicity [29].

Prolyl hydroxylases are members of an iron-dependent and 2-oxoglutarate-dependent dioxygenase enzyme family. The HIF prolyl hydroxylase enzymes, termed the prolyl hydroxylase domain, play an important role in oxygen regulation in different tissues and organs [30]. Cells recognize and respond to hypoxia by accumulating the transcription factor HIF-1, composed of oxygen-sensitive inducible HIF-1α and constitutive HIF-1β subunits. Prolyl hydroxylase domain enzymes are involved in the degradation of the HIF-1α subunit. Under hypoxic conditions, the lack of oxygen leads to stabilization of HIF-1α to form a HIF heterodimer that is subsequently translocated to the nucleus, triggering the transactivation of target genes. The nature of the target gene and type of expressed proteins may vary depending upon the type of tissues and disease conditions [31]. Similar effects can be obtained using the prolyl hydroxylase inhibitor, dimethylxalylglycine (DMOG). HIF-1α in turns activates the transcription of a number of angiogenic and survival genes such as VEGF, glucose transporter 1 (Glut-1), and EPO [32-36]. Our previous study showed that DMOG enhanced cell survival against the apoptotic insult of serum deprivation in a dose-dependent manner in BMSC cultures, mediated by the mechanisms of HIF-1α stabilization, regulation of the mitochondrial pathway and phosphatidylinositol-4, 5-bisphosphate 3-kinase (PI3K)/Akt activation [37]. DMOG treatment reduced mitochondrial cytochrome c release, prevented nuclear translocation of apoptosis inducing factor, and promoted Akt phosphorylation [37]. In a subsequent investigation, we demonstrated in neuronal cultures that DMOG (100 μM) reduced cell death induced by oxygen glucose deprivation [34]. In a cerebral ischemic stroke model, DMOG treatment (50 mg/kg, intraperitoneally) enhanced HIF-1α activation and transcription of its downstream genes VEGF, EPO, endothelial nitric oxide synthesis, and pyruvate dehydrogenase kinase-1. The DMOG treatment reduced caspase-3 activation and infarct formation and improved functional recovery after stroke [34]. These data suggest that the oxygen-sensing pathway is a
potent protective mechanism. The present investigation explores the hypothesis that DMOG, as an inhibitor of the oxygen-regulated enzyme prolyl hydroxylase, can mimic the hypoxic environment to prime BMSCs for an effective transplantation therapy after heart ischemia.

**Methods and materials**

**Bone marrow mesenchymal stem cell cultures**

BMSCs from green fluorescent protein transgenic rats were isolated and harvested as described previously [38]. In brief, bone marrow tissues were acquired by flushing the cavities of femurs and tibias with basal Dulbecco’s modified Eagle’s medium. Collected bone marrow cells were seeded into flasks with Dulbecco’s modified Eagle’s medium supplemented with 10% fetal bovine serum. Cells were cultured at 37°C in a humidified environment of 5% carbon dioxide. Nonadherent cells were removed 24 hours later, and adherent cell colonies were washed three times with phosphate-buffered saline solution (PBS). Fresh complete medium was added and changed every 3 to 4 days. Cells were subcultured 1:2 or 1:3 when they reached 80% confluence. BMSCs of four to six passages were used in this study.

For identification of BMSCs, cell surface markers CD90, CD34 and CD45 were detected by fluorescence-activated cell sorting. The detailed characterization of our isolated bone marrow cells and multipotency of identified BMSCs were confirmed in our previous investigations [39]. In the present investigation, CD90+/CD34−/CD45− cells were selected and tested.

**Dimethlyoxalylglycine preconditioning of BMSCs**

Cells were subcultured at 1:2/1:3 and cultured for 2 to 3 days until they reached 70 to 80% confluence. Cells were then exposed to fresh complete medium supplemented with 1 mM DMOG for 24 hours. The DMOG concentration was selected based on our previous investigations [37]. For the nonpreconditioned control, culture medium was changed to fresh complete medium at the same time as DMOG treatment. Before transplantation, cells were labeled with Hoechst 33342 by adding a final concentration of 10 μM/ml in the culture medium and were incubated for 2 hours to trace BMSCs after transplantation. BMSCs were then washed six times with PBS to remove unbound Hoechst dye, and digested with 0.25% (w/v) trypsin ethylenediamine tetraacetic acid, followed by suspension in complete medium. After several centrifugations and PBS washes, cells were suspended in a serum-free medium at 1 × 10^6 cells per 150 μl. This solution was injected into five sites (30 μl each) in peri-infarct myocardium 30 minutes after the ligation of the left anterior descending coronary artery as described previously [40]. For control, MI rats received the same volume of serum-free/cell-free medium. Four groups of 10 rats each were randomly divided as follows: sham-operated control group; MI with injection of medium as MI-medium control; MI with transplantation of nonpreconditioned BMSCs as N-BMSC control; and MI with transplantation of DMOG-pretreated BMSCs as DMOG-BMSC control.

**Myocardial infarction model of rats**

All animal experiments and surgical procedures were approved by the University Animal Research and Use Committee (IACUC, Emory University; No. 2001421) and met National Institutes of Health standards. Wistar rats were subjected to general anesthesia with 4% chloral hydrate (4 mg/kg intraperitoneally) and ventilated with room air using a small animal ventilator (Vetronics, Lafayette, IN, USA). MI was induced by ligation of the left anterior descending coronary artery with a 6–0 silk suture [10,40]. Successful performance of coronary artery occlusion was verified by the blanching of the myocardium distal to the coronary ligation.

**Cell death assessments of Trypan Blue staining in vitro**

To evaluate the protective effect of the DMOG preconditioning in vitro, cell death induced by hydrogen peroxide (100 μM) in a serum-free medium (90 minutes) was tested. This combination insult was intended to mimic the microenvironment of a myocardial ischemic attack [41]. Trypan Blue staining was applied to assess cell death. Serum-free medium containing 0.05% Trypan Blue was added for 10 minutes and phase contrast images were taken in five randomly chosen fields per well. The cell death percentage was determined by the ratio of Trypan Blue-positive cells to the total number of cells.

**Terminal deoxynucleotidyl transferase biotin-dUPT nick end labeling in heart sections**

The terminal deoxynucleotidyl transferase biotin-dUPT nick end labeling (TUNEL) staining kit (DeadEnd™ Fluorometric TUNEL system; Promega, Madison, WI, USA) was used for detecting cell death in heart sections. After fixing with 10% buffered formalin phosphate (Fisher Scientific, Pittsburgh, PA, USA) for 10 minutes, heart sections were pretreated with −20°C ethanol:acetic acid (2:1) and 0.2% Triton X-100. The slices were then incubated with an equilibration buffer as specified by the TUNEL kit. Sequentially, the TdT enzyme and nucleotide mix were added at proportions instructed by the kit for 75 minutes in a humidified chamber at room temperature. The slices were then washed with 2× SSC washing buffer for 15 minutes, and finally washed three times with PBS. In the end, the sections were slip covered and examined under a fluorescent microscope.
Western blot analysis

Cells were harvested and lysed with modified RIPA buffer (50 mM HEPES, pH 7.3, 1% sodium deoxycholate, 1% Triton X-100, 0.1% SDS, 150 mM NaCl, 1 mM ethylenediamine tetraacetic acid, 1 mM Na3VO4, 1 mM NaF, and protease inhibitor cocktail (Roche, Nutley, NJ, USA)). Cells were vortexed repeatedly until completely lysed, followed by centrifugation at 14,000 × g for 20 minutes. The concentration of proteins from different groups was determined using the Bicinchoninic Acid Assay (Sigma, St Louis, MO, USA). All of the above steps were performed at 4°C.

Proteins (30 to 50 μg) per sample were electrophoresed on a 6 to 18% gradient gel by SDS-PAGE in a Hoefer Mini-Gel system (Amersham Biosciences, Piscataway, NJ, USA) and transferred in a Hoefer Transfer Tank (Amersham Biosciences) to a PVDF membrane (BioRad, Hercules, CA, USA). After blockage by 5% milk in Tris-buffered saline with 0.1% Tween at room temperature for 2 hours, membranes were incubated with specific primary antibodies overnight at 4°C. Alkaline phosphatase/horseradish peroxidase-marked secondary antibodies were then conjugated at room temperature for 2 hours. Finally, the expression signals were detected with BCIP/NBT solution (Sigma) and analyzed by Image-Pro Plus software (NIH, Bethesda, MD, USA).

Tube formation test

The tube formation test is a well-established assay used to detect the formation of three-dimensional vessels and used widely to assess angiogenesis in vitro. Matrigel (Sigma) was prepared in 24-well plate according to the manufacturer's instructions. DMOG-preconditioned (DMOG-BMSCs) or nonpreconditioned BMSCs were seeded into the coated wells at a density of 50,000 cells, and were incubated for 6 hours at 37°C for full development of vessel-like tube structures. Tube formation was quantified by the cumulative tube length. In the case where several tube-like structures merged together or branched, the total length of tubes was calculated as the sum of the length of the individual branches.

Immunofluorescence staining

For immunofluorescence staining, heart slices were fixed with 10% formalin for 10 minutes, followed by permeabilization with 0.2% Triton X-100 for 5 minutes and blocked with 1% fish gelatin (Sigma) for 1 hour at room temperature. Specific primary antibodies were incubated overnight at 4°C in a humidified environment, and washed three times with PBS. Specimens were then incubated with Cy3-conjugated Donkey anti-rabbit IgG (1:500; Jackson ImmunoResearch, West Grove, PA, USA) or Alexa Fluor 488 anti-goat IgG (1:200; Molecular Probes, Carlsbad, CA, USA) for 1 hour at room temperature. Nuclear staining was performed by treatment with Hoechst 33342 (1:20,000; Molecular Probes, Carlsbad, CA, USA) for 5 minutes. Slices were then mounted and observed under a florescent microscope (BX51: Olympus, Center Valley, PA, USA).

Evaluation of cardiac function

Transthoracic echocardiography and hemodynamic measurement were performed for the analysis of cardiac function 4 weeks after myocardial ischemia. Rats were anesthetized with 4% chloral hydrate by intraperitoneal injection. Echocardiography was performed using a VisualSonics Vevo 2100 system (VisualSonics, Inc., Toronto, ON, Canada) with a 12.0 MHz transducer. The left ventricular end-diastolic diameters and left ventricular end-systolic diameters were measured by two-dimensional targeted M-mode, and the left ventricular ejection fraction (LVEF) was calculated:

\[
\text{LVEF} = \frac{\text{left ventricular end-diastolic volume} - \text{left ventricular end-systolic volume}}{\text{left ventricular end-diastolic volume}} \times 100
\]

Using echocardiographic evaluation, hemodynamic measurement was performed. After right carotid artery exposure, a microtip catheter was cannulated into the left ventricle and was connected with a MLT0699 disposable pressure transducer (AD Instrument, Colorado Springs, CO, USA). The left ventricular systolic pressure (LVSP), left ventricular end-diastolic pressure (LVEDP), maximum change rate of left ventricular pressure rise and fall (±dp/dt), time constant of the isovolumic pressure decline (Tau) and heart rate were monitored and recorded using the Powerlab/800 data acquisition system (AD Instrument).

Measurement of infarct size

Rats were sacrificed after measuring hemodynamics by an overdose of anesthetic. Hearts were harvested quickly and split into transverse sections: apex, mid left ventricle, and base. The tree sections of the hearts were embedded in optimal cutting temperature compound (Sakura Finetek USA Inc., Torrance, CA, USA). Samples cut at 10 μm thickness were stained with Masson’s trichrome and images of each slide were digitized through the NIH image analysis system. The percentage of infarct size/fibrotic area was calculated by dividing the sum of epicardial and endocardinal circumference of the infarcted area by the sum of the total endocardial and epicardial circumference of the left ventricle [42].

Statistical analysis

Data were analyzed with SPSS 13.0 (IBM Corp. Armonk, NY, USA) and expressed as mean ± standard error of mean. Student’s two-tailed t test was applied for
comparison of two independent experimental groups and one-way analysis of variance followed by the Tukey post hoc test for multiple comparisons. Statistical significance was defined as $P < 0.05$.

**Results**

**Upregulation of survival and angiogenic factors in DMOG-preconditioned BMSCs**

To evaluate the effect of DMOG preconditioning on BMSCs, we first analyzed the expression of survival and angiogenic proteins in control BMSCs (C-BMSCs) and DMOG-treated BMSCs (DMOG-BMSCs) using western blot analysis. HIF-1α, VEGF, Glut-1 and phospho-Akt were detected in both groups of BMSCs. However, significantly higher expressions of HIF-1α and the downstream factors Glut-1 and VEGF were seen in DMOG-BMSCs compared with C-BMSCs. Akt activation was markedly increased by DMOG preconditioning so that the ratio of phospho-Akt/Akt increased approximately 10-fold compared with C-BMSCs (Figure 1).

**Dimethyloxalylglycine preconditioning reduced BMSC cell death in vitro and after transplantation**

To examine whether DMOG-induced preconditioning could increase the tolerance of BMSCs, cell death was first examined in BMSC cultures, which helped to determine the protective effect of different durations of DMOG pretreatment. In the Trypan Blue assay, hydrogen peroxide (6 to 24 hours) caused significantly more than 50% cell death in C-BMSC cultures, while DMOG pretreatment of 6, 12 and 24 hours showed time-dependent protective effects. Trypan Blue-positive cells were significantly reduced in DMOG-treated cells, with the strongest protection induced by 24-hour DMGO pretreatment (Figure 2). The cell death rate in this group decreased from 52.4 ± 4.8% to 18.1 ± 0.4% ($n = 4$, $P < 0.05$). Based on this observation, we selected the 24-hour DMOG exposure in our preconditioning procedure in the following experiments.

To evaluate whether DMOG-BMSCs could show enhanced survival *in vivo*, C-BMSCs and DMOG-BMSCs were implanted into the peri-infarct region 30 minutes after the left anterior descending ligation in rats. Animals were sacrificed 24 hours later to identify the fate of transplanted BMSCs. This time point was selected because the majority of cell death after transplantation occurred within 24 hours [3]. Cell death was identified by the ratio of TUNEL/Hoechst/green fluorescent protein co-labeled cells versus total Hoechst/green fluorescent protein-positive cells (Figure 2B,C,D,E,F,G). In the

![Figure 1](http://stemcellres.com/content/5/5/111)

**Figure 1** Effect of dimethyloxalylglycine preconditioning on the expression of prosurvival and angiogenic factors. (A) Rat bone marrow mesenchymal stem cells (BMSCs) were cultured with or without dimethyloxalylglycine (DMOG, 1 mM) for 24 hours. Hypoxia-inducible factor (HIF)-1α, glucose transporter 1 (Glut-1), vascular endothelial growth factor (VEGF) and phospho-Akt were then detected by western blotting. Beta-actin was used as the loading control. (B) Densitometric analysis was applied for comparison of the relative expression levels of different proteins in DMOG-BMSCs with respect to vehicle control BMSCs (C-BMSCs) that is arbitrarily presented as 1. $n = 3$ independent assays. *$P < 0.05$ compared with C-BMSC group. (C) Ratio of phospho-Akt against total Akt expression, quantified from the western blot gels. A large increase of 10-fold Akt phosphorylation was seen in DMOG-BMSCs compared with control BMSCs. $n = 3$ independent assays. *$P < 0.05$ vs. control cells.
C-BMSC group, 87.3 ± 2.2% of total transplanted cells were TUNEL-positive. While transplanted DMOG-BMSCs showed significantly less death, there were 62.8 ± 3.1% TUNEL-positive cells among the total transplanted cells (Figure 2F).

**Dimethyloxalylglycine preconditioning enhanced angiogenic activities of BMSCs in vitro and after transplantation**

The effect of DMOG preconditioning on BMSC-induced angiogenesis was examined in vitro and in vivo. For in vitro studies, tube formation stimulated by Matrigel was tested. Although C-BMSCs were able to form vessel tubes in response to Matrigel, DMOG-BMSCs formed more tubes and the total tube length was significantly longer (1.9-fold) than that formed by C-BMSC (Figure 3A,B,I). To assess angiogenic activity after transplantation, we stained the endothelial marker von Willebrand factor in heart sections. Both BMSC transplantation groups presented increased vessel density/area compared with the MI control group. Transplantation of DMOG-BMSCs showed even...
greater angiogenesis in the ischemic heart. More vessels were observed in the DMOG-BMSC group compared with the C-BMSC group (Figure 3C,D,E,F,G,H,J).

**Reduced infarct size after transplantation of BMSCs**
Masson’s trichrome staining was used to assess fibrosis and scar formation 4 weeks after myocardial infarction and BMSC transplantation. Obvious infarct and scar formation was observed in the ischemic hearts that received cell-free medium injection. In comparison, the infarct size in both C-BMSC and DMOG-BMSC implantation groups was significantly smaller. Transplantation of DMOG-BMSCs resulted in the smallest infarct size in the ischemic heart (Figure 4).

![Figure 3](http://stemcellres.com/content/5/5/111)

**Figure 3** Effect of dedimethyloxalylglycine preconditioning on angiogenesis in vitro and after transplantation. (A), (B) Tube formation test stimulated by Matrigel was performed to identify the angiogenic activity of control bone marrow mesenchymal stem cells (C-BMSCs) and dedimethyloxalylglycine-preconditioned BMSCs (DMOG-BMSCs) in vitro. (C) to (H) Angiogenesis was inspected using von Willebrand factor staining (red) in heart sections from the myocardial infarction (MI), C-BMSC and DMOG-BMSC groups 4 weeks after MI. Hoechst staining (blue) shows the total cells. (I) Summary of total tube length measured in (A) and (B). The total tube length in the C-BMSC group was arbitrarily presented as 1. n=3 independent measurements. (J) Summary of total vessel density in different groups of in vivo experiments. n=8 animals in each group. *P<0.05 compared with C-BMSC group; #P<0.05 compared with MI control group.
Improved cardiac function after BMSC transplantation

The ischemic heart functional recovery after BMSC transplantation was measured by the changes in LVSP, LVEDP, +dp/dt, −dp/dt, Tau and LVEF 4 weeks after ischemia and treatments. Compared with the MI control group, transplantation of both C-BMSCs and DMOG-BMSCs facilitated better cardiac recovery, shown as increased LVSP and absolute values of ± dp/dt, and decreased Tau and promotion of LVEF (Figure 5). However, C-BMSC transplantation had no effect on the value of LVEDP, while DMOG-BMSC transplantation significantly reduced the LVEDP value to a near-normal level. Furthermore, even better recovery shown as significantly higher LVSP, enhanced ± dp/dt, lower LVEDP and Tau, and ameliorated LVEF was seen in MI rats receiving DMOG-BMSCs (Figure 5).

Discussion

The present investigation demonstrates for the first time that the prolyl hydroxylase inhibitor DMOG can be used to prime BMSCs in cell-based transplantation therapy after heart ischemia. We show that DMOG-pretreated BMSCs are more resistant to death both in vitro and after transplantation into the ischemic heart. DMOG-BMSCs show enhanced angiogenic activities. Survival and angiogenic factors such as HIF-1α, VEGF, Glut-1 and phospho-Akt increased after DMOG preconditioning, a typical gene regulation usually seen with hypoxic preconditioning. Transplantation of DMOG-BMSCs also leads to better recovery of the heart function in the rat’s MI model. It is thus likely that, as a regulator of the oxygen sensing system, DMOG can be used as an alternative way of preconditioning stem cells/progenitors. Like hypoxic preconditioning, DMOG preconditioning can optimize BMSC viability and regenerative capability for better engraftment and/or functional benefit in the harsh environment of myocardial infarction.

We and others have shown that the prolyl hydroxylase inhibitor DMOG prevents acute damage in the ischemic heart and brain as well as ameliorating BMSC cell death under pathological conditions [34,37,43]. However, DMOG has not been tested as a preconditioning reagent for stem cell therapy. So far, there has been only one related report showing that intraperitoneal injection of DMOG 48 hours before a skin ischemic insult could improve skin flap survival [44]. Fewer apoptotic cells were present in the ischemic flaps of DMOG-treated mice. Marked increases in circulating endothelial progenitor cells and bone marrow proliferative progenitor cells were observed after DMOG treatment [44]. We can now consistently
Figure 5 Effect of bone marrow mesenchymal stem cell transplantation on functional recovery after heart ischemia. Heart function was measured 4 weeks after myocardial infarction (MI) and cell transplantation treatment. MI rats showed functional deficits in all measurements of (A) left ventricular systolic pressure (LVSP), (B) left ventricular end-diastolic pressure (LVEDP), (C) maximum change rate of left ventricular pressure rise and fall (+dp/dt), (D) minimum change rate of left ventricular pressure rise and fall (–dp/dt), (E) time constant of the isovolumic pressure decline (Tau), and (F) left ventricular ejection fraction (LVEF). BMSC transplantation improved functional parameters of MI rats. Rats received dedimethyloxalylglycine-preconditioned BMSCs (DMOG-BMSCs) showed the enhanced functional recovery in all four functional parameters. *P <0.05 compared with MI-only group. #P <0.05 compared with the control BMSC (C-BMSC) group.
show that modulation of prolyl hydroxylases provides a new method of triggering endogenous beneficial mechanisms without applying hypoxic treatments.

Current strategies of BMSC transplantation achieve only modest recovery of cardiac deficits largely because of acute-phase cell death after implantation into the infarcted myocardium [3]. One limitation is that, as multipotent cells, the transdifferentiation efficiency of BMSCs into cardiac lineage cells is low and their engraftment in the ischemic heart is not clear [45]. The present investigation focuses on the survival and increased paracrine factors in preconditioned cells. Whether transplanted BMSCs became cardiomyocytes or vascular endothelial cells and whether DMOG preconditioning enhances the transdifferentiation in vivo remain to be examined.

In cell transplantation therapy, besides selections of ideal timing and the route of cell transplantation, it has also become important to test gene modification and cell preconditioning for increased cell survival [2,7,25,46,47]. A concern with regards to gene modification is whether or not permanent gene modification would increase the risk of tumorigenesis. Permanent or long-term gene modification in cell-based therapy may therefore be of limited use in clinical applications [48]. Since preconditioning strategies usually modify related genes in a relatively short period (days to weeks), this approach may be more feasible for clinical applications. To this end, hypoxic preconditioning has been shown highly effective in transplantation therapy using several stem cells and neural progenitor cells after ischemic stroke and other disorders [21,39]. For example, our group was among the first to report that hypoxic preconditioning could be applied to stem cell therapy for enhanced cell survival and optimized regenerative capabilities including enhanced angiogenesis, neurogenesis, suppressing inflammatory response, improved directed cell migration, homing to the ischemic region, and finally better functional recovery after heart and brain ischemia [7,15-17]. Since then a number of preconditioning mediators and pharmacological reagents have been tested for the preconditioning of different cells [18,19].

Our previous findings showed that both the HIF-1α pathway and the PI3K/Akt signaling were involved in the DMOG protective mechanisms [37]. In accordance with this, we found HIF-1α and Akt pathways were both activated in DMOG-BMSCs. HIF-1α plays an important role in vascular development and embryonic lethality. In HIF-1α−/− mice, defects in angiogenesis have been observed in both the yolk sac and the developing embryonic tissues [49]. HIF-1α stabilization and enhanced VEGF expression followed by prolyl hydroxylase inhibition increased lung angiogenesis in the primate model of bronchopulmonary dysplasia, a chronic form of lung disease [50]. HIF-1α stabilization is a major stimulus for increased VEGF production that plays a pivotal role in angiogenesis. Activation of the PI3K/Akt pathway can also increase VEGF secretion both by HIF-1-dependent and HIF-1-independent mechanisms [51]. These observations are consistent with our observation that HIF-1α and the PI3K/Akt pathway, as well as some downstream molecules such as VEGF, contribute to the protective and regenerative benefits with DMOG preconditioning of BMSCs.

Conclusion

We conclude that targeting an oxygen sensing system such as prolyl hydroxylase provides a new promising pharmacological approach for enhanced survival of BMSCs, increased paracrine signaling, augmented regenerative activities and improved functional recovery in cell transplantation therapy for the ischemic heart. The present data and previous evidence support a possible preconditioning benefit of increased transdifferentiation of BMSCs into cardiac lineage cells.

Abbreviations
BMSC: bone marrow mesenchymal stem cell; C-BMSCs: control BMSCs; DMOG: dedimethyloxalylglycine; DMOG-BMSCs: DMOG-preconditioned BMSCs; EPO: erythropoietin; Glut-1: glucose transporter 1; HIF: hypoxia-inducible factor; LVEDP: left ventricular end-diastolic pressure; LVEF: left ventricular ejection fraction; LVSP: left ventricular systolic pressure; MI: myocardial infarction; PBS: phosphate-buffered saline solution; PI3K: phosphatidylinositol-4,5-bisphosphate 3-kinase; TUNEL: terminal deoxynucleotidyl transferase biotin-DUPT nick end labeling; VEGF: vascular endothelial growth factor.

Competing interests
The authors declare that they have no competing interests.

Authors’ contributions
XL was responsible for study conception and design, carrying out experiments, acquisition and analysis of data and interpretation of data, and wrote the original manuscript. J-AW was responsible for conception development, data analysis, manuscript revision and funding support. X-YJ performed experiments, data analysis and manuscript revision. SPY was responsible for conception and experimental design, data analysis and interpretation, manuscript writing and revision, and funding support. LW was responsible for conception and project design, image analysis, data interpretation, manuscript organization and revision, and funding support. All authors read and approved the final manuscript, and agree to be responsible for the originality and reliability of the data in the paper.

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