Human Amnion-Derived Mesenchymal Stem Cells Protect Human Bone Marrow Mesenchymal Stem Cells against Oxidative Stress-Mediated Dysfunction via ERK1/2 MAPK Signaling

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Epidemiological evidence suggests that bone is especially sensitive to oxidative stress, causing bone loss in the elderly. Previous studies indicated that human amnion-derived mesenchymal stem cells (HAMSCs), obtained from human amniotic membranes, exerted osteoprotective effects in vivo. However, the potential of HAMSCs as seed cells against oxidative stress-mediated dysfunction is unknown. In this study, we systemically investigated their antioxidative and osteogenic effects in vitro. Here, we demonstrated that HAMSCs significantly promoted the proliferation and osteoblastic differentiation of H₂O₂-induced human bone marrow mesenchymal stem cells (HBMSCs), and down-regulated the reactive oxygen species (ROS) level. Further, our results suggest that activation of the ERK1/2 MAPK signal transduction pathway is essential for both HAMSCs-mediated osteogenic and protective effects against oxidative stress-induced dysfunction in HBMSCs. U0126, a highly selective inhibitor of extracellular ERK1/2 MAPK signaling, significantly suppressed the antioxidative and osteogenic effects in HAMSCs. In conclusion, by modulating HBMSCs, HAMSCs show a strong potential in treating oxidative stress-mediated bone deficiency.

INTRODUCTION

Bone deficiency is an emerging medical and socioeconomic challenge among patients requiring dental implants. It not only impairs the long-term stability of dental implants, but also the osseointegration of implants and bone (Maire, 1997). Further, bone deficiency significantly affects patients’ appearance, decreases the stability of dentures, and increases the difficulty of restoring oral function (Hertz, 1956). Increasing evidence suggests a biochemical link between increased oxidative stress and bone deficiency (Basu et al., 2001). Increase in oxidative stress and decreased capacity for antioxidant defense (Muthusami et al., 2005) results in decreased bone mineral density (BMD) in postmenopausal women (Ozgocmen et al., 2007; Sendur et al., 2009; Witko-Sarsat et al., 1996). Excessive production of reactive oxygen species (ROS), due to oxidative stress triggers bone resorption with a direct effect on osteoclast-induced superoxide generation and bone deficiency (Sontakke and Tare, 2002; Yang et al., 2001). The results indicate that oxidative stress is a crucial factor underlying bone deficiency (Sendur et al., 2009), inhibits osteoblastic differentiation (Fatouk et al., 2008; Mody et al., 2001) and increases osteoclast differentiation and function (Sontakke and Tare, 2002).

For instance, antioxidants such as ophiopogonin D (Huang et al., 2015), resveratrol (Haigis and Sinclair, 2010) and flavones (Zeng et al., 2014), have been used to treat oxidative stress-induced bone deficiency. The disadvantages include remarkable individual differences and limited availability. Tissue engineering using appropriate seed cells has shown great potential in the treatment of bone deficiency (Ito et al., 2006; Kim et al., 2009; Wang et al., 2009; Zhao et al., 2009). Human amnion-derived mesenchymal stem cells (HAMSCs), obtained from human amniotic membrane (AM) are a readily available and highly abundant tissue (Bourne, 1960), with substantial benefits as seed cells. Their low anti-inflammatory properties and fewer ethical concerns compared with other sources of stem cells are clear advantages (Leyva-Leyva et al., 2013). A previous study has confirmed that HAMSCs provide an environment conducive to osteogenic differentiation in human bone marrow mesenchymal stem cells (HBMSCs)(Wang et al., 2014). We, therefore, hypothesized that HAMSCs play a role in oxidative stress-induced bone deficiency. In this study, we used a Transwell co-culture system to investigate the in vitro effects of HAMSCs on osteogenic differentiation in H₂O₂-induced HBMSCs. Interestingly, we found that HAMSCs promoted proliferation, activated alkaline phosphatase activity (ALP), enhanced the expression of osteogenic marker proteins and stimulated mineralized matrix deposition. We showed that HAMSCs were not only capable of promoting osteogenic differentiation, but also reversing...
the oxidative stress-induced suppression of osteogenic differentiation.

Bone formation, associated with the differentiation of mesenchymal stem cells, is regulated by multiple signaling pathways, including Wnt-β-Catenin, MAPK, Hedgehog and NF-κB (Galli et al., 2010; Hammond and Schulte-Merker, 2009; Krum et al., 2010; Schindeler and Little, 2006; Simmons et al., 2003). Signal transduction pathways involved in oxidative stress-induced inhibition of osteogenic differentiation consist of MAPK, Akt/mTOR/4EBP1, p53 and NF-κB (Martindale and Holbrook, 2002; Zeng et al., 2014). Previous studies revealed that ERK1/2 MAPK signaling regulates the differentiation and proliferation of HBMSCs (Salasynyk et al., 2004; Xia et al., 1995; Zhang and Liu, 2002). Further, the transcriptional activity of Runx-related transcription factor 2 (Runx2) is activated by ERK1/2 MAPK signaling (Franceschi et al., 2003; Xiao et al., 2002). The phosphorylation of Runx2 is essential for subsequent bone formation and osteoblast differentiation (Franceschi and Xiao, 2003; Salasynyk et al., 2004; Zhang and Liu, 2002). To clarify the underlying mechanisms, we further investigated the role of HAMSCs on ERK1/2 MAPK signaling pathway and Runx2 in H2O2-induced HBMSCs.

**MATERIALS AND METHODS**

**Chemicals and reagents**

Fetal bovine serum (FBS), α-minimum essential medium (αMEM), trypsin-EDTA, phosphate-buffered saline (PBS) and penicillin G-streptomycin sulfate were purchased from Gibco® Life Technologies. U0126 (ERK1/2 inhibitor), 2,7-dichlorodihydrofluoresceindiacetate (DCFH-DA), β-glycerophosphate, ascorbic acid, and dexamethasone were purchased from Sigma–Aldrich (USA). The goat anti-phospho-Runx2 (Ser451) was purchased from Abcam (China). The rabbit anti-phospho-p44/42 MAPK, p44/42 MAPK (ERK1/2), rabbit mAb, and Runx2 (D1L7F) rabbit mAb were purchased from Celle-Bio (China). The anti-rabbit IgG, phospho-p44/42 MAPK rabbit mAb(p-ERK1/2), p44/42 MAPK (ERK1/2) rabbit mAb, and Runx2 (D1L7F) rabbit mAb were purchased from Cell Signaling Technology (USA). The protein assay kit, RIPA buffer, and DAPI were purchased from Beyotime (China). The alkaline phosphatase (ALP) assay kit and bicinechonic acid (BCA) assay kit were purchased from the Jiangchong Corp (China). Other reagents used were of the highest commercial grade available.

**Cell culture**

The HBMSCs were obtained from the American Type Culture Collection (ATCC, USA). HAMSCs were collected from abandoned amniotic membrane samples obtained within 24 h using the pancreatin/collagenase digestion method as previously described (Zhang et al., 2011). Both cells were cultured in αMEM supplemented with 10% FBS, 100 U/L penicillin and 100 mg/L streptomycin in a humidified atmosphere of 5% CO2 at 37°C. Cells from passages 3-5 were used in this study and the culture medium was changed every 3 days.

**The co-culture system**

A Transwell co-culture system was used to investigate the effects of HAMSCs on proliferation and osteogenic differentiation in H2O2-induced HBMSCs. HBMSCs were seeded at an initial density of 5 × 10^4 cells/cm² in 6-well culture plates. Transwells were placed in other 6-well culture plates and seeded at increasing HBMSCs/HAMSCs ratios (5 × 10^4 cells/transwell, 10 × 10^4 cells/transwell and 15 × 10^4 cells/transwell). After approximately 24 h, HBMSCs were incubated with serum-free medium or 200 μM H2O2 for 12 h (Bai et al., 2004; Liu et al., 2004) to induce oxidative injury as previously described. After washing with PBS, transwells containing HAMSCs were transferred into the corresponding wells of the 6-well culture plate containing HBMSCs to create the HAMSC/HBMSCs Transwell co-culture system. HBMSCs in wells with transwells served as the treatment groups, while HBMSCs without transwells were designated as the control groups.

**Analysis of cellular proliferation**

Cellular proliferation was measured by flow cytometry at 3, 5 and 7 days and via immunofluorescence staining of 5-ethynyl-2′-deoxyuridine (EdU) at 5 days. Briefly, after starvation in serum-free medium or 200 μM H2O2 for 12 h, HBMSCs were washed with PBS. Transwells containing HAMSCs were moved into the corresponding wells of the 6-well culture plate containing HBMSCs to create the HAMSC/HBMSCs Transwell co-culture system. HBMSCs in wells with transwells served as the treatment groups, while HBMSCs without transwells were designated as the control groups.

**Assessment of ROS production**

The level of ROS induced by H2O2 in HBMSCs was measured using DCFH-DA as a fluorescent probe. After co-culturing with HAMSCs for 48 h, transwells containing HAMSCs were removed and HBMSCs were washed twice with PBS and fixed with 75% ice-cold ethanol at 4°C in the dark (Wang et al., 2013), cell cycle fractions (G0, G1, S, and G2 M phases) were determined by flow cytometry. The Cell-Light™ EdU Apollo®488 In Vitro Imaging Kit was used to investigate the EdU levels in each group at 5 days. Briefly, 100 μL EdU medium (50 μM) was added to each well and incubated for 2 h. The EdU medium was removed and HBMSCs were washed twice with PBS and fixed with 75% ethanol. After lysing with Triton X-100 (0.5%) for 15 min, staining solution was added to each well. After incubating for 30 min, EdU levels were examined by immunofluorescence staining and the area of stained EdU-positive nodules relative to the total culture surface was measured by Image-pro Plus analysis (IPP). DAPI was used to stain the cell nuclei.

**In vitro osteogenic differentiation**

After incubating in serum-free medium or 200 μM H2O2 for 12 h, HBMSCs were washed with PBS. Transwells containing HAMSCs were removed and HBMSCs were washed twice with PBS, incubated with DCFH-DA (10 mM) for 30 min at 37°C and again washed three times with PBS. Macrographs of DCFDA fluorescence were immediately recorded and the area of stained ROS-positive nodules relative to the total culture surface was measured by Image-pro Plus analysis (IPP). The intensity of 2',7'-dichlorofluorescin (DCF) fluorescence was determined by flow cytometry.
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Assessment of ALP activity and mineralized matrix formation
Fourteen days after co-culturing, Transwells containing HAMSCs were removed. HBMSCs in each group were then washed twice with PBS and lysed with Triton X-100 (0.5%) for 15 min. A bicinchoninic acid (BCA) assay kit was used to determine the protein concentration. ALP activity was performed using the alkaline phosphatase (ALP) assay kit (Jiancheng Corp, China) according to the manufacturer's instructions and estimated based on the absorbance at 405 nm as previously described (Ozeki et al., 2008; Schneider et al., 1999; Wang et al., 2012). The enzyme activity was expressed as micromoles of reaction product per minute per total protein.

HBMSCs co-cultured with HAMSCs for 21 days were stained with Alizarin red S to assess mineralized matrix deposition for bone nodule formation (Reinholz et al., 2000; Wang et al., 2012). Cells were washed twice with PBS, fixed with 75% dehydrated alcohol and then stained with 40 mM alizarin red S (pH 4.4) for 10 min at room temperature. After rinsing with PBS, nodules were visualized using an inverted microscope and the area of mineralized nodules was measured by Image-pro Plus analysis (IPP). Ten images were captured for each group, and the mean percentage was calculated.

Assessment of osteoblast (OB)–related protein
At the end of the 14-day co-culture, Transwells containing HAMSCs were removed and HBMSCs in each group were lysed in RIPA buffer containing 1 mM phenylmethane sulfonyl-fluoride according to the manufacturer’s instructions. The total protein concentration was determined using a bicinchoninic acid (BCA) assay kit. Protein lysates (20 μg) were separated by sodium dodecyl sulfate-polyacrylamide gel electrophoresis (SDS-PAGE) and then transferred onto 0.22 μm polyvinylidene difluoride membranes (Millipore, USA). After blocking, membranes were incubated overnight at 4°C with specific antibodies for the detection of p-ERK1/2(1:500), ERK1/2(1:500), and Runx2(1:1,000). After three washes with PBST (0.5% Tween 20 in PBS), the membranes were incubated with the relevant secondary antibodies (1:1,000) for 1 h at 37°C, washed and visualized with an ECL detection kit (Amersham Pharmacia, USA). The β-actin (1:500) served as an internal control.

The level of phospho-Runx2 was measured by flow cytometric analysis. Fourteen days after co-culturing, Transwells containing HAMSCs were removed. HBMSCs in each group were fixed with 1% paraformaldehyde for 10 min at 37°C and then permeabilized with Triton X-100 (0.5%) for 30 min on ice. After washing with PBS, cells were stained with anti-phospho-Runx2(Ser451) antibody (1:200) for 20 min at 37°C in the dark. The cells were incubated with the relevant secondary antibodies and analyzed on a flow cytometer after washing with PBS.

Statistical analysis
All data were expressed as mean ± S.D. from at least three independent experiments, each performed in quintuplicate. Data were analyzed using one-way analysis of variance (ANOVA). P-values < 0.05 were considered statistically significant.

RESULTS
HAMSCs promoted H2O2-induced HBMSCs proliferation
Flow cytometry and immunofluorescence staining of 5-ethyl-2'-deoxyuridine (EdU) were used to measure the proliferation of H2O2–induced HBMSCs seeded in the transwell co-culture system. Cell cycle fractions (G0, G1, S, and G2M phases) were determined by flow cytometry at 3, 5 and 7 days. At 3 and 7
days, the S-phase showed no significant difference between each groups. However, the proliferation of HBMSCs was significantly inhibited by H2O2 and the S-phase checkpoints increased slightly with the HBMSCs:HAMSCs ratio at 5 days (Fig. 1A). Further, immunofluorescence staining of EdU, a cell proliferation marker, revealed a statistically significant increase of EdU-positive HBMSCs nodules with the HBMSCs:HAMSCs ratio in co-culture groups compared with the single-culture groups at 5 days (Figs. 1B and 1C). Previous study had shown that co-culturing with HAMSCs promoted HBMSCs proliferation in the absence of hydrogen peroxide (Wang et al., 2015). Our results further demonstrated that co-culturing with HAMSCs also accelerated H2O2-induced HBMSCs proliferation.

**HAMSCs inhibited ROS generation in H2O2-induced HBMSCs**

To elucidate whether the beneficial effects of HAMSCs were linked to their antioxidant properties, the ROS generation in H2O2-induced HBMSCs was measured. Subsequently, macrographs of DCFDA fluorescence were analyzed qualitatively using fluorescence microscopy and the area of stained ROS-positive nodules relative to the total culture surface was measured by Image-pro Plus analysis (IPP). The intensity of fluorescence was determined by flow cytometry at 48 h. Scale bar, 100 μm; #P < 0.01 vs. the HBMSCs group; *P < 0.05 and **P < 0.01 compared with the group treated with H2O2 alone.

**HAMSCs promoted ALP activity and mineralized matrix formation in H2O2-induced HBMSCs**

Figure 3A shows the proportion of cultured HBMSCs staining positively for extracellular matrix in the presence or absence of H2O2 after 21 days co-culturing with and without HAMSCs. In contrast to H2O2-untreated HBMSCs, H2O2 significantly inhibited the mineralized matrix formation in other groups. When H2O2-induced HBMSCs were co-cultured with HAMSCs, higher levels of mineralization were observed after 21 days compared with the H2O2-induced single-culture groups and the percentage of mineralized nodules increased with the HBMSCs:HAMSCs ratio.
HAMSCs ratio (Fig. 3B). These observations demonstrated that HAMSCs positively regulated the mineralization of H2O2-induced HBMSCs in vitro.

ALP activity in HBMSCs co-cultured with and without HAMSCs was measured after 14 days irrespective of H2O2. Compared with untreated HBMSCs, ALP activity decreased significantly in the H2O2-induced groups and gradually increased with the HBMSCs:HAMSCs ratio (Fig. 3C), indicating the potential effect of HAMSC on osteoblastic differentiation in H2O2-induced HBMSC.

**HAMSCs activated ERK1/2 MAPK signaling and Runx2 in H2O2-induced HBMSCs**

ERK1/2, a member of the MAPK family, regulates the differentiation, mineralization and proliferation of HBMSCs (Salasznyk et al., 2004; Xia et al., 1995; Zhang and Liu, 2002). The transcriptional activity of Runx2 is essential for osteoblastic differentiation and subsequent bone formation. Therefore, we investigated the effect of HAMSCs on ERK1/2 signal and Runx2 activation in H2O2-induced HBMSCs. Figures 4A and 4B, respectively, show ERK1/2 and Runx2 phosphorylation in HBMSCs irrespective of H2O2 treatment after 14 days with and without co-culture with HAMSCs. In contrast to the HBMSCs without treatment of H2O2, H2O2 significantly inhibited the ERK1/2 and Runx2 phosphorylation in HBMSCs. However, when H2O2-induced HBMSCs were co-cultured with HAMSCs, higher levels of phosphorylation were observed compared with H2O2-induced single-culture groups and increased with the HBMSCs:HAMSCs ratio. These results suggested that HAMSCs enhance H2O2-induced inhibition of ERK1/2 and Runx2 phosphorylation, which might play a role in regulating H2O2-induced HBMSC differentiation and mineralization.

**U0126 inhibited the effect of HAMSCs on H2O2-induced HBMSCs**

Based on the results that ERK1/2 and Runx2 phosphorylation was activated in H2O2-induced HBMSCs cultured with HAMSCs, we used U0126, a highly selective inhibitor of ERK1/2, to clarify whether the effect of HAMSCs was ERK1/2-dependent. An appropriate HBMSCs:HAMSCs ratio was designed based on previous results. To eliminate the influence of U0126 on HAMSCs, H2O2-induced HBMSCs were treated with U0126 for 24 h after incubation with 200 μM H2O2 and prior to co-culture with HAMSCs. Figures 5A and 5B show that HAMSCs had little effect in the downregulation of ROS generation in the presence of U0126. As expected, U0126 inhibited mineralized matrix deposition and ALP activity in H2O2-induced HBMSCs co-cultured with HAMSCs (Figs. 6A-6C). Further, we found no significant difference in ERK1/2 and Runx2 phosphorylation between H2O2-induced HBMSCs with or without HAMSCs treatment consistent with the effects of U0126 (Figs. 6D and 6E). The results suggested that by inhibiting the ERK1/2 MAPK signaling, HAMSCs failed to reverse the oxidant injury induced by H2O2.

**DISCUSSION**

Approximately 50% to -70% of patients requiring dental implants...
exist at various degrees of bone inadequacy clinically (Karlin, 1971; Nakano and Yatani, 2007; Stoler, 1984). A growing body of evidence suggests that increased oxidative stress may be a significant pathogenic factor of bone loss, leading to a decrease in bone formation (Baek et al., 2010; Maggio et al., 2003). Previous studies have shown that H$_2$O$_2$ causes cell injury and inhibits osteogenic differentiation in osteoblastic cells (Liu et al., 2004). In this study, an H$_2$O$_2$-induced oxidative cell injury model was built using HBMSCs and 200 μM H$_2$O$_2$ management for 12 h. However, pre-treatment with H$_2$O$_2$ suppressed proliferation, decreased alkaline phosphatase activity (ALP), downregulated expression of osteogenic marker protein and inhibited mineralized matrix deposition, which is similar to a previous report (Huang et al., 2015).

HAMSCs are considered a potential resource in the field of bone engineering because of their beneficial properties (Marcus et al., 2008; Tsuji et al., 2010). A previous study has suggested that HAMSCs were capable of promoting proliferation and osteogenic differentiation in HBMSCs (Wang et al., 2014), which prompted us to hypothesize the role of HAMSCs in reversing the oxidative stress-induced oxidative cell injury. First, we found that HAMSCs promoted proliferation in H$_2$O$_2$-induced HBMSCs at 5 days, which was confirmed by flow cytometry and immunofluorescence staining of EdU, indicating that HAMSCs exerted a critical role in the prevention of H$_2$O$_2$-derived oxidative injury. ROS generation in H$_2$O$_2$-induced HBMSCs was determined to measure the anti-oxidant properties of HAMSCs. The excessive production of ROS, such as superoxides and H$_2$O$_2$ severely damages the DNA, protein and lipids. Oxidative stress not only affects cell proliferation, but also affects differentiation (Huang et al., 2015; Lippuner, 2012; Moriwaki et al., 2014; Ollivere et al., 2012). Our findings suggest that HAMSCs inhibited ROS generation in H$_2$O$_2$-induced HBMSCs. Further investigations suggested that co-culturing

![Image](image1.png)

**Fig. 4.** Expression of p-ERK1/2, ERK1/2, Runx2 and phospho-Runx2 proteins in H$_2$O$_2$-induced HBMSCs cultured with or without HAMSCs at 14 days. (A) Protein expression of p-ERK1/2 and total ERK1/2 were determined by Western blot, β-actin served as an internal control, (B) The level of phospho-Runx2 was measured by flow cytometric analysis.

![Image](image2.png)

**Fig. 5.** U0126 inhibited the effect of HAMSCs in downregulating ROS generation. (A) Macrographs of DCFDA fluorescence were detected by fluorescence microscopy and the intensity of fluorescence was determined by flow cytometry at 48 h, (B) Area of stained ROS-positive nodules relative to the total culture surface at 48 h was measured by Image-pro Plus analysis (IPP). Scale bar, 100 μm; ###P < 0.01 versus the HBMSCs group.
with HAMSCs significantly promoted ALP activity and mineralized matrix formation in H₂O₂-induced HBMSCs. Thus, we deduced that the anti-oxidant property induced a protective effect of HAMSCs. By improving osteogenic differentiation against oxidative stress, HAMSCs might represent an appropriate therapeutic alternative against bone inadequacy.

The core purpose of our present study was to investigate the potential molecular signaling pathways engaged by HAMSCs. Signaling pathways involved in oxidative stress-induced inhibition of osteogenic differentiation consist of MAPK, Akt/mTOR/4EBP1, p53 and NF-κB (Martindale and Holbrook, 2002; Zeng et al., 2014). The MAPK signaling including JNK, ERK and p38 pathways have emerged as major regulators of cellular physiology and the activation of MAPK is a crucial trigger of osteogenic differentiation (Herbert et al., 2015; Hu et al., 2015; Ki et al., 2013; Qin et al., 2015). Runx2, an early-stage osteoblastic differentiation marker, typically plays an important role in osteogenesis (Franceschi et al., 2009; Phiphilai et al., 2006). Researchers believe that Runx2 overexpression enhances HBMSCs differentiation along the OB lineages, increasing new bone formation (Hu et al., 2013; Kang et al., 2013). Our study showed that HAMSCs activated ERK1/2 MAPK signaling and Runx2 in H₂O₂-induced HBMSCs, while neither the JNK nor p38 pathways appeared to be involved (data not shown). Furthermore, by using U0126, the anti-oxidant function of HAMSCS was significantly downregulated by suppression of the ERK1/2 signaling pathway.

The present study highlights the antioxidant role of HAMSCs in promoting ALP activity, mineralized matrix deposition, ERK1/2 and Runx2 phosphorylation in H₂O₂-induced HBMSCs. (A, B) Mineralized matrix deposition was measured at 21 days by Alizarin red S staining and Image-pro Plus analysis (IPP), (C) ALP activity was measured at 14 days using ALP assay kit, (D) Protein expression of Runx2, p-ERK1/2 and ERK1/2 was determined by Western blot at 14 days, using β-actin as an internal control, (E) The level of phospho-Runx2 was measured by flow cytometric analysis at 14 days. Double bar, 12.5 mm; #P < 0.01 versus the HBMSCs group.

**ACKNOWLEDGMENTS**

This study was supported by the National Basic Research Program of China (2012CB966902), the National Natural Science Foundation of China (81271109) and a project funded by the Priority Academic Program Development of Jiangsu Higher Education Institutions (PAPD).

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