Isolation, culture and induced differentiation of rabbit mesenchymal stem cells into osteoblasts

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Abstract. Mesenchymal stem cells (MSCs) may be easily isolated from the bone marrow, and possess multi-lineage differentiation potential and various therapeutic applications. The differentiation of MSCs into osteoblasts is a complex process that is regulated by multiple internal and external factors. In the present study, the differentiation of MSCs isolated from rabbit bone marrow into osteoblasts using different osteoblast inductive media in the presence of dexamethasone, bone morphogenetic protein 2 (BMP-2), 1,25-dihydroxyvitamin D3, transforming growth factor β (TGFβ), platelet lysate and cyclooxygenase 2 (COX2), respectively. Alkaline phosphatase (ALP) activity, mineralization, collagen type (Ct) I and osteocalcin activities, and the mRNA and protein expression levels of vascular endothelial growth factor (VEGF), BMP-2 and Ct II were measured during the differentiation process in MSCs treated with different inducers. Rabbit MSCs were successfully isolated and were observed to be predominantly circular in shape after culture for 24 h. Following subculture for 5 days, the cells demonstrated a spindle shape. ALP, Ct I and osteocalcin activities were higher in cells cultured in dexamethasone, BMP-2 and TGFβ compared with the activities in control cells. Following differentiation, the dexamethasone, BMP-2 and TGFβ groups demonstrated significantly enhanced mineralization of MSCs detected by Alizarin Red S staining. The mRNA and protein expression levels of VEGF, BMP-2 and Ct II were significantly increased in the same groups compared with the levels in the control group. In conclusion, rabbit MSCs were successfully isolated from bone marrow and differentiated into osteoblasts indicated by raised ALP, Ct I and osteocalcin activities, mineralization and expression of osteogenesis-inducing genes and proteins. The present study revealed that dexamethasone, BMP-2 and TGFβ have a positive effect on cell differentiation.

Introduction

Marrow mesenchymal stem cells (MSCs) derived from the bone marrow stromal system have different and distinct lineage potential (1). Bone marrow MSCs may be easily isolated and exhibit a multipotent nature, including the ability to be greatly expanded in vitro and induced to differentiate into multiple mesenchymal cell types (2). MSCs serve as precursors for various mesoderm-type cells, including osteoblasts, chondroblasts and adipocytes (3-5). As MSCs are easily obtained and have strong osteogenic differentiation capabilities, they are widely applied in cellular therapy, tissue repair and regenerative medicine (6,7).

Differentiation of MSCs into osteoblasts is a complex process that is regulated by the expression of various transcription factors, predominantly vascular endothelial growth factor (VEGF) and bone morphogenetic protein 2 (BMP-2), and expression of osteoblast-specific proteins, including alkaline phosphatase (ALP) and collagen type (Ct) I (8-10). Dexamethasone and other factors, including transforming growth factor β (TGFβ), cyclooxygenase 2 (COX2) and platelet lysate, as pluralistic osteogenic inducers have attracted widespread attention from researchers (11). Ahmad and Shakoori (12) have illustrated that dexamethasone treatment accelerated murine MSC proliferation and induced early differentiation of osteoblasts. BMP-2 is the most widely studied and is the most potent inducer of osteoblastic differentiation (13,14). It has previously been reported that BMP-2 enhanced TGFβ3-mediated chondrogenesis of human bone marrow MSCs, which suggested that the combination of BMP-2 and TGFβ3 is superior to promote MSC osteogenic differentiation using TGFβ alone in MSCs chondrogenesis (15).

Multiple studies have demonstrated that environmental and hormonal factors as well as factors secreted by the cells themselves, serve an important role in the differentiation of MSCs into osteoblasts (16,17). Various approaches and methodologies have been utilized to study the molecular mechanism of

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MSC differentiation into osteoblasts (18-20). In the present study,MSCs were isolated from bone marrow of rabbits and the differentiation process of osteoblasts was studied using various cell culture media. Prior to differentiation, the first, second and third passages of cells were performed to observe cell morphology alterations, and the positive rate of MSC antigen presentation of cluster of differentiation CD44 and CD105 was measured for cell identification. Following various cultures, the activity of ALP by Alizarin Red S staining was determined, and Ct I and osteocalcin activities were measured using immunohistochemical staining. Reverse transcription-quantitative polymerase chain reaction (RT-qPCR) and western blotting were used to evaluate the expression of VEGF, BMP-2 and Ct II.

Materials and methods

**Experimental animals.** One male New Zealand white rabbit (age, 4 months; weight, 2.5-3.0 kg) was obtained from Hubei Provincial Center for Disease Control and Prevention (Wuhan, China). The rabbit was housed in a cage at 25±3°C in a 12 h light/dark cycle with 50% relative humidity and received dry pellets *ad libitum* with the intermittent addition of green fodder. All animal experiments were performed according to the Policies on the Use of Animals and Humans in Neuroscience Research, revised and approved by the Society for Neuroscience in 1995 (21). All protocols involving animal specimens were approved by the Ethical Committee of Tongji Hospital, Tongji Medical College, Huazhong University of Science and Technology Institution (Wuhan, China).

**Isolation and culture of rabbit MSCs.** A total of 10-15 ml femoral supracondylar bone marrow of New Zealand rabbits was extracted by bone marrow cavity puncture as previously described (12). MSCs were flushed out using Dulbecco’s modified Eagle’s medium (DMEM; incomplete) (Gibco; Thermo Fisher Scientific, Inc., Waltham, MA, USA). A single cell suspension was obtained by passing the flushed-out cells through a 22-gauge needle several times and counting the number of cells using a hemocytometer. DMEM supplemented with 10% fetal bovine serum (Invitrogen; Thermo Fisher Scientific, Inc.), glutamine 10 mmol/l, 100 U/ml penicillin and 100 μg/ml streptomycin was added to the cell suspension in 12-well culture plates. The plates were incubated at 37°C with 95% air and 5% CO₂ for 10 min at room temperature. Following centrifugation, the cells were washed with PBS three times, and the cell concentration was adjusted to 1x10⁶ cells/ml. The cells were subsequently incubated with fluorescein isothiocyanate conjugated goat-anti-rabbit immunoglobulin (Ig) G secondary antibodies (ab97050; 1:200; Abcam) for 40 min at room temperature. Cell surface antigens were detected using a flow cytometer (Cytomics FC 500; Beckman Coulter, Inc., Brea, CA, USA) and the results analyzed by CXP 2.1 software (Beckman Coulter, Inc.).

**Differentiation of induced, cultured MSCs into osteoblasts.** For osteoblast induction, second passage MSCs (inoculation concentration, 1x10⁶ cells/ml) were cultured in each well of 24-well plates and divided into six experimental groups (n=3) and a control group (n=3). The six experimental groups were created by the addition of one of the following to the complete culture medium: Dexamethasone 1x10⁻³ mol/l, dexamethasone (D4902) + 10 mmol/l sodium β-glycerophosphate (G9422) + 50 μg/l vitamin C (A4544); BMP-2 (B1814); COX2 (C0858); platelet lysate (SCM141); 1,25-dihydroxyvitamin D₃[1,25-(OH)₂D₃] (H107) (all Sigma-Aldrich; Merck KGaA); and TGFβ (14-8348-62; Thermo Fisher Scientific, Inc.). The cells of the control group were cultured in complete medium. Cells were incubated at 37°C with 95% air and 5% CO₂ and the culture medium in each group was changed every 3 days.

**ALP activity.** ALP activity staining was measured in all experimental groups and the control group after 7 days of culture. For this purpose, MSC layers were washed twice with PBS, fixed with 4% paraformaldehyde for 10 min at 4°C, rinsed with PBS and stained with nitro blue tetrazolium and 5-bromo-4-chloro-3-indolyl phosphate for 2 h in darkness at room temperature. The chromogenic reaction was stopped by washing twice with dH₂O and following drying the samples were observed using a light microscope at magnification, x100.

**Alizarin Red S staining.** Alizarin Red S staining was performed after growing the cells for 21 days in induction media. MSCs were washed twice with PBS and fixed with ice-cold 70% ethanol for 1 h at 4°C, and then rinsed with dH₂O twice. Alizarin Red S solution was added to cover the cells and incubated at room temperature for 30 min. The wells were washed four times with dH₂O and images were taken using an inverted microscope at magnification, x100.

**Immunohistochemical staining.** The cell pellet was fixed in 4% paraformaldehyde for 20 min at 4°C and washed twice with PBS. Cells were incubated in 0.5% Triton X-100 for 20 min at room temperature, washed twice with PBS, and then treated with 3% H₂O₂ for 15 min for the quenching of endogenous peroxidase activity at room temperature. The sections were washed twice using PBS for 2 min and antigen retrieval was performed with 2 mg/ml protease at 37°C for 30 min. Sections were washed with PBS for 5 min and non-specific staining was blocked with 5% goat serum (Sigma-Aldrich; Merck KGaA; 1:20 dilution with PBS containing 1% BSA) for 20 min at room temperature, followed by incubation with specific antibody to Ct I and osteocalcin. Following cell cultured induction for 7 days, Ct I was detected. Sections were incubated with a mouse monoclonal antibodies against Ct I (ab21286; 1:200 dilution).
dilution; Abcam) overnight at 4°C. The spatial localization of Ct I was visualized by incubation with mouse IgG horseradish peroxidase (HRP)-conjugated secondary antibodies (ab6728; 1:200 dilution; Abcam) for 1 h at room temperature, followed by 3,3′-diaminobenzidine tetrahydrochloride (Sigma-Aldrich; Merck KGaA) in the presence of H₂O₂ for 5-10 min at room temperature.

Following induction for 14 days, osteocalcin was detected. Sections were incubated with a mouse monoclonal antibody to osteocalcin (1:200; Abcam) overnight at 4°C. The spatial localization of osteocalcin was visualized by incubation with goat IgG HRP-conjugated secondary antibody (ab6728; 1:200 dilution; Abcam) for 1 h at room temperature, followed by 3,3′-diaminobenzidine tetrahydrochloride (Sigma, Missouri, USA) in the presence of H₂O₂ for 5-10 min at room temperature.

Following this, the sections were rinsed with PBS, counterstained with hematoxylin for 3 min at room temperature, dehydrated with graded ethanol (80% alcohol 1 min, 95% alcohol 1 min twice and dehydrated alcohol 1 min) and xylene for 1 min twice, and mounted with Entellan® (KGaA). The sections were observed by light microscopy (CX41; Olympus Corporation, Tokyo, Japan) at magnification, x200.

### Table I. Primers used for reverse transcription-quantitative polymerase chain reaction.

| Gene                          | Forward            | Reverse            |
|-------------------------------|--------------------|--------------------|
| Bone morphogenetic protein 2 | GTGAGGATTAGCAGGTCTT | CTGGATTTGAGGCAGTTT |
| Vascular endothelial growth factor | GACATCTTCCAGGAGTACCC | GAGGTTTGATCCGCATGAT |
| Collagen type II             | AACACTGCCAACGTCCAGAT | AGTGGATATGGCAGACGTC |
| GAPDH                        | ACCCACTCCTCTACCTTCG | CACCACCCCTGTCTGT |

**Figure 1. Morphology of rabbit MSCs.** (A) Primary MSCs were cultured for 1 day and circular-shaped cells of uniform size. (B) MSCs at the first passage were cultured for 5 days. (C) MSCs at the second passage were cultured for 5 days. (D) MSCs at the third passage were cultured for 5 days. MSCs were spindle-shaped (fibroblast-like cells) in first, second and third passages after cultured for 5 days. MSCs, mesenchymal stem cells. Magnification, x100.

**RNA extraction and RT-qPCR.** Total RNA was extracted from cells in each group with TRIzol reagent (Takara Biotechnology Co., Ltd., Dalian, China) and detected by an ultraviolet spectrophotometer and 1% agarose electrophoresis and visualized with ethidium bromide. For each sample, 1 µg RNA was reverse transcribed to obtain first-strand cDNA using a PrimeScript® RT reagent kit with gDNA Eraser (Takara Biotechnology Co., Ltd.) according to the manufacturer’s protocol at 42°C for 2 min,
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Expression levels of target genes BMP2, VEGF and Ct II were analyzed using RT-qPCR. Primer Premier 5.0 (PREMIER Biosoft International, Palo Alto, CA, USA) was used to design the fluorescent primers (Table I). Each reaction (20 µl total volume) contained 10 µl 2X SYBR Premix Ex Taq™ (Takara Biotechnology Co., Ltd.), 0.50 µmol/l each primer and 0.2±0.02 µg cDNA template. The following three-step RT-qPCR reaction was performed: Pre-denaturation

37°C for 15 min and 85°C for 15 sec. Expression levels of target genes BMP2, VEGF and Ct II were analyzed using RT-qPCR. Primer Premier 5.0 (PREMIER Biosoft International, Palo Alto, CA, USA) was used to design the fluorescent primers (Table I). Each reaction (20 µl total volume) contained 10 µl 2X SYBR Premix Ex Taq™ (Takara Biotechnology Co., Ltd.), 0.50 µmol/l each primer and 0.2±0.02 µg cDNA template. The following three-step RT-qPCR reaction was performed: Pre-denaturation
at 95°C for 30 sec, followed by 40 cycles of denaturation at 95°C for 5 sec, annealing at 60°C for 20 sec and elongation at 72°C for 20 sec. The transcriptional levels of genes were calculated using the $2^{-\Delta\Delta Cq}$ method (22). By genome analysis, the results demonstrated that GAPDH was the most stable gene among the three commonly used housekeeping genes GAPDH, 18s ribosomal RNA and β-actin in all cells. The threshold cycle (Ct) was determined for each reaction by using the $2^{-\Delta\Delta Cq}$ method,
which generated Ct values for each gene of interest normalized to the endogenous control gene (GAPDH). The quantification of target and reference genes was evaluated using standard curves, and the ratio between the target and reference gene represented the relative expression levels of the target gene. For each group, three samples were measured, and three technical replicates of each measurement were obtained.

**Western blot analysis.** Western blot analysis was performed to determine the expression of VEGF, BMP-2 and Ct II proteins in MSCs cultured in different cell culture media. The cells were homogenized with RIPA lysis buffer (Beijing Solarbio Science & Technology Co., Ltd., Beijing, China) and centrifuged at 12,000 x g for 20 min at 4°C. The protein concentration was determined using a bicinchoninic acid protein assay kit (Bio-swamp; Wuhan Beinglay Biological Technology Co., Ltd., Wuhan, China). Equal amounts of protein (30 µg) were separated by 10% SDS-PAGE and subsequently transferred to a polyvinylidene difluoride membrane (EMD Millipore, Billerica, MA, USA). The membranes were blocked for 2 h at room temperature with 5% skim milk in Tris-buffered saline (20 mmol/l Tris, 500 mmol/l NaCl and 0.05% Tween 20). Subsequently, the membrane was incubated with specific antibodies overnight at 4°C, including rabbit anti-BMP-2 antibody (ab14933; 1:1,000; Abcam), rabbit anti-VEGF antibody (ab32152; 1:4,000; Abcam) and human anti-Ct II antibody (ab159157; 1:2,000; Abcam). Anti-GAPDH antibody (ab181602; 1:10,000; Abcam) was selected as an internal reference. Following this, the membranes were washed with Tris-buffered saline and incubated with goat anti-rabbit secondary antibody (PAB160009; 1:10,000; Bio-swamp; Wuhan Beinglay Biological Technology Co., Ltd.) for 2 h at room temperature. Immunoreactivity was visualized by colorimetric reaction using an enhanced chemiluminescence substrate buffer (EMD Millipore). Membranes were scanned with a Gel Doz EZ imager and the bands were quantified using Quantity One 5.0 software (Bio-Rad Laboratories, Inc., Hercules, CA, USA).

**Statistical analysis.** All data were expressed as the mean ± standard deviation. Statistical differences were analyzed by one-way analysis of variance using SPSS 18.0 (SPSS, Inc., Chicago, IL, USA). Significant differences were detected by the Duncan's test. P<0.05 was considered to indicate a statistically significant difference.

**Results**

**Observation and identification of rabbit MSCs.** Rabbit MSCs were successfully isolated following the incubation of flushed-out cells from bone marrow. The MSCs were attached to the surface of the plate and were predominantly circular-shaped cells of uniform size after culture for 24 h (Fig. 1A). At the first, second and third passages, after 5 days, the cells were spindle-shaped (fibroblast-like cells), which is typical morphology of MSCs (Fig. 1B-D).

**Detection of rabbit MSC surface antigens.** Flow cytometric results demonstrated that the CD44- and CD105-positive rates of MSCs at the fourth passage were 97.07 and 96.54%, respectively (Fig. 2A and B). The results revealed that the purity of the isolated and cultured MSCs in the experiment was high,
confirming that the cells were derived from differentiated BMSCs following induction.

**ALP staining.** Following culture in osteogenic medium for 7 days, ALP staining was positive. Cells demonstrated gray-black and black granule precipitate in the cytoplasm precipitation (Fig. 3). Compared with the control group, ALP activity was markedly increased in the dexamethasone group, BMP-2 group and TGFβ group (Fig. 3A, B and D). The ALP activity in the COX2 group was higher than that in the 1,25-(OH)2VD3 group and platelet lysate group, but the COX-2 group was similar to the control group (Fig. 3E). These results indicated that treated with dexamethasone, BMP-2 or TGFβ notably increased ALP activity.

**Alizarin Red S staining.** The control cells and experimental groups with various osteoblast induction media were stained with Alizarin Red S following 21 days of differentiation. Alizarin Red S staining indicated dark red precipitated calcium deposits (Fig. 4). The dexamethasone, BMP-2 and TGFβ groups demonstrated markedly enhanced mineralization of MSCs compared with that observed in the other groups (Fig. 4A, B and D). Whereas cells in 1,25-(OH)2VD3, platelet lysate, COX-2 and control media were negative. The results demonstrated that dexamethasone, BMP-2 and TGFβ promoted mineralization of MSCs.

**Ct I and osteocalcin are increased in MSCs following culture in osteogenic media.** Following culture in osteogenic medium for 7 days, compared with the control group, Ct I and osteocalcin levels were markedly increased in the dexamethasone group, BMP-2 group and TGFβ group (Fig. 3A, B and D).
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in the dexamethasone, BMP-2, TGFβ and COX2 groups (Figs. 5 and 6). However, compared with the control group the Ct I and osteocalcin levels in the 1,25-(OH)2VD3 and platelet lysate group revealed no notable difference. This suggests that dexamethasone, BMP-2, TGFβ and COX2 may have induced the activity of Ct I and osteocalcin in MSCs.

**BMP-2, VEGF and Ct II mRNA and protein expression levels are increased in MSCs following culture in osteogenic induction media.** Compared with the control group, rabbit MSCs cultured in osteogenic medium (dexamethasone, BMP-2, TGFβ and COX2 groups) for 7, 14 and 21 days demonstrated significantly increased mRNA expression levels of BMP-2, VEGF and Ct II (P<0.05; Fig. 7). The expression level of BMP-2 was also significantly increased in the 1,25-(OH)2VD3 group compared with the level in the control group at days 7, 14 and 21 (P<0.05; Fig. 7).

The protein expression levels of BMP-2, VEGF and Ct II in rabbit MSCs are demonstrated in Figs. 8-10. In the dexamethasone, BMP-2, TGFβ and COX2 groups, the BMP-2, VEGF and Ct II protein expression levels were significantly higher than those in the control group in rabbit MSCs at 7, 14 and 21 days (P<0.05). The protein expression levels of BMP-2, VEGF and Ct II were consistent with the results of their gene expression. These results indicated that dexamethasone, BMP-2, TGFβ and COX2 induced the expression of BMP-2, VEGF and Ct II in rabbit MSCs.

**Discussion**

MSCs may be induced to differentiate into multiple cell types, such as osteoblasts (23). In the present study the differentiation of rabbit MSCs was studied and markers of differentiation were analyzed. Additionally, MSCs were differentiated in different culture media, including media containing dexamethasone, BMP-2, COX2, TGFβ, platelet lysate and 1,25-(OH)2VD3, respectively, and its effect was analyzed by measuring ALP activity, mineralization and expression of related genes and proteins. There are various factors that affect the osteogenic differentiation of MSCs (24). In the present study, common osteogenic induction factors were selected to explore their ability to induce osteoblasts, which laid a foundation for subsequent osteogenic induction.

Several methods are used for isolation of MSCs, including negative and positive selection of cells (25), cell sorting (26)...
and applications of cytotoxic materials (27); however, these methods have varying impacts on MSC proliferation and differentiation. In the present report, a simple method was utilized for the isolation of MSCs. Cells were successfully isolated and were attached to the surface of plates. Mostly circular-shaped cells of uniform size were observed after culture for 24 h and, following additional days of culture, the cells demonstrated spindle-shaped morphology.

The increase of ALP activity is an important marker of MSC differentiation into osteoblasts. ALP is highly expressed at an early stage of osteogenic differentiation to promote cell maturation and calcification, which is therefore considered as an important indicator for early osteogenesis (28). Mineralized nodules are typically observed as a marker for terminal differentiation, and MSCs may be stained for mineral deposition by Alizarin Red S (29). As indicated in the present study, the expression level of ALP was markedly higher in differentiating cells that were cultured in dexamethasone, BMP-2 and TGFβ1 media. The present study demonstrated the positive impact of dexamethasone, BMP-2 and TGFβ1 on the development of mineralized nodules by detecting the calcium influx. Previous studies have revealed that dexamethasone suppressed the proliferation and accelerated the differentiation process of the MSCs (12,30). BMP-2 has emerged as a key regulator of stem cell commitment and serves an essential role in osteogenic differentiation (31). Several studies have applied BMP-2 to MSCs for the repair of cartilage tissue (32,33). TGFβ1 is also known to promote chondrogenic differentiation (34). In the present study, immunohistochemical staining of Ct I and osteocalcin also demonstrated that the osteogenic potential of dexamethasone, BMP-2 and TGFβ1 was more profound than that of 1,25-(OH)2VD3, platelet lysate and COX2.

The expression of osteogenesis-inducing genes and proteins (BMP-2, VEGF and Ct II) in the present study were detected by RT-qPCR and western blotting. The results indicated that the mRNA and proteins expression levels of BMP-2, VEGF and Ct II were significantly increased in rabbit MSCs induced by dexamethasone, BMP-2, TGFβ1 and COX2. These results suggest that dexamethasone, BMP-2, TGFβ1 and COX2 promote the expression of osteogenesis-inducing genes and proteins during MSC chondrogenic differentiation. The osteogenic differentiation of MSCs involves a complex process orchestrated by multiple regulatory factors and proteins. BMPs are master regulators of osteoblast differentiation (35). It has been reported that increased levels of BMP-2 promote the chondrogenic effect, as indicated by increased expression levels of Ct II and expression of chondrogenic markers (36). VEGF was initially recognized as an endothelial-specific growth factor, which increased vascular permeability and angiogenesis, and it is now apparent that this cytokine regulates multiple biological functions in the endochondral ossification of mandibular condylar growth, as well as long bone formation (37). A study by Peng et al (38) reported that the addition of a specific antagonist of VEGF significantly inhibited BMP-2-induced bone formation and the associated angiogenesis indicated that endogenous VEGF activity is important for bone formation. BMP-2, VEGF and Ct II are important regulators that promote osteogenic differentiation of MSCs. Accordingly, it may be inferred from the present study that dexamethasone, TGFβ1 and COX2 may have the potential to increase the expression of osteogenesis-inducing genes and proteins to accelerate the osteogenic differentiation of MSCs.

In conclusion, rabbit MSCs were successfully isolated from bone marrow using a simple procedure and differentiated into osteoblast-like cells as indicated by raised ALP, Ct I and osteocalcin activities, mineralization and increased expression of osteogenesis-inducing genes and proteins. The present study also revealed that dexamethasone, BMP-2 and TGFβ1 have a positive effect on MSC differentiation. These results may provide a basis for the development of sequential delivery systems for multiple growth factors in bone engineering.

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Competing interests

The authors declare that they have no competing interests.

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