# Osteogenic differences in cultured rat periosteal cells under hypoxic and normal conditions

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Abstract. The aim of the present study was to investigate the osteogenic capability of rat calvarial periosteal cells in hypoxic conditions in vitro. Periosteum was obtained from the calvarial bone of Sprague-Dawley rats. Following primary tissue culture, subcultured cells were used in hypoxic or normal conditions. On days 1, 2, 3 and 4 following the cell culture, cell proliferation and mRNA and protein expression levels were evaluated. No significant difference in the cell proliferation rate was found between the normal and hypoxic condition groups. The hypoxic condition group exhibited a stronger expression of hypoxia-inducible factor (HIF) $1\alpha$ , vascular endothelial growth factor (VEGF), Runx2, alkaline phosphatase (ALP), bone sialoprotein (BSP), osteocalcin (OCN) and periostin at the mRNA level compared to that of the normal condition group. The hypoxic condition group also exhibited a stronger expression of HIF1 $\alpha$ , VEGF, bone morphogenetic protein (BMP)2, Runx2, ALP and BSP at the protein level compared to that of the normal condition group. In conclusion, periosteal cells cultured in hypoxic conditions demonstrated activated osteogenic capability in vitro.

### Introduction

Bone regeneration therapy is becoming common in regenerative medicine, and is used in the treatment of bone defects caused by periodontal disease and mandibular tumor resection. Bone marrow is often used as the typical vital material for bone regeneration; however, this requires an intricate procedure for harvesting. Recently, the periosteum has been cited as a bone supplement material that could be used as an alternative to bone

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marrow (1). There have also been reports that the periosteum has an osteogenic capability that is as high as bone marrow, which is often used for transplants in the maxillofacial area (2). A number of studies have been performed on the osteogenic capability of periosteal transplants (3,4). At present, collected periosteum is starting to be cultivated and clinically applied as a bone supplement (5-8).

However, it is not so easy to obtain the quantity of periosteum required for bone regeneration. To proliferate a sufficient quantity of cells for transplant, cell culture is known to be a good method. Furthermore, it is known that osteogenic capability is accelerated by changes in the environment, such as oxygen conditions. Miescher et al (10) as well as others (9,11) have reported that a hypoxic condition increased red blood cells and had various other effects on cells, such as the activation of glycolytic pathways and the induction of angiogenesis. Amemiya K et al (12) and Amemiya H et al (13) reported on the accelerated osteogenic capability of pulp cells and periodontal ligament cells in hypoxic conditions. However, no comparisons have been made to date regarding the osteogenic capability of periosteal cells in hypoxic and normal conditions. The purpose of this molecular biological study was to investigate the osteogenic capability of the cultured periosteal cells of rats when incubated under hypoxic conditions.

#### Materials and methods

This study was conducted in compliance with the Guidelines for the Treatment of Experimental Animals at the Tokyo Dental College (approval number 226102).

Animals and cell culture. Periosteal explants were harvested from the calvaria of 20 male Sprague-Dawley 7-week-old rats, each weighing approximately 250 g (Sankyo Labo Service, Tokyo, Japan). The skin incision was made and underlying muscular fibrous connective tissue was removed to expose the periosteum. The periosteum was stripped off mechanically using fine forceps. The obtained periosteum was mechanically cut into sections approximately 2x2 mm in size. The cut periosteum was placed with the osteogenic (cambium) side down onto the surface of 35-mm culture dishes for 30 min, then culture medium was added and the dishes were cultured

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for 4 days. For the culture, Medium 199 containing 10% fetal bovine serum and 50  $\mu$ g/ml gentamicin, supplemented with 10<sup>-8</sup> M dexametasone, 10 mM  $\beta$ -glycerophosphate and 50  $\mu$ g/ml ascorbic acid was used (13).

To set the oxygen conditions, a BL-40 M CO<sub>2</sub> incubator (JuujiField Labo; Bio Labo, Tokyo, Japan) which regulates the condition of oxygen in the air to nitro-oxygen was used. For the hypoxic condition group, cells were incubated in a humidified atmosphere at conditions of 5% O<sub>2</sub>, 5% CO<sub>2</sub> and 90% N<sub>2</sub> at 37°C. For the normal condition group, cells were incubated in a humidified atmosphere at normal conditions of 20% O<sub>2</sub>, 5% CO<sub>2</sub> and 75% N<sub>2</sub> at 37°C (15).

*Cell proliferation assay.* Subcultured cells from each group were harvested with 0.25% trypsin and 0.02% EDTA, and approximately  $3x10^3$  cells were seeded onto the 35-mm culture dishes. At 1, 2, 3 and 4 days after the culture, the cells were washed with phosphate-buffered saline (PBS) and harvested using 0.25% trypsin and 0.02% EDTA. The number of cells was counted using a Vi-CELL<sup>TM</sup> Coulter counter (Beckman Coulter, Inc., Fullerton, CA, USA).

Quantitative reverse transcription-polymerase chain reac*tion (RT-PCR).* Approximately  $3x10^4$  cells from each group were seeded in 35-mm dishes and cultured as detailed above. Total RNA was extracted from each sample using the acid guanidium thiocyanate/phenol-chloroform method as follows. The culture medium was removed and cells were rinsed twice using PBS. The cells were homogenized in 1 ml TRIzol reagent (Invitrogen, Grand Island, NY, USA) after 1, 2, 3 and 4 days of incubation. Each solution was transferred to a 1.5 ml tube containing chloroform and mixed. The tubes were centrifuged at 13,200 rpm at 4°C for 20 min, after which the supernatants were placed in 1.5 ml tubes containing 250 µl 100% isopropanol (half the amount of the TRIzol reagent) at -80°C for 1 h. Following centrifugation at 13,200 rpm at 4°C for 20 min, the supernatants were discarded and the remaining total RNA pellets were washed with 70% cold ethanol. Total RNAs were dissolved in 50  $\mu$ l RNase-free (diethylpyrocarbonate-treated) water and then reverse transcribed and amplified in 20  $\mu$ l volumes using a reverse transcription kit (QuantiTect; Qiagen, Germantown, MD, USA) containing RNA polymerase chain reaction (PCR) buffer (2 U/µl RNase inhibitor, 0.25 U/µl reverse transcriptase, 0.125  $\mu$ M oligo(dT) adaptor primer and 5 mM MgCl<sub>2</sub> in RNase-free water) (13). RT-PCR products were analyzed by quantitative real-time RT-PCR using a TaqMan gene expression assay (Applied Biosystems, Inc., Foster City, CA, USA) for the target genes: Hypoxia-inducible factor (HIF)1 $\alpha$ (Rn00577560\_m1, 72 bp), vascular endothelial growth factor (VEGF) (Rn00582935\_m1, 75 bp), alkaline phosphatase (ALP) (Rn01516028\_m1, 68 bp), bone sialoprotein (BSP) (Rn01450118\_m1,89bp), osteocalcin (OCN) (Rn01455285\_g1, 81 bp), Runx2 (Rn01512296\_m1, 116 bp). The TaqMan endogenous control (Applied Biosystems) for the target gene β-actin (Rn01768120\_m1, 63 bp) was used as an endogenous control. All PCR reactions were performed using a real-time PCR 7500 fast system (Applied Biosystems). Gene expression quantification using TaqMan gene expression assays was performed as the second step in a two-step RT-PCR. Assays

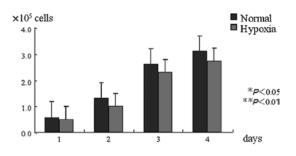
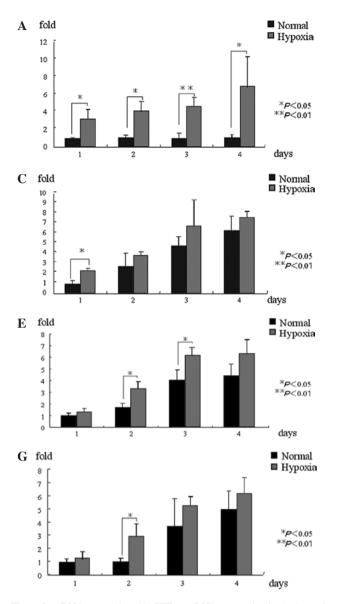


Figure 1. Cell proliferation rate. No difference was noted between the hypoxic and normal condition groups at any of the time-periods.

were performed in 20  $\mu$ l single-plex reactions containing TaqMan Fast Universal PCR Master mix, TaqMan gene expression assays, distilled water and complementary DNA according to the manufacturer's instructions. Reaction conditions consisted of 50 cycles at 95°C for 3 sec and at 62°C for 30 sec.

ALP activity. ALP activity was measured using a colormetric assay kit (Alkaline Phosphatase Opt; Roche Diagnostics Japan, Tokyo, Japan). Cultured cells (3500 cells per well) were washed with calcium- and magnesium-free PBS at each timepoint, and harvested with demineralized and distilled water for 60 sec using a sonicator (Sonifier 250D; Branson, Rochester, MI, USA) on ice. Each homogenate was centrifuged at 800 x g for 5 min and the supernatants were used for assay. One milliliter of premixed solution (1 M diethanolamine buffer, pH 9.8, with 0.5 mM MgCl<sub>2</sub> and 10 mM p-nitrophenylphosphate, kept at 37°C) was added to 10  $\mu$ l supernatant. Absorption at 405 nm for p-nitrophenol was measured using a spectrophotometer (Ultrospec 3000; Amersham Pharmacia Biotechnologies, Rochester, NY, USA). To determine the specific activity of ALP, protein concentrations in each lysate were determined using the Pierce bicinchoninic acid protein assay (Pierce, Rockford, IL, USA). A volume of 100  $\mu$ l of each cell lysate was added to 100  $\mu$ l bicinchoninic acid working reagent (kept at 37°C for 30 min). Absorbance was measured at 595 nm using a microplate reader. ALP activity was calculated according to the manufacturer's instructions.

Western blot analysis. The expression of HIF1a, bone morphogenetic protein (BMP)2, Runx2, BSP, OCN, VEGF, Glut1 and periostin in periosteal cells was assessed by western blot analysis. Blocking was performed by PVDF blocking reagent to confirm specific and non-specific bands. Periosteal cells were seeded in 60-mm cell culture dishes and were cultured for 4 days. After each culture period, the cells were washed with CMF-PBS, and resuspended in radio-immunoprecipitation assay buffer for 60 sec using a sonicator (Branson) on ice. Each homogenate was further stirred using a tube rotator (As One) for 20 min and then centrifuged at 9,300 x g for 20 min at 4°C. The protein concentration of each supernatant was measured using the Nowak method (16). Samples (30  $\mu$ g protein each) were electrophoresed in 7.5% sodium dodecyl sulfate polyacrylamide gels and were transferred to nitrocellulose membranes using standard methods (12). Primary antibodies were used at a dilution of 1:1000. These antibodies comprised rabbit polyclonal antibodies against



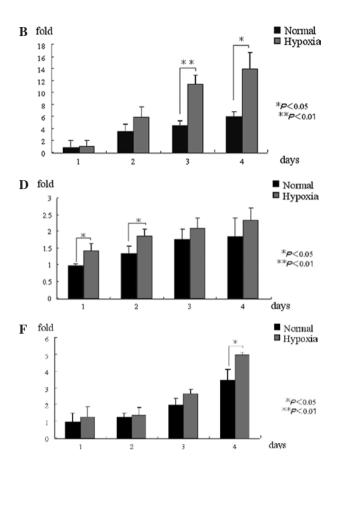


Figure 2. mRNA expression. (A) HIF1 $\alpha$  mRNA expression in the hypoxic condition group was significantly higher than that in the normal condition group on days 2 and 3, but was not significantly different at other time-periods. (C) Runx2 mRNA expression in the hypoxic condition group was significantly higher than that in the normal condition group on days 2 and 3, but was not significantly different at other time-periods. (C) Runx2 mRNA expression in the hypoxic condition group was significantly higher than that in the normal condition group on day 1, but was not significantly different at other time-periods. (D) ALP mRNA expression in the hypoxic condition group was significantly higher than that in the normal condition group on days 1 and 2, but was not significantly different at other time-periods. (E) BSP mRNA expression in the hypoxic condition group on days 2 and 3, but was not significantly higher than that in the normal condition group was significantly higher than that in the normal condition group on days 1 and 2, but was not significantly different at other time-periods. (F) OCN mRNA expression in the hypoxic condition group on days 4, but was not significantly different at other time-periods. (G) Periostin mRNA expression in the hypoxic condition group on day 4, but was not significantly different at other time-periods. (F) QCN mRNA expression in the hypoxic condition group on day 2, but was not significantly different at other time-periods. (G) Periostin mRNA expression in the hypoxic condition group was significantly higher than that in the normal condition group on day 2, but was not significantly different at other time-periods. (F) QCN mRNA expression in the hypoxic condition group on day 4, but was not significantly different at other time-periods. (G) Periostin mRNA expression in the hypoxic condition group was significantly higher than that in the normal condition group on day 2, but was not significantly different at other time-periods. \*P<0.05, \*\*P<0.01. HIF1 $\alpha$ , hypoxia-inducible factor;

HIF1 $\alpha$ , BMP2, Runx2, BSP, OCN, VEGF, Glut1 and periostin. As a secondary antibody, horseradish peroxidase (HRP) conjugated anti-rabbit IgG antibody was used. Specifically bound antibodies were detected on the film with an enhanced chemiluminescence detection system (ECL Western blot detecting system; Amersham Pharmacia Biotech, Piscataway, NJ, USA). Images were analyzed using NIH Image (Scion Corporation, Frederick, MD, USA) and each density was normalized with actin from the same sample.

*Statistical analysis*. The results were analyzed using one-way analysis of variance (ANOVA) and then compared by Scheffe's test.

### Results

*Cell proliferation rate*. There was no significant difference in the cell proliferation rate between the normal and hypoxic condition groups (Fig. 1).

mRNA expression (RT-PCR). HIF1 $\alpha$  expression was higher in the hypoxic condition group than in the normal condition group on days 1, 2, 3 and 4 (P<0.05). VEGF expression was significantly higher in the hypoxic condition group than in the normal condition group on days 3 and 4 (P<0.05). In particular, a significantly high value was indicated on day 3 (P<0.01). Runx2 expression was significantly higher in the

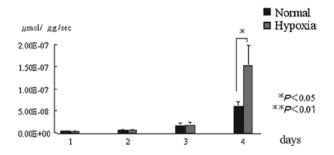


Figure 3. Alkaline phosphatase (ALP) activity. Compared to the normal condition group, the hypoxic condition group demonstrated a higher expression of ALP activity on day 4. \*P<0.05, \*\*P<0.01

hypoxic condition group than in the normal condition group on day 1 (P<0.05). ALP expression was significantly higher in the hypoxic condition group than in the normal condition group on days 1 and 2 (P<0.05). BSP expression was significantly higher in the hypoxic condition group than in the normal condition group on days 2 and 3 (P<0.05). OCN expression was significantly higher in the hypoxic condition group than in the normal condition group on day 4 (P<0.05). In particular, a significantly high value was indicated on day 3 (P<0.01). Periostin expression was significantly higher in the hypoxic condition group than in the normal condition group on day 2 (P<0.05) (Fig. 2).

ALP activity. In comparison to the normal condition group, the hypoxic condition group demonstrated a higher (P<0.05) expression of ALP activity on day 4. However, there were no significant differences at the other time-periods (Fig. 3).

*Protein expression*. HIF1 $\alpha$ , VEGF, BMP2, Runx2 and BSP were all expressed more strongly in the hypoxic condition cell group than in the normal condition group. Glut1, OCN and periostin were all expressed in both the normal and hypoxic condition groups (Fig. 4).

## Discussion

It has been reported that osteogenic capability is just as remarkably high in periosteal cells as it is in bone marrow cells, which are the material typically used in bone regeneration therapy (8). Pittenger et al reported that SH2 and SH3 are mesenchymal stem cell markers that differentiate into osteoblasts within bone marrow (17). Since then, the same markers have also been reported in the periosteum (18,19). Furthermore, Sakaguchi et al reported that there was no significant difference between periosteal and bone marrow cells in terms of the expression of CD44, CD90 and CD105, which are also mesenchymal stem cell markers (20). We observed that rat cultured bone marrow cells expressed the same markers as cultured periosteal cells using exactly the same antibodies as were used in this study (data not shown). Taken together, these results suggest that both periosteal and bone marrow cells have similar characteristics, in terms of their quality and quantity of stem cells and strong osteogenic capability.

In the maxillofacial area, the mandible has been cited as a collection site for actual periosteal transplants. The calvarial periosteum used in this study also originates from intramembranous ossification as in the mandibular bone. Jeroen *et al* (22)

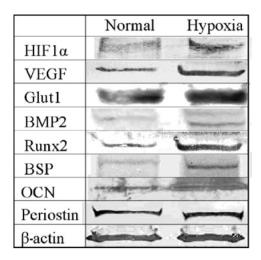


Figure 4. Protein expression. We carried out an evaluation of the normal and the hypoxic condition group on day 4 after the incubation. HIF1 $\alpha$ , VEGF, BMP2, Runx2 and BSP were all expressed more strongly in the hypoxic condition cell group than in the normal condition group. Glut1, OCN and periostin were all expressed in both the normal and hypoxic condition groups. HIF1 $\alpha$ , hypoxia-inducible factor; VEGF, vascular endothelial growth factor; BMP2, bone morphogenetic protein 2; ALP, alkaline phosphatase; BSP, bone sialoprotein; OCN, osteocalcin.

conducted comparisons of the periosteal osteogenic capability of rabbits and humans, and Krzysztof *et al* (21) conducted comparisons in chickens, dogs, mice and rats. Furthermore, Wei *et al* compared periosteum of the same size collected from the diaphyseal and epiphyseal region of the femurs of rats according to age (23,24). In these studies, there were differences in the speed of differentiation and the degree of osteogenic capability according to species, site and age, but they were mostly in agreement regarding cell kinetics. Therefore, this molecular biological study of cell kinetics using calvarial periosteal cells of rats may assist in the clinical application of human mandible periosteal cells.

Generally, it is known that the cell proliferation ratio changes according to the type of hypoxic condition that the cells are placed in (25-30). Also, the hypoxic condition affects the cell kinetics even when the same cells are used in the condition. Yoshida *et al* studied mouse embryonic fibroblasts *in vitro* with the oxygen condition set to 1, 5 and 20%, and found the number of colonies to be highest at the condition of 5% oxygen (31). Senzui *et al* reported that the cell count in rat pulp cells decreased at an oxygen condition of 5% (16). No significant difference was noted in the hypoxic condition at 5% oxygen in this study. These results shows that there is a marked difference in cell proliferation between rat periosteal cells and mouse embryonic fibroblasts under hypoxic conditions.

HIF1 $\alpha$  is a hypoxia-inducible factor, and expressed in hypoxic conditions. Hanada *et al* reported that the expression of BMP2 raised the osteogenic capability of the periosteum (32). In this study, a stronger protein expression of BMP2 was noted in the hypoxic condition group than in the normal condition group. HIF1 $\alpha$  induces hypoxia via integrin-linked kinase (ILK), Akt and the mammalian target of rapamycin (mTOR), causing BMP2 to be strongly expressed (33) and accelerating the cell differentiation.

In this study, the hypoxic condition group exhibited significantly and chronologically higher values than the normal

condition group which may suggest that periosteal cells are resistant to hypoxia. In this study, Glut1 and VEGF have been cited as being strongly expressed at either the protein or mRNA level in hypoxic conditions. Glut1, which comprises 5% of red blood cell membrane protein, activates the glycolytic pathways, and VEGF activates angiogenesis and HIF1 $\alpha$  is closely related to osteogenic capability (34-38). HIF1a binds to hypoxia response elements located in the gene promoters to regulate the transcription of VEGF and Glut1 (39). Runx2 is an initial transcription factor for the differentiation process of osteoblasts. Komori et al (40) and Otto et al (41) proved that Runx2 gene is vital to the differentiation and osteogenesis of osteoblasts since in Runx2 knockout mice, osteoblast differentiation is clearly suppressed and osteoblast formation does not take place. Kobayashi et al proved that although neither endochondral ossification nor intramembranous ossification occur in Runx2 knockout mice, since the differentiation of damaged mesenchymal cells into adipocytes and chondrocytes is possible, it is an essential factor in the initial stage of osteoblast differentiation (42). Liu et al reported that although Runx2 functions as an accelerator in the initial stage of osteoblast differentiation, it inhibits mature osteoblasts and osteocytes (43). In this study, both mRNA and protein levels of Runx2 were strongly expressed in the hypoxic incubation group. This suggests that periosteal cells, which might include pre-osteoblast stage cells, react to the hypoxic condition. ALP is known to be a marker for early osteogenesis, BSP is known to be a marker for pre-calcification, and OCN is known to be a marker for post-calcification (44). In this study, both these markers showed higher expression at both the protein and mRNA level in the hypoxic condition group. In particular, BSP was more strongly expressed on days 2 and 3 and OCN was more strongly expressed on day 4 in the hypoxic condition group. It would appear that a hypoxic condition is also effective for osteoblasts to mature to osteocytes.

Periostin, which is known to be a specifically expressed protein of the osteoblastic cells of the periodontal ligament and periosteum, was strongly expressed at the mRNA level in the hypoxic condition group in this study. Hiouchi *et al* reported that periostin existed only in the periosteum, and was expressed in the periodontal ligament induced by some kind of mechanical stress (45). Kudo *et al* (47) as well as others (46,48) reported that periostin affects the ligands  $\alpha$ -V/ $\beta$ 3 and  $\alpha$ -V/ $\beta$ 5, and is involved in the angiogenesis of malignant tumors. Another report suggests that periostin is involved in the phosphorylation of FAK and Akt in  $\alpha$ v-integrin pathways during the recovery process following a myocardial infarction (49). The fact that Akt is a factor in the upregulation of HIF1 $\alpha$  (50) suggests that there must be a significant correlation between periostin and hypoxic conditions.

In conclusion, hypoxic conditions activate the osteogenic capability of periosteal cells in rats.

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