REGULAR ARTICLE

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Isolation and multilineage differentiation of bovine bone marrow mesenchymal stem cells

Abstract The bone marrow harbors a population of mesenchymal stem cells (MSCs) that possess the potential to differentiate into bone, cartilage, and fat, and along other tissue pathways. To date, MSCs from various species have been studied. Despite the bovine experimental model being widely used in experiments in vivo and in vitro, only a limited amount of information regarding bovine MSCs is available. The aim of this study was to isolate and induce the multilineage mesenchymal differentiation of bovine MSCs, thereby initiating further research on these cells. Bovine MSCs were isolated from eight calves, and osteogenic, chondrogenic, and adipogenic differentiation was induced by using a combination of previously reported protocols for other species. The level of differentiation was evaluated by histological examination and by analyzing the expression of tissue-specific genes by a quantitative "real time" reverse transcription/polymerase chain reaction technique. Following osteoinduction, the isolated fibroblast-like cells transformed into cuboidal cells and formed alkaline-phosphatase-positive colonies; during differentiation, these colonies transformed into mineralized nodules. In addition, osteogenesis was followed by osteocalcin and collagen type I mRNA expression. Chondrogenesis was confirmed by the demonstration of collagen type II, aggrecan, and sox9 mRNA expression in the cells stimulated by transforming growth factor β1 in monolayer culture. After being cultured in an adipogenesis-inducing medium, the MSCs responded by the accumulation of lipid vacuoles and the expression of adipocyte-specific genes. We have therefore demonstrated that cells harvested from bovine bone marrow are capable of in vitro extensive multiplication and multilineage differentiation, making them a relevant and invaluable model in the field of stem cell research.

Keywords Mesenchymal stem cells · Multilineage differentiation · Bovine experimental model

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Introduction

Bone marrow mesenchymal stem cells (MSC) have potential to differentiate into cells with mesenchymal, mesodermal, neuroectodermal or endodermal characteristics (Jiang et al. 2002a, 2002b). In addition, the pluripotent character of adult stem cells present in the muscle, brain, adipose tissue, and dermis of the skin has also been confirmed (Jiang et al. 2002a, 2002b; Zuk et al. 2002; Toma et al. 2001). Pluripotent cells are an attractive promising source in the field of tissue regeneration and engineering, particularly because the use of human embryonic stem cells is limited by ethical considerations (Frankel 2000).

MSCs can be directly isolated from bone marrow aspirates (1) based on their ability to adhere to plastic when plated in a monolayer culture or (2) with micro-magnetic beads by using specific antibodies for stem cells (STRO1, CD105; positive selection) or for hematopoetic cells (CD45, GlyA; negative selection; Gronthos and Simmons 1995; Majumdar et al. 2000; Reyes et al. 2001). Under appropriate cell culture conditions and stimulation by certain exogenous or endogenous (gene transfer) bioactive factors, MSCs can differentiate into various cell lineages, such as osteocytes, adipocytes, chondrocytes, myocytes, hepatocytes, astrocytes, or neurons (Wang et al. 2003; Lagasse et al. 2000; Woodbury et al. 2000).

MSCs transplanted into bone and cartilage defects are able to differentiate into osteoblast and cartilage and can repair damaged tissue by newly synthesized bone or hyaline cartilage, respectively (Bruder et al. 1998; Wakitani et al. 1994, 2002; Liu et al. 2002). Similarly, when injected into cardiac muscle, MSCs acquire the phenotype of cardiac myoblasts (Orlic et al. 2001). These characteristics indicate that MSCs can be used as powerful tools in reconstructive medicine.

The apparent plasticity of adult stem cells in vivo may be attributed to de-differentiation, trans-differentiation, or cell–cell fusion (Wagers et al. 2004). Whereas most studies have not sufficiently fulfilled the criteria proposed as the requisite proof of de-differentiation and trans-differentiation (Wagers et al. 2002; Goodell 2003; Rodic et al. 2004), cell fusion has recently been well presented as a mechanism involved in the contribution of transplanted bone marrow cells to liver hepatocytes, cardiac myocytes, and Purkinje neurons (Alvarez-Dolado et al. 2003; Vassilopoulos et al. 2003; Weimann et al. 2003). However, differentiation by cell fusion has been reported to occur only in some types of cells, whereas in other cell types, cell-fusion-independent differentiation has been reported, making all of the presently proposed mechanisms debatable (Ianus et al. 2003; Wurmser et al. 2004).

Ever since Friedenstein et al. (1976) first observed that bone marrow contains fibroblastic cells that are capable of differentiating into bone and adipose tissue, numerous studies have advanced our understanding of the phenotype, physiology, differentiation potential, and possible clinical application of these cells. MSCs used in these studies have been sourced from various species, including humans and small experimental animals such as rat, mouse, and rabbit, and large animals such as dog, cat, pig, and horse (Kadiyala et al. 1997; Martin et al. 2002; Ringe et al. 2002; Fortier et al. 1998; Pittenger et al. 1999). However, limited information regarding bovine MSCs is available, despite the wide application of the bovine experimental model in both in vivo and in vitro experiments, particularly in research on chondrocyte metabolism and cartilage resurfacing.

In the present study, we have isolated bovine MSCs and induced chondrogenesis, adipogenesis, and osteogenesis, thereby establishing a new experimental animal model for this type of research. The use of a large animal experimental model in this type of research provides numerous advantages, particularly prior to the transfer of regeneration technology to human medicine, because large animals are biomechanically more relevant to humans than small experimental animals (Aerssens et al. 1998; Liu et al. 2002; Bosnakovski et al. 2004; Kon et al. 2000).

Materials and methods

Harvesting and isolation of bovine MSC

Bone marrow was aspirated from eight calves (2–6 months old). The marrow was drawn into syringes containing heparin (1,000 units) and kept on ice until later use. Bovine MSC were isolated by previously described methods (Worster et al. 2000a, 2000b; Mastrogiacomo et al. 2001; Holzer et al. 2002; Bosnakovski et al. 2004). Briefly, the bone marrow sample was washed twice with phosphate-buffered saline (PBS) and twice with Dulbecco's Modified Eagle Medium (DMEM; GIBCO BRL, Grand Islands, N.Y., USA). Following the determination of cell viability and number, nucleated cells ($5 \times 10^4/cm^2$) were plated in T-75 culture flasks in control medium consisting of DMEM (low glucose) with 100 U/ml penicillin G, 100 μg/ml streptomycin, 0.25 μg/ml amphotericin B, 2.4 mg/ml HEPES, 3.7 mg/ml $NaHCO_3$, and 10% fetal bovine serum (FBS; lot no. 5300C, ICN, Biomedicals, Ohio, USA). The cells were incubated at 37°C in a humidified atmosphere containing 5% CO_2. After being cultured for 4 days, non-adherent cells were removed by changing the culture medium. Following the initial 4 days, which were required for the cells to attach to the flask, the medium was changed every 2–3 days. On day 12 or 13, the cells were detached by using 0.25% trypsin in 0.1% EDTA and either used in experiments or stored by deep-freezing in 10% dimethylsulfoxide until further use.

Cell proliferation assay

Proliferation of the cells was assayed by the Hoechst 33258 fluorescent dye technique (Kim et al. 1988). Briefly, the cells, which were cultured in monolayers, were detached by trypsin, collected in a 1.5-ml centrifuge tube, and digested overnight at 60°C with papain (125 μg/ml; Sigma, St Louis, Mo., USA) in PBS containing 5 mM EDTA, 5 mM L-cysteine hydrochloride. The amount of DNA was measured in triplicate by using Hoechst 33258 dye (Sigma) in a spectrofluorometer (FP777; Jasco, Tokyo, Japan) with excitation (365 nm) and emission (440 nm).

Osteogenic differentiation

Osteogenic differentiation was induced in the cells from the primary culture and from the first passage by using combinations of previously published protocols in order to enhance the expression of an osteogenic lineage (Jaiswal et al. 1997; Pittenger et al. 1999). In the primary culture (Fig. 1), osteogenesis was initiated from the first day of culturing by using osteogenic medium (Dex), viz., control medium containing 100 nM dexamethasone, 10 mM sodium β-glycerophosphate, 0.05 mM ascorbic acid (all from Sigma). Bone marrow nucleated cells ($5 \times 10^4/cm^2$) were plated in T-60 culture dishes. The dishes were divided into two groups based on the culture medium; in one group (Cont), the cells were cultured in control medium alone, whereas in the other group, they were cultured in the osteogenic medium (Dex). The cells were cultured for 24 days with the medium being changed every

3 days. On day 24, the cells from half of the dishes in each group were exposed to 10 nM 1,25-dihydroxyvitamin D_3 (vitamin D; Biomol Research Laboratory, Plymouth Meeting, Pa., USA) added to control or osteogenic medium for three more days, thereby forming the following four experimental groups: control, control + vitamin D, Dex, and Dex + vitamin D (Fig. 1).

To analyze osteogenic gene expression during differentiation, MSCs ($3 \times 10^3/cm^2$) from the first passage were plated in 6-well plates and cultured until subconfluence (80%). Subsequently, three experimental groups, namely, the control, Dex, and control + vitamin D groups, were formed based on the culture medium used. Differentiation was closely followed during the 9 days of treatment.

During the experiment, the level of osteogenic differentiation was analyzed by the visualization of alkaline phosphatase (ALP) activity and by the demonstration of mineralized materials in the cell culture by using von Kossa staining. Total RNA was harvested on day 27 (at the completion of the differentiation protocol) from the cells in the primary culture and on days 3, 6, and 9 from the cells treated in the first passage.

Chondrogenic differentiation in monolayer culture

MSCs were plated with a density of $2–5 \times 10^3$ cells/cm^2 in 6-well plates and cultured until the cells reached subconfluence. To induce chondrogenic differentiation, the cells were treated with 1 ng/ml or 5 ng/ml transforming growth factor beta 1 (TGF-β1; R&D Systems, Minneapolis, Minn., USA) in plain medium (Worster et al. 2000a, 2000b). In the control group, the cells were cultured in control medium alone. The level of differentiation was determined by analyzing the expression of cartilage-specific genes by quantitative "real time" reverse transcription/polymerase chain reaction (qRT-PCR) on days 3 and 6 of treatment.

To analyze the effect of various concentrations of TGF-β1 (0, 1, 5, and 10 ng/ml) on bovine MSC proliferation, the cells were cultured under the culture conditions previously described for cell differentiation. Proliferation was analyzed by the measurement of DNA content during the 9 days of treatment.

Adipogenic differentiation

MSC from the first or second passages were plated at a density of $2–5 \times 10^3$ cells/cm^2 in 6-well plates and cultured in control medium until the cell culture reached confluence. At this point, adipogenic stimulation was initiated by culturing the cells in adipogenesis-inducing medium containing DMEM (high glucose, 4.5 g/l), 1 μM dexamethasone, 0.5 mM indomethacin, 10 μg/ml insulin, 100 mM 3-isobutyl-1-methylxantine (all from Sigma), 10% FBS for 3 days, and for 3 more days in adipogenesis-maintenance medium containing DMEM (high glucose, 4.5 g/l), 10 μg/ml insulin, and 10% FBS. This stimulation was twice repeated (total 18 days treatment; Janderova et al. 2003). The cells in the control group were cultured in control medium (high glucose). The level of differentiation was closely followed by means of a phase-contrast microscope and by histological examination. The expression of the adipogenic-specific genes, peroxisome proliferator-activated receptor gamma 2 (PPAR-γ2) and adipocyte fatty-acid-binding protein (aP2), was analyzed on day 18 of differentiation.

Histological staining

The deposition of minerals in the cultures was demonstrated by von Kossa staining. Cell cultures in each experimental group were rinsed twice in PBS, fixed with 3.7% paraformaldehyde for 15 min, and rinsed three times in distilled water. The cells were stained with 2% silver nitrate in a dark environment for 10 min followed by washes and exposure to light for 10 min.

In glutaraldehyde-fixed cells, ALP activity was visualized by the azo-dye method by using naphthol AS-BI phosphate and Fast blue BB salt as the substrate.

Accumulation of the lipid vacuoles was visualized by Oil red O staining as follows. In brief, 10% formalin-fixed cells were stained with 0.2% Oil red O for 2 h, thoroughly rinsed in water, and counterstained with hematoxylin.

Measurement of mRNA level by qRT-PCR method

Total RNA was isolated by using Trizol (Invitrogen, Life Technologies, Carlsbad, Calif., USA) in accordance with the manufacturer's instruction. After the samples were treated with DNase to remove possible contamination by genomic DNA, first-strand complementary DNA (cDNA) was synthesized from 2 μg total RNA by using M-MLV reverse transcriptase (Invitrogen) with oligo $(dT)_{20}$ as a primer in 20 μl reaction mixture. The amount of cDNA was measured by the qPCR method by using the Smart Cycler System (Cepheid, Calif., USA). PCR was carried
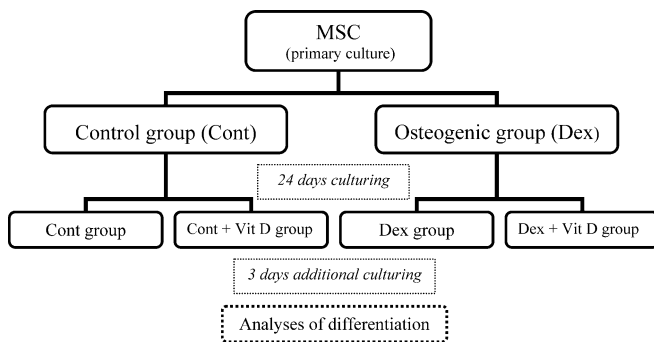
Fig. 1 Scheme of the osteogenic differentiation protocol in primary culture. Cells in the control group (*Cont*) were cultured in control medium; cells in the osteogenic group (*Dex*) were cultured in osteogenic medium; cells in the *Cont + Vit D* and *Dex + Vit D* groups were cultured in control and osteogenic medium, respectively, supplemented with 10 nM 1,25-dihydroxyvitamin D_3 (*Vit D*)

out in a 25-μl final volume containing PCR buffer, 3 mM MgCl$_2$, 0.3 mM dNTP mixture, 0.3 μM each of the relevant bovine-specific primer (Table 1), and *Taq* polymerase (1.25 units/tube; Takara Biomedicals, Otsu, Japan). Forty-five cycles of thermal cycling were carried out at 95°C for 5 s, 55°C-60°C for 15 s, and 75°C for 15 s. The amount of PCR product was estimated by measurement of the intensity of fluorescence of SYBR Green I interacting with the PCR product. The mRNA expression level of target genes was normalized by dividing with the mRNA level of glyceraldehyde-3-phosphate dehydrogenase (GAPDH; osteogenesis and chondrogenesis) or β-actin (adipogenesis). The quality of the PCR products was checked by melting curve analysis and electrophoresis. In addition, the identity of each PCR product was confirmed by sequence analysis (ABI-PRISM 310 Genetic Analyzer; PE Biosystems, Foster City, Calif., USA).

Statistical analyses

Data from a minimum of four independent experiments were evaluated and expressed as mean ± SD. An unpaired Student's *t* test was used to test significance between the two groups. The results were considered significantly different at $P<0.05$.

Results

Cell culture

MSCs were isolated from bovine bone marrow based on their characteristic property of attaching to plastic culture flasks. The isolated cells that were cultured in monolayer had a bipolar to polygonal fibroblast-like shape (Fig. 2). During monolayer culturing (up to 6 passages) by standard cell culture techniques, no remarkable loss was noted in the cell proliferation potential (data not shown).

Osteogenic differentiation

Bovine MSCs grown in the primary culture with the osteogenic medium (Dex) formed colonies, and their cellular morphology changed from a fibroblastic shape to a cuboidal form. The cell colonies were found to be ALP-positive on days 12 and 27 of differentiation (Fig. 3a, b). During the culture period, the colonies transformed into mineralized nodules that were demonstrated by von Kossa staining (Fig. 3c). On the other hand, the cells cultured in control medium did not form colonies and retained their fibroblastic shape. A 3-day additional treatment with vitamin D in cell culture had no notable influence on cell shape, ALP activity, or mineralized nodule formation. After 27 days of osteogenesis, significant differences in the expression of collagen type I mRNA were found between the two groups ($P<0.01$); the expression was 2.8 times higher in the group cultured with dexamethasone
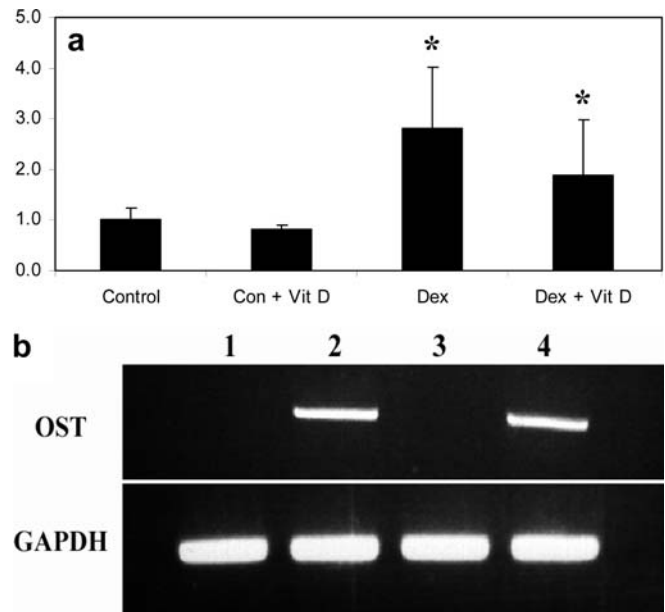
Table 1 Marker genes used in qRT-PCR

Gene	Primer nucleotide sequence	Product size (bp)	Accession number
GAPDH	Forward 5′ CCTTCATTGACCTTCACTACATGGTCTA Reverse 5′ TGGAAGATGGTGATGGCCTTTCCATTG	127	U85042
β-Actin	Forward 5′ CGCACCACTGGCATTGTCAT Reverse 5′ TCCAAGGCGACGTAGCAGAG	227	K00622
Collagen type I	Forward 5′ TGCTGGCCAACCATGCCTCT Reverse 5′ CGACATCATTGGATCCTTGCAG	120	AB008683
Collagen type II	Forward 5′ ATCCATTGCAAACCCAAAGG Reverse 5′ CCAGTTCAGGTCTCTTAGAG	147	X02420
Aggrecan	Forward 5′ CACTGTTACCGCCACTTCCC Reverse 5′ GACATCGTTCCACTCGCCCT	303	U76615
Osteocalcin	Forward 5′ TGACAGACACACCATGAGAACCC Reverse 5′ AGCTCTAGACTGGGCCGTAGAAG	320	X53699
Sox9	Forward 5′ CATGAAGATGACCGACGAG Reverse 5′ CGTCTTCTCCGTGTCGGA	118	AF278703
PPAR γ2	Forward 5′ CGCACTGGAATTAGATGACAGC Reverse 5′ CACAATCTGTCTGAGGTCTGTC	214	Y12420
aP2	Forward 5′ CTGGCATGGCCAAACCCA Reverse 5′ GTACTTGTACCAGAGCACC	186	X89244

Fig. 2 Morphology of bovine MSCs cultured in monolayer; phase-contrast microscopy. The cells exhibited a fibroblast-like shape. *Bar* 100 μm

Fig. 4 **a** Quantitative analysis of collagen type I gene expression by MSCs in primary culture on day 27 in the control (*Control*), control + vitamin D (*Con + Vit D*), Dex (*Dex*), and Dex + vitamin D (*Dex + Vit D*) groups. The expression level was normalized with the level of GAPDH mRNA. In the control group, expression of collagen type I mRNA was represented as 1. Data of four independent experiments were evaluated and expressed as mean ± SD. *$P<0.05$ when compared with the control group. **b** Electrophoresis of the PCR products from RT-PCR analysis of osteocalcin (*OST*) and GAPDH mRNA (*GAPDH*) expression in cells from primary culture (day 27) in the control (*lane 1*), control + vitamin D (*lane 2*), Dex (*lane 3*), and Dex + vitamin D (*lane 4*) groups. No osteocalcin was detected in cells untreated with vitamin D for 3 days and in the final stage of osteogenic differentiation.

than in the control group (Fig. 4a). Gene expression of osteocalcin, a typical marker of the osteoblastic phenotype, was only detected in the cells treated with vitamin D (Fig. 4b). In contrast, it was not detected in the cells treated with dexamethasone alone or cultured in control medium for 27 days.

The first-passage cells that were cultured in the osteogenic medium also transformed into cells with a cuboidal-shape; however, colonies and mineralized nodules were not formed. These cells showed 6.9 and 5.5 times higher expression level of collagen type I mRNA on days 3 and 9, respectively, when compared with the controls ($P<0.01$). The expression level of collagen type I mRNA in osteogenic medium increased with time and was 1.5 times higher on day 9 than on day 3 of differentiation (Fig. 5a). The cells in the groups treated with vitamin D showed an increased expression level of collagen type I mRNA when compared with the control group; however, this was negligible when compared with the groups treated with dexamethasone. Osteocalcin was detected after 6 days of stimulation of cells with vitamin D, and its expression had significantly ($P<0.01$) increased (by 6.3 times) after 3 days of additional treatment (Fig. 5b).

Chondrogenic differentiation

Bovine MSCs cultured in monolayer and treated with TGF-β1 exhibited a more rounded shape than the cells from the control group. By day 3, cultures treated with TGF-β1 showed a prominent increase in cellular density
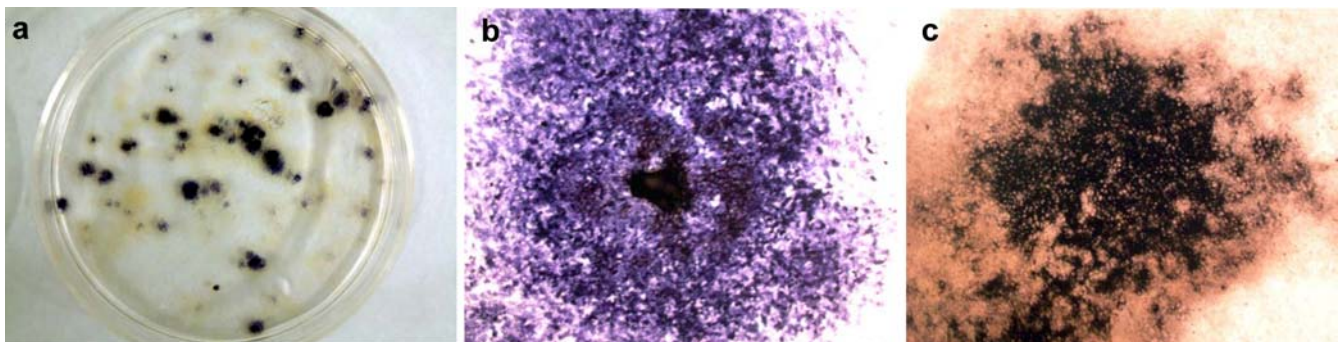
Fig. 3 Osteogenic differentiation demonstrated by histological methods. MSCs cultured in osteogenic medium formed colonies that were found to be positive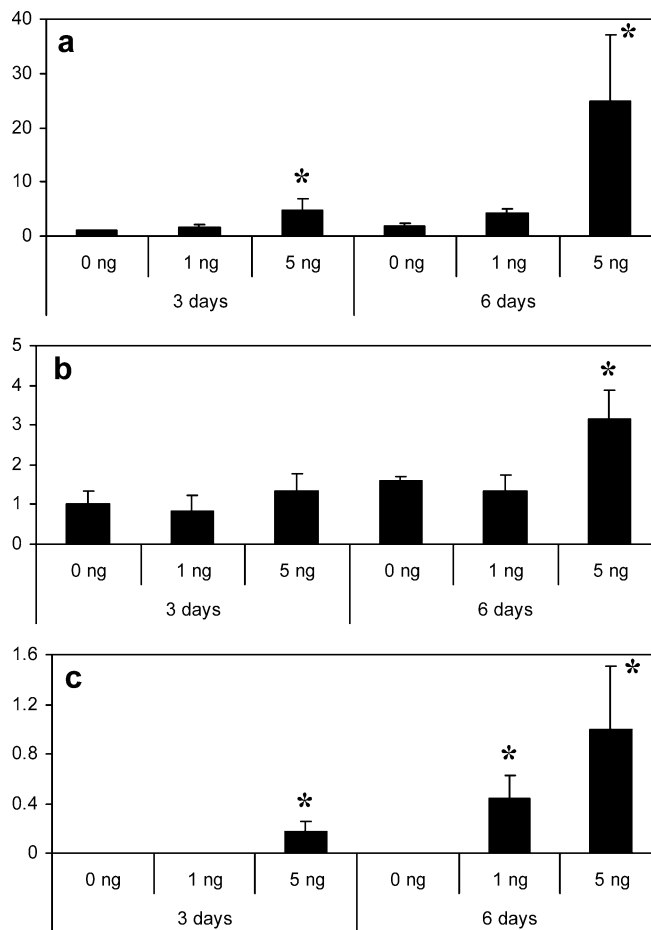 for alkaline phosphatase (ALP) staining (**a**). During culture, the colonies transformed into nodules that were found to be positive for ALP (**b**) and von Kossa staining (**c**)
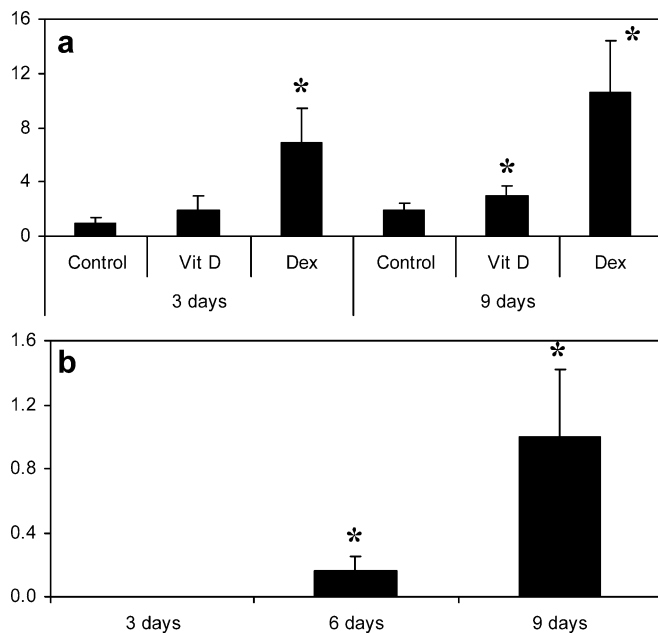
Fig. 5 a Quantitative analysis of collagen type I mRNA in the cells from the first passage in the control (*Control*), Dex (*Dex*), and control + vitamin D (*Vit D*) groups. **b** Osteocalcin gene expression by MSC from the first passage treated with 10 ng/ml vitamin D for 9 days. The expression level was normalized for the level of GAPDH mRNA. The expression of collagen type I mRNA in the control group on day 3 and that of osteocalcin in the treatment group on day 9 are represented as 1. Values are expressed as mean ± SD. *$P<0.05$ when compared with the control on the same day (**a**) and on day 3 (**b**)

Fig. 6 Quantitative analysis of chondrogenic gene expression by MSCs treated with TGF-β1. The expression level was normalized for the level of GAPDH mRNA. The expression of collagen type II (**a**) and aggrecan (**b**) in the cells in the control group on day 3 are represented as 1. The expression of sox9 (**c**) in the cells in the group treated with 5 ng/ml TGF-β1 on day 6 is represented as 1. Data of four independent experiments were evaluated and expressed as mean ± SD. *$P<0.05$ when compared with the control (0 ng/ml TGF-β1) on the same day

with multilayering; this was more pronounced on day 6. The effect on cell morphology was correlated with the concentration of TGF-β1 in the medium. The most pronounced response was observed in cultures treated with >5 ng/ml TGF-β1. The MSCs started detaching on the second day of treatment with 10 ng/ml TGF-β1 (data not shown). The detachment was more prominent during the culture period; therefore, this group was excluded from further studies. The expression of chondrocyte-specific genes that occurred following TGF-β1 treatment was found to be dose-dependent and time-dependent. Treatment with 5 ng/ml TGF-β1 increased type II collagen expression by 4.7 and 14.2 times on days 3 and 6, respectively, when compared with the cells cultured in the control groups ($P<0.05$ and $P<0.0001$; Fig. 6a). On day 6, the expression of aggrecan in the cells treated with 5 ng/ml TGF-β1 was twice that in the control cells ($P<0.01$; Fig. 6b). Sox9, an early marker of chondrogenic differentiation, was detected as early as the third day of treatment with 5 ng/ml TGF-β1. On day 6, expression increased 5.6 times compared with that on day 3 ($P<0.05$; Fig. 6c).

The effects of TGF-β1 on cell proliferation were also analyzed. After 3 days of treatment with 1 ng/ml or 5 ng/ml TGF-β1, cell proliferation significantly ($P<0.01$) increased 1.9 and 2.3 times, respectively, when compared with the controls (Table 2). During the 9 days of treatment with TGF β1, the rate of increased proliferation persisted at approximately the same level.

Table 2 Influence of TGF-β1 on the proliferation of cells during culture in monolayer for 9 days. The treatment of cell cultures with various concentrations of TGF-β1 was started at the stage of subconfluence (day 0). DNA content on day 0 is represented as 1, and the data is presented as n-fold differences (±SD).
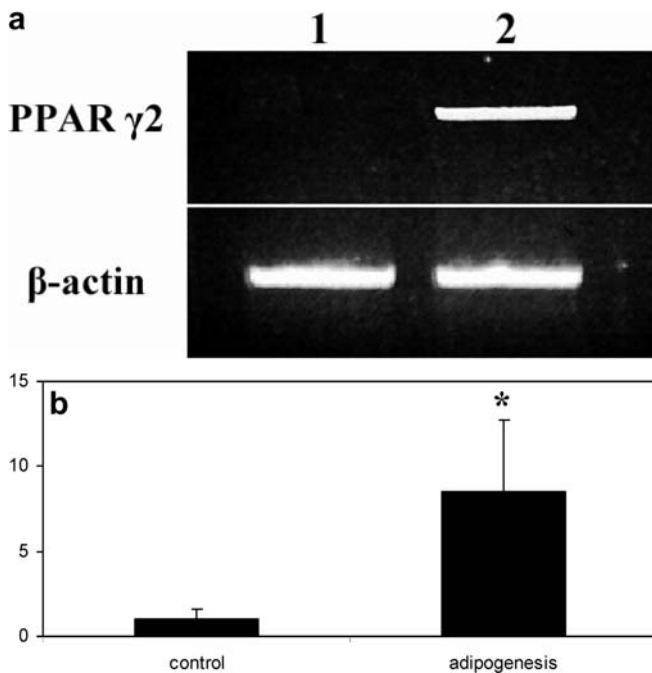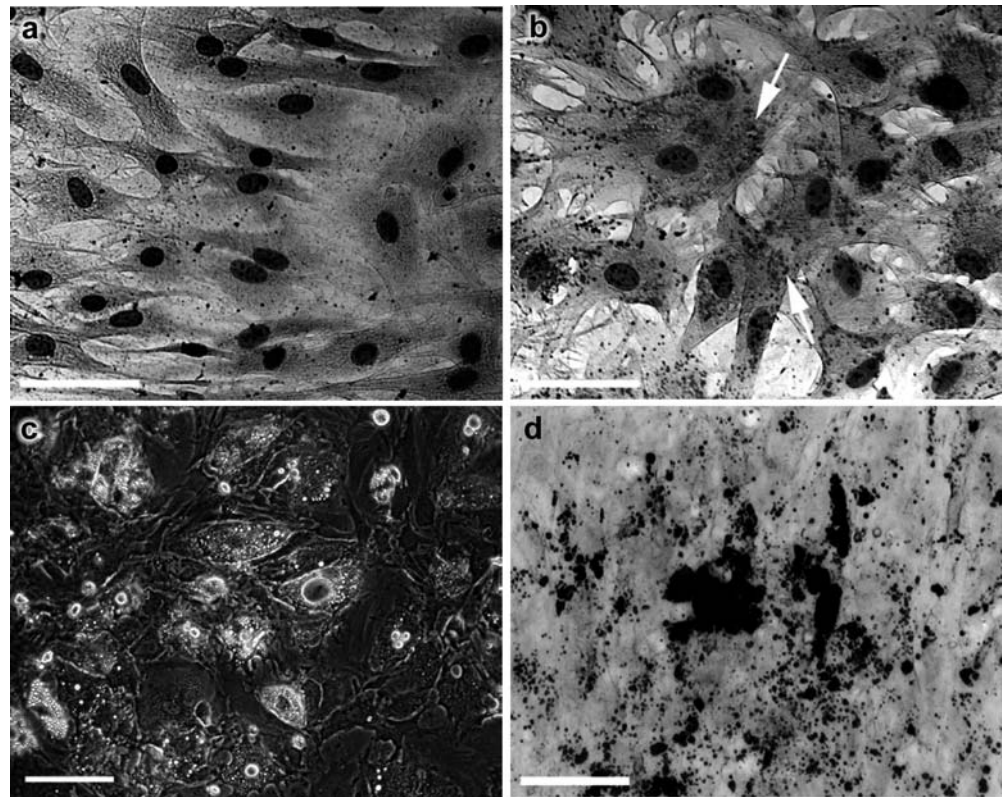
TGF β1	Day 0	Day 3	Day 6	Day 9
0 ng/ml	1±0.16	1.14±0.23	1.46±0.26	1.96±0.24
1 ng/ml	1±0.16	2.21±0.30*	3.40±0.42*	3.56±0.47*
5 ng/ml	1±0.16	2.58±0.27**	4.20±0.46**	4.24±0.48**

*$P<0.01$, **$P<0.001$ when compared with the control group (0 ng/ml TGF-β1) on the same day

Adipogenic differentiation

Adipogenic differentiation was revealed by changes observed in the cell morphology that followed the formation of lipid vacuoles and by the expression of adipose-specific genes. Following the first induction

Fig. 7 Adipogenic differentiation of MSCs; Oil red O staining of the cells in the control group on day 6 (**a**) and in the adipogenesis-induced group (**b**) after the first adipogenic treatment (3 days culturing in adipogenesis-inducing medium followed by 3 days culturing in the adipogenesis-maintaining medium). Small lipid vacuoles were detected in the cells in the treated group (*white arrows*). Morphology of the cells by phase-contrast microscopy (**c**) and Oil red O staining (**d**) after three cycles of adipogenesis induction (total 18 days) showing the increase in size of the lipid vacuoles, and the typical morphology of adipocytes. *Bar* 100 μm

treatment, small vacuoles were observed in the oval-transformed cells by phase-contrast microscopy. Oil red O staining confirmed that these were lipid droplets (Fig. 7b). The number of vacuoles and their size increased during the culture period, reaching a maximum following the third induction treatment (Fig. 7c, d). After 18 days of differentiation, 50%–90% of the cells contained lipid vacuoles of varying sizes. This estimation was carried out by counting the cells that had acquired an adipocytic phenotype in a minimum of five microscopic fields. On the other hand, cells in the control groups maintained their fibroblastic shape with a sporadic incidence of cells containing small vacuoles stained by Oil red O. On day 18 of differentiation, the expression of adipocyte-specific genes was analyzed by qRT-PCR. Expression of PPAR-γ2 was observed only in the adipogenesis-induced cells (Fig. 8a). The expression of aP2 was significantly ($P<0.001$) increased, viz., 8.4-fold (Fig. 8b), when compared with the control group.

Fig. 8 Gene expression prior to adipogenic differentiation. **a** Electrophoresis of the PCR products from RT-PCR analysis of PPAR-γ2 and β-actin mRNA expression in the cell culture for 18 days in the control (*lane 1*) and adipogenesis-induced (*lane 2*) groups. PPAR-γ2 was detected only in the treated cells. **b** Quantitative analysis of aP2 mRNA expression in the adipogenesis-induced cells and in the control cells on day 18. The expression level was normalized for the level of β-actin mRNA. The expression of aP2 in the control group is represented as 1. Data have been expressed as mean ± SD. *$P<0.05$

Discussion

The capability of fibroblast-like cells, isolated from bovine bone marrow, to undergo osteogenic, chondrogenic, and adipogenic differentiation under adequate culture conditions has been demonstrated in the present study. These fibroblast-like cells cells, referred to as MSCs, have been found to be pluripotent, as are their rodent, feline, porcine, and human counterparts.

The techniques used for the isolation and differentiation MSCs were a combination of those previously used in other species together with minor modifications aimed at achieving a better expression of lineage markers (Ringe et al. 2002; Martin et al. 2002; Heng et al. 2004).

Osteogenesis was confirmed by the presence of the ALP enzyme, a tissue non-specific (bone/liver/kidney/cartilage/fat) but well-known early osteo-differentiation marker, by the expression of osteocalcin and collagen type I mRNA, and by the mineralization of the matrix. Currently, a definitive marker for cells of the osteogenic lineage is unknown; however, the three criteria mentioned above are widely accepted as being characteristic of osteoblasts (Andrades et al. 1999; Mizuno et al. 2000; Walsh et al. 2003; De Jong et al. 2004; Chen 2004). On exposure to osteoinductive medium, bovine MSCs, like those from other species, transformed from a fibroblastic to a cuboidal shape, thereby exhibiting osteoblastic morphology. In primary culture, changes in the cell morphology were followed by cell growth into colonies and the formation of ALP-positive mineralized nodules, a characteristic of osteogenesis in rat and human MSCs (Mizuno et al. 2000; Bruder et al. 1997) but not in feline or ovine MSCs (Martin et al. 2002; Kon et al. 2000). Bovine MSCs expressed osteocalcin mRNA in response to vitamin D but not to dexamethasone treatment. The same behavior was reported in human MSCs, which when treated with dexamethasone, expressed considerably less osteocalcin than did cells that were not cultured in osteoinductive medium prior to treatment with vitamin D. This could be attributable to the glucocorticoid-induced reduction of vitamin D receptors in osteoblastic cells, leading to decreased osteocalcin expression during vitamin D treatment (Jaiswal et al. 1997). The expression of collagen type I mRNA increased in the cells treated with osteoinductive medium, a result indicating the occurrence of osteodifferentiation in the bovine MSCs.

During osteogenesis in the first passage, the bovine MSCs lost the ability to form mineralized nodules; however, the potential to express osteocalcin and collagen type I mRNA was maintained. This modification in the manner of osteogenic phenotype expression does not necessarily imply that the cells lose their potential to differentiate. Satomura et al. (2000) have shown that the formation of mineralized nodules in vitro does not predict actual osteogenic capability in vivo because clonal lines of certain murine cells do not form nodules in culture but are able to form bone when transplanted into immunocompromised mice. However, the loss in their potential for differentiation during culturing could be attributed to several factors, including species character, cell isolation, and culture techniques, and conditions in the culture medium (Kadiyala et al. 1997; Sekiya et al. 2002; Digirolamo et al. 1999).

Close cell-to-cell contact and certain chondrogenic bioactive factors are widely accepted as being two main principles involved in enhancing chondrogenic differentiation. The best method to obtain strong cell-to-cell interaction is to culture MSCs in a pellet or micromass culture system (Johnstone et al. 1998; Jones et al. 2002). The bioactive factors that stimulate chondrogenesis include TGF-β1, bone morphogenic protein, fibroblast growth factor, and insulin-like growth factor (Mastrogiacomo et al. 2001). TGF-β1 acts via high-affinity interaction with a heteromeric receptor complex comprising two structurally-related serine-threonine kinases, the type I and II receptors, and transduces its signals through Smad proteins intracellularly (Miyazono 2000). In addition to the pellet and micromass culture systems, Worster et al. (2000a) have induced chondrogenesis in equine MSCs in monolayer culture by using TGF-β1. We have reported the spontaneous chondrogenic differentiation of bovine MSCs, independent of exogenous bioactive substances, in a pellet culture system and thereby have confirmed the importance of culture conditions prior to chondrogenesis (Bosnakovski et al. 2004). In the present study, the potential of bovine MSCs to undergo chondrogenesis has been analyzed after treatment with TGF-β1 in monolayer culture. The chondrogenic differentiation of MSCs in monolayer culture apears to be dose-dependent and time-dependent in relation to TGF-β1, despite its suppressive effect on chondrogenesis in pellet culture (Bosnakovski et al. 2004). In our system, TGF-β1 induces increased collagen type II expression (14.2 times), which clearly demonstrates the chondrogenic effect of this factor. Cells cultured with no treatment with any bioactive substance do not express sox9, an early transcription factor of chondrogenesis. However, it is detectable after treatment with TGF-β1. Sox9 has been shown to regulate the rate of chondrocyte differentiation by controlling the expression of a series of chondrocyte-specific genes, including those for aggrecan and collagens type II, IX, and XI (De Crombrugghe 2000; Tchetina et al. 2003). In our previous study, the expression of chondrogenic genes was not observed in cells cultured in monolayer (Bosnakovski et al. 2004). However, with the improvement in the sensitivity of RT-PCR techniques, a basal level of gene expression of collagen type II and aggrecan was detected in MSCs cultured in medium containing FBS. A low level of gene expression of collagen type II has also been detected in untreated human and equine MSCs (Majumdar et al. 2000; Fortier et al. 1998; Worster et al. 2000a, 2000b). Moreover, murine bone marrow stromal cells express several markers that are characteristic of osteoblastic, chondrogenic, and adipogenic phenotypes at some point during the extended culture period; this suggests that individual stromal cells are inherently flexible with respect to phenotype and demonstrate the potential for plasticity (Satomura et al. 2000). The effect of TGF-β1 on chondrocyte proliferation and the synthesis of extracellular matrix is contextual and may be stimulatory or inhibitory, depending on the stage of cellular differentiation, growth condition, and other factors (Glansbeek et al. 1993).

In addition to the promotion of chondrogenesis, TGF-β1 also stimulates the proliferation of bovine MSCs in monolayer culture. This is evident even at a low concentration (1 ng/ml) after 3 days of stimulation. These results can be correlated with previously published

reports on the stimulating effect of TGF-β1 on chondro-cytes and rat MSC proliferation (Qi and Scully 1998; Andrades et al. 1999); however, they are not in agreement with studies in which human MSCs have been used (Walsh at al. 2003). This could be attributable to species difference, the use of different doses of TGF-β1, or differences in culture conditions.

Adipogenic differentiation was induced by using a combination of MIX and dexamethasone, a routinely used method to stimulate the differentiation of pre-adipocytes and stem cells. The mechanism of action of both stimulants is not completely understood; however, certain studies indicate that the effect is based on the mediation of cAMP metabolism (Gregoire et al. 1998). In our system, differentiation was apparent by the accumulation of lipid vacuoles within cells that developed, coalesced, and eventually filled the entire cell over a period of time. Increased expression of aP2 and PPAR-γ2 was detected in these cells. PPAR-γ2 is an adipocyte-specific gene belonging to the PPARs family, and together with C/EBP, is one of the transcription factors that are induced early during the differentiation of adipocytes (Rosen et al. 1999). AP2 is also an adipose-tissue-specific gene and is synthesized in the terminal phase of differentiation (Gregoire et al. 1998). In addition, we have confirmed an increased expression of lipoprotein lipase in treated cells (data not shown); however, this marker is not adipose-tissue-specific and is also synthesized by macro-phages and other cells of mesenchymal origin (Tengku-Muhammad et al. 1996). During the screening of adipo-genesis, aP2 expression has been detected in uncommitted cells. This may be attributable to the minor contamination of cell cultures with pre-adipocytes and adipocytes or to spontaneous adipogenesis in cells in a state of growth arrest or may be characteristic of the cells of the species. The method of isolation of MSCs used in this study results in a cell culture composed of a heterogenic cell population. Cell clones from this type of culture are reported to grow into colonies with a wide range of sizes and morphologies at various proliferation rates and exhibit a potential for multilineage differentiation. Cells that spontaneously dif-ferentiate into osteocytes, adipocytes, or chondrocytes have also been detected in some colonies (Bianco et al. 2001).

A limitation of the present study is the use of an in vitro model that contains a heterogeneous population of cells. However, this well-documented system has been pre-viously used for the confirmation of the presence of MSCs in cultures of bone marrow stromal cells from various species by inducing multilineage differentiation at the single cell level.

In conclusion, we have demonstrated that bovine MSCs, similar to corresponding cells from other species, exhibit high proliferation rates and multilineage differentiation potential. In addition, they can be easily harvested and isolated, thereby making them attractive tools for further studies into development and physiology and for clinical application.

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