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CHONDROGENIC DIFFERENTIATION OF INDUCED PLURIPOTENT STEM CELLS FROM OSTEOARTHRITIC CHONDROCYTES IN ALGINATE MATRIX

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Abstract

Induced pluripotent stem cells (iPSCs) have the potential to revolutionise cell therapy; however, it remains unclear whether iPSCs can be generated from human osteoarthritic chondrocytes (OCs) and subsequently induced to differentiate into chondrocytes. In the present study, we investigated the differentiation potential of OCs into iPSCs using defined transcription factors and explored the possibility of using these OC-derived iPSCs for chondrogenesis. Our study demonstrates that iPSCs can be generated from OCs and that these iPSCs are indistinguishable from human embryonic stem cells (hESCs). To promote chondrogenic differentiation, we used lentivirus to transduce iPSCs seeded in alginate matrix with transforming growth factor- β 1 (TGF- β 1) and then *in vitro* co-cultured these iPSCs with chondrocytes. Gene expression analysis showed that this combinational strategy promotes the differentiation of the established iPSCs into chondrocytes in alginate matrix. Increased expression of cartilage-related genes, including collagen II, aggrecan, and cartilage oligomeric matrix protein (COMP), and decreased gene expression of the degenerative cartilage marker, vascular endothelial growth factor (VEGF), were observed. The histological results revealed a dense sulphated extracellular matrix in the co-culture of TGF- β 1-transfected iPSCs with chondrocytes in alginate matrix. Additionally, *in vivo* chondroinductive activity was also evaluated. Histological examination revealed that more new cartilage was formed in the co-culture of TGF β 1-transfected iPSCs with chondrocytes in alginate matrix. Taken together, our data indicate that iPSCs can be generated from OCs by defined factors and the combinational strategy results in significantly improved chondrogenesis of OC-derived iPSCs. This work adds to our understanding of potential solutions to osteoarthritic cell replacement problem.

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Keywords

Chondrocytes; chondrogenesis; differentiation; co-culture

Introduction

Articular cartilage (AC) is an avascular tissue composed of chondrocytes, responsible for abundant matrix synthesis and maintenance (Merceron et al., 2011). When damaged due to traumatic or pathological conditions, AC does not heal spontaneously under physiological circumstances that result in Osteoarthritis (OA). OA is the most common musculoskeletal disease in the elderly, and could become the fourth leading cause of disability by the year 2020 (Reginster, 2002; Woolf and Pfleger, 2003; Koelling et al., 2009). Current treatment strategies are restricted to short-term symptomatic relief by pharmaceutical interventions and surgical procedures (Lohmander and Roos, 2007). Thus, treatment of cartilaginous damage poses a significant clinical challenge. Recent advances in regenerative medicine suggest that using tissue engineering-based constructs to enhance OA-cartilage repair by mobilising chondrogenic cells is a promising approach for the restoration of AC structure and function (Dehne et al., 2009; Hattori and Ohgushi, 2009; Ahmed et al., 2011).

A number of strategies have been used to identify potential cell sources for AC regeneration. It is widely agreed that tissue regeneration from autologous cells can be achieved by taking advantage of the natural course of embryonic development (DeLise et al., 2000; Tuan, 2004; Ingber et al., 2006; Goepfert et al., 2010). Therefore, autologous stem cells would be an ideal choice (Yoshida and Yamanaka, 2010). Stem cells are mainly classified into two kinds of cells: somatic stem cells and pluripotent stem cells. Somatic stem cells reside in various organs and show multipotency to regenerate damaged tissues, but have limited *in vitro* proliferative potential (Yoshida and Yamanaka, 2010). Therefore, pluripotent stem cells may be a better source for tissue regeneration. hESCs, the first pluripotent cells isolated from embryos, have high proliferation potential and can generate differentiated progeny of all three embryonic germ layers. However, the derivation of hESCs from early embryos raises technical and ethical limitations for their use in research and the clinic (Gaspar-Maia et al., 2011). Engineered stem cells, known as iPSCs, generated from somatic cells by transduction of defined reprogramming transcription factors, typically OCT4, SOX2, KLF4, and c-MYC, open a new avenue to avoid the controversy of using hESCs (Takahashi and Yamanaka, 2006; Takahashi et al., 2007). Because iPSCs have high proliferative and differentiation capabilities similar to those of hESCs, they hold great potential for regenerative medicine (Itoh et al., 2011).

Recent evidence indicates that OC-secreted soluble morphogens may induce chondrogenic differentiation of human mesenchymal stem cells (MSCs) in a 3-dimensional (3D) environment while preventing their hypertrophic differentiation (Aung et al., 2011). Moreover, human OCs responded similarly non-diseased passaged dedifferentiated chondrocytes by the strategy of co-culture (Ahmed et al., 2010). In addition, a recent report suggested that bone morphogenetic protein 4-expressing dedifferentiated chondrocytes recovered a chondrocytic phenotype *in vitro* and that some of the dedifferentiated

chondrocytes in the defect sites could undergo redifferentiation and form matrix that displayed positive toluidine blue staining for glycosaminoglycans after transplantation into the joint (Lin et al., 2008). Based on above findings, we believe that OCs are likely to serve as a cell source for the generation of iPSCs to be used in reconstruction of damaged AC. In the present study, we investigate whether iPSCs can be generated from OCs by the expression of defined transcription factors and whether these OC-derived iPSCs can be induced to differentiate into chondrocytes through a combination of lentivirus-mediated TGF- β 1 transduction and co-culture of AC in an alginate scaffold.

Materials and Methods

Tissue sources and preparation

Normal and osteoarthritic cartilage tissue fragments (Ruijin Hospital, Shanghai, China) were collected according to an Institutional Review Board approved protocol. Osteoarthritic cartilage samples, without signs of rheumatoid involvement, were obtained from the knee joints of a patient diagnosed according to the criteria of the American College of Rheumatology (Altman et al., 1990) with OA after total knee replacement. Control normal cartilage samples were obtained from the knee joints of an amputated patient with severe injury. Tissue samples were fixed in 4 % paraformaldehyde, decalcified with 10 % ethylenediaminetetraacetic acid (EDTA, pH 7.4, Sigma-Aldrich, St. Louis, MO, USA), and embedded in paraffin. Paraffin sections were stained with haematoxylin/eosin (H&E) and toluidine blue.

Chondrocyte isolation and culture

Chondrocytes from osteoarthritic and healthy cartilage were harvested following digestion for 6 h at 37 °C with collagenase I (152 U/mL; Invitrogen, Carlsbad, CA, USA), collagenase II (280 U/mL; Invitrogen), and dispase (15 U/mL; Invitrogen) and underwent subsequent filtration through a 40 μ m mesh sieve (BD Falcon, Franklin Lakes, NJ, USA) (Koelling et al., 2009). OCs were cultured in Dulbecco's modified Eagle's medium/Ham's F-12 (DMEM/F-12, Invitrogen) medium supplemented with 10 % foetal bovine serum (FBS, Invitrogen). To maintain the chondrogenic phenotype, normal chondrocytes were cultured in chondrogenic medium containing high-glucose DMEM, 20 % FBS, 100 nM dexamethasone (Sigma-Aldrich), 50 μ g/mL ascorbate-2-phosphate (Sigma-Aldrich), ITS (insulin, transferrin, and selenous acid, BD), and 1 % penicillin/streptomycin (Invitrogen). Cells were seeded on coverslips, fixed in 4 % paraformaldehyde and stained with toluidine blue.

Lentiviral production

Lentiviral vectors were constructed and packaged, and virus titers were measured according to a previous report (Honda et al., 2010). Human OCT4, SOX2, KLF4, c-MYC and TGF- β 1 cDNAs (Shanghai Biotechnology Co., Ltd, Shanghai, China) were inserted into the pENTR/D-TOPO entry vector plasmid (Invitrogen). The cDNAs in pENTR/D-TOPO were then transferred to the pCSII-EF-MCS-IRES2-Venus lentiviral vector plasmid using Gateway LR clonase (Invitrogen). Lentiviral vectors pseudotyped with the vesicular stomatitis virus G glycoprotein (VSV-G) were produced by transient transfection of three plasmids into 293T cells: the packaging plasmid (pCAG-HIVgp), the VSV-G glycoprotein-

and Rev-expressing plasmid (pCMV-VSV-G-RSV-Rev), and the lentiviral vector plasmid. The culture supernatant was concentrated using ultracentrifugation, and the viral pellet was resuspended in Hanks' balanced salt solution. Vector titers were determined by infection of 293T cells with serial dilutions of the viral concentrates supplemented with 1 µg/mL Polybrene (Sigma-Aldrich) and followed by fluorescence-activated cell sorting analysis of the transduced cells expressing EGFP (human OCT4, SOX2, KLF4 and c-MYC). Titers were expressed as 293T-transducing units per mL.

iPSC induction and culture

Approximately 3×10^5 OCs were transduced for 24 h with a lentiviral cocktail carrying GFP and reprogramming factors (OCT4, SOX2, KLF4, and c-MYC), and cultured for 6 days in DMEM (Invitrogen) containing 10 % FBS. Then, the cells were passaged onto a feeder layer of human amniotic epithelial cells (HAmEpiC) and cultured for 24 h in DMEM containing 10 % FBS. One day later, the medium was changed using an ESCs-supportive medium with high glucose DMEM, 15 % FBS, 4 ng/mL basic fibroblast growth factor (R&D, Minneapolis, MN, USA), 2mM L-Gin (Sigma-Aldrich), 1×10^{-4} M nonessential amino acids (Sigma-Aldrich), 1×10^{-4} M 2-mercaptoethanol (Sigma-Aldrich) and 1 % penicillin/streptomycin (Invitrogen). After three weeks, the human iPSC colonies were picked, dissociated with 0.05 % trypsin-EDTA (Sigma-Aldrich) digestion, plated into new culture dishes and identified. All ESC-like colonies were picked by hand and clonally expanded.

Differentiation of iPSCs *in vitro*

In vitro differentiation of the human iPSCs was performed using the standard embryoid body (EB) differentiation method (Li et al., 2009). Human iPSCs were cultured in culture dishes, and the cells were washed with 1× phosphate-buffered saline (PBS) without Ca or Mg (Invitrogen) before being dissociated with 0.05 % trypsin-EDTA. The cell suspension was pipetted and spun, and then re-suspended in ESC-supportive medium. The cell suspension was added back into the old dishes and incubated for 30 min to reattach HAmEpiC in the dishes. After that, the cells were collected from the dishes again and seeded onto the ultra-low attachment 100 mm dish (2×10^6 /dish) (Corning, Corning, NY, USA) in DMEM supplemented with 10 % FBS to form EBs. Subsequently, the EBs were plated onto Matrigel-coated slides (BD Falcon), and 3-7 days later, the cells were fixed for immunocytochemical analysis.

Directed differentiation of human iPSCs into specific cell lineage *in vitro*

For 2-3 days prior to treatment, the EBs were maintained in the three different differentiation media on ultra-low plates. Following this precondition step, the EBs were transferred to gelatine-coated 6-well plates, cultured in differentiation media for 14 days with the media changed every other day and subsequently processed for gene expression analysis and histology. The 6-well plates (Corning) were coated for at least 30 min at room temperature with 0.1 % gelatine (Sigma-Aldrich) in dH₂O (1 mL per well). The differentiation protocol was performed according to a previous report (Miki et al., 2005). Standard culture media was DMEM supplemented with 10 % FBS, 2 mM L-glutamine (Sigma-Aldrich), 1 % nonessential amino acid (Sigma-Aldrich), 55 µM 2-mercaptoethanol (Sigma-Aldrich), 1 mM sodium pyruvate (Sigma-Aldrich), and 1 % antibiotic-antimycotic (Invitrogen). For

pancreatic differentiation (endoderm), we used the standard medium supplemented with nicotinamide (10 mM); for cardiomyocyte differentiation (mesoderm), we used the standard media supplemented with 1 mM ascorbic acid 2-phosphate (Sigma-Aldrich); and for neural differentiation (ectoderm), we used the standard media supplemented with 5×10^{-5} M all-trans retinoic acid (Sigma-Aldrich) and fibroblast growth factor-4 (R&D) at 10 ng/mL.

Cytochemistry and immunofluorescence assay

Alkaline phosphatase staining was performed according to the manufacturer's instructions using the Alkaline Phosphatase Detection Kit (Millipore, Billerica, MA, USA). For the immunofluorescence assay (Miki et al., 2005), undifferentiated iPSCs and their differentiated derivatives were fixed in cold acetone for 30 min at 4 °C, then cells were rinsed with PBS three times and incubated in protein blocking agent for 20 min. Samples were incubated with primary antibodies (Table 1) overnight at 4 °C, then rinsed with PBS three times and incubated with Cy3-conjugated secondary antibodies (R&D) for 2 h at room temperature. In some experiments, cells were rinsed with PBS and mounted in aqueous mounting medium containing the nuclear counterstain DAPI (4,6 diamidino-2-phenylindole; Sigma-Aldrich). Images were captured using an Axio Observer A1 microscope (Carl Zeiss MicroImaging, Goettingen, Germany).

Teratoma formation

Human iPSCs were injected intramuscularly into nonobese diabetic/severe combined immune deficient (NOD/SCID) mice (4×10^6 cells per site). After 4-6 weeks, all mice ($n = 3$) developed teratomas, which were removed and processed for H&E staining. All animal experiments were conducted in accordance with the Guide for the Care and Use of Animals for Research Purposes and were approved by the Shanghai Jiaotong University School of Medicine Animal Care Committee.

Transfection of iPSCs with lentivirus-TGF- β 1

Human iPSCs were infected with lentivirus carrying RFP and TGF- β 1 in the presence of polybrene (8 μ g/mL) and plated onto a feeder layer of HamEpiC for 5 days, after which red-fluorescing human iPSCs were picked and transferred onto gelatine-coated 6-well plates, where they were cultured in DMEM medium containing 20 % FBS. The observed transfection rate was about 70 % for iPSCs.

Co-culture of transfected iPSCs with chondrocytes in alginate matrix

The transfected iPSCs (iPSCs/TGF- β 1) were released from the culture substratum and resuspended at a concentration of 0.417×10^6 cells/mL. Then iPSCs/TGF- β 1 were seeded onto the alginate matrix (Invitrogen) by pipetting the iPSCs/TGF- β 1 suspension (300 μ L) onto the matrix. iPSCs/TGF β 1/alginate matrix (iPSCs/TGF β 1/Alginate) constructs were incubated for an additional 20 min to allow cell attachment *in vitro*, and then the composites were seeded into a 24-well culture plate and cultured for 12 h. Normal chondrocytes were digested, counted and plated in a 24-well culture plate at a density of 1.25×10^5 cells and cultured in chondrogenic medium for 12 h.

After the above procedure was performed, the Transwell insert (0.4 μm pore size, Corning) containing the iPSCs/TGF- β 1/Alginate construct was fitted into the well containing chondrocytes, which was used as the experimental group (iPSCs/TGF- β 1/ Alginate/ chondrocytes). The co-culture was grown in chondrogenic medium for 14 days.

Additionally, the following parallel groups were cultured in chondrogenic media for 14 days: non-transfected iPSCs/alginate matrix group (iPSCs/Alginate), non-transfected iPSCs/ alginate matrix/chondrocytes group (iPSCs/Alginate/ chondrocytes) and transfected iPSCs/ alginate matrix group (iPSCs/ TGF- β 1/Alginate) according the same procedures (a schematic is available from the author upon request).

Western blotting for TGF- β 1

To confirm the presence of TGF- β 1 in lentivirus-mediated TGF- β 1 transduced iPSCs, western blotting was performed using a polyclonal rabbit anti-TGF- β 1 antibody (ab53169, Abcam, Cambridge, UK). Protein was extracted from the non-transfected and transfected iPSCs, and the protein concentration of the extracts was assessed using a Bradford assay kit. The optical density at 595 nm was measured using a spectrophotometer. The protein concentration was a standard curve obtained from a serial dilution of bovine serum albumin. The cell extracts, normalised for total protein content, were resolved using polyacrylamide gel electrophoresis and electrophoretically transferred to a supported nitrocellulose membrane. Afterwards, the membranes were blocked in 5 % non-fat milk in Tris-buffered saline with 0.1 % Tween (TBST) for 3 h and incubated with the primary antibody (1:500) at 4 °C overnight. The membranes were then washed with TBST three times for 10 min each, followed by the addition of anti-rabbit IgG (1:5000) and enhanced chemiluminescence (ECL) visualisation of the bands.

Histological analysis of iPSCs-chondrogenic differentiation

All samples were cultured for 14 days and fixed in 4 % paraformaldehyde (pH 7.4, Sigma-Aldrich) overnight at room temperature, and sections were stained with toluidine blue.

Analysis of mRNA expression using reverse transcription-polymerase chain reaction (RT-PCR) and quantitative real-time PCR

Total RNA was isolated using TRIzol (Invitrogen). RNA concentrations were determined by measuring the absorbance at 260 nm with a spectrophotometer. Conventional RT-PCR was performed using the Superscript One-Step RT-PCR system (Invitrogen). Quantitative real-time PCR was carried out using SYBR green qPCR Supermix UDG (Invitrogen) and analysed with the 7300 real-time PCR system (Applied Biosystems, Foster City, CA, USA). Amplification of specific genes was performed using the primers shown in Table 2.

Ectopic cartilage formation

Five-week-old BALB/c nude mice (Shanghai Jiaotong University School of Medicine) were anaesthetised using Sumianxin (2 mL/kg, Agricultural University, Changchun, China) administered intraperitoneally. The following groups were cultured in chondrogenic media for 14 days *in vitro*: iPSCs/Alginate group, iPSCs/Alginate/chondrocytes group, iPSCs/ TGF- β 1/Alginate group and iPSCs/TGF- β 1/Alginate/ chondrocytes group according to the

above procedure. Then, four groups were implanted subcutaneously into the dorsal region of the mice. All animal experiments were performed in accordance with the institutional animal guidelines. Specimens from each group were harvested 6 weeks after implantation. Explants harvested from the mice were fixed, dehydrated, and embedded in paraffin. Sections were subjected to immunohistochemical analysis for collagen type II (diluted at 1:100, 53047, Abcam).

Statistical analysis

Results are reported as the mean \pm SD. Values with $P < 0.05$ were considered statistically significant.

Results

Histological analysis of normal and osteoarthritic cartilage-derived cells

Healthy AC exhibited typically four intact cell layers including a superficial, transitional, deep radial, and deep calcified zone (Fig. 1Aa), and uniformly-distributed proteoglycan as the main extracellular matrix component (Fig. 1Ab). Osteoarthritic cartilage showed disappearance of cell organelles and the existence of a cell outline in some cells, and some necrotic cells (Fig. 1Ba) as well as a loss of proteoglycan (Fig. 1Bb). Chondrocytes isolated from healthy AC exhibited the typical round, polygonal shape (Fig. 1Ca) and had an abundance of proteoglycan, the main extracellular matrix component of chondrocytes (Fig. 1Cb). Most OCs had an elongated shape (Fig. 1Da) and exhibited a loss of proteoglycan (Fig. 1Db).

Characterisation of reprogrammed human OCs

Seven days after transduction with the four reprogramming factors, colonies with an ESC-like morphology were visible. After two weeks, colonies with an ESC-like morphology were large and round, with clear boundaries. To confirm that reprogramming of the patient OCs had occurred and that the putative iPSCs were pluripotent, we assayed the expression of undifferentiated ESC markers by RT-PCR and immunocytochemistry. RT-PCR results revealed that two of the putative human iPSCs expressed many predicted markers of undifferentiated hESCs, including OCT-4, SOX-2, REX-1 and NANOG, which were not expressed in the other putative iPSCs (Fig. 2A). Two of the three colonies exhibited strong alkaline phosphatase activity and the expressed ESC-specific surface markers including NANOG, SSEA-1, SSEA-4, TRA1-60, and TRA1-81 (Fig. 2B, the image shown is of the C1 colony. An image of the C2 colony is available from the author upon request). Together, these results indicate that human iPSCs expressed pluripotent markers and shared a similar gene expression profile with hESCs.

Multipotency of reprogrammed human OCs

Next, we evaluated whether the human iPSCs could differentiate into cell types of all three germ layers *in vitro* through EB-mediated differentiation. The iPSCs from the osteoarthritic patient formed ball-shaped EBs with high efficiency (Fig. 3Aa) and then readily differentiated into cells with various morphologies. Immunocytochemical analyses of the EBs confirmed that human iPSCs could differentiate into cells from all three germ layers, as

evidenced by the expression of α -foetoprotein (AFP, endoderm marker), nestin (ectoderm marker), and desmin (mesoderm marker) (Fig. 3Ab,c,d).

To induce lineage-directed differentiation of human iPSCs *in vitro*, previously reported protocols were followed for endoderm, mesoderm and ectoderm differentiation, and the associated differentiation markers were tested by conventional RT-PCR (Miki et al., 2005) (Fig. 3B). Human iPSCs were induced to differentiate into a pancreatic (endodermal) lineage, and the differentiated cells expressed PDX-1, PAX-6, NKX2.2, and insulin. Human iPSCs were induced to differentiate into a cardiac (mesodermal) lineage, and the differentiated cells expressed GATA-4, NKX2.5, MLC-2A, and MLC-2V. Finally, human iPSCs were induced to differentiate into a neural (ectodermal) lineage, and the differentiated cells expressed neural-related markers, such as NSE, NF-M, MBP, GAD, and nestin. These data demonstrate that human iPSCs can be induced directly to differentiate into cell types of all three germ layers.

Histological observation (Fig. 3C, image shown is of the C1 colony; an image of the C2 colony is available from the author upon request) showed that these teratomas possessed tissues of the three germ layers, including gut-like and respiratory-like epithelium (endoderm), bone and muscle tissues (mesoderm), and neural tissue (ectoderm).

***In vitro* chondrogenic differentiation of reprogrammed human OCs**

Fluorescence microscopy indicated that iPSCs appeared green at day 5 (Fig. 4Aa), while lentivirus-mediated TGF- β 1 transduced iPSCs appeared red (Fig. 4Ab) at day 5 and day 14 (Fig. 4Ac). Increased TGF- β 1 expression in the transduced iPSCs was confirmed using western blotting (Fig. 4Ad).

Histological analysis showed that the iPSCs/TGF- β 1/Alginate/chondrocytes had a round morphology, an obvious cartilage lacuna and a dense sulphated extracellular matrix that stained strongly for toluidine blue compared to the other groups (Fig. 4B).

Real-time PCR was carried out to determine the status of iPSC differentiation at transcription level. As can be seen in Fig. 4C, at day 14, the expression of the cartilage markers collagen II, aggrecan and COMP was significantly higher in iPSCs/ TGF- β 1/ Alginate/chondrocytes than in their respective control cultures including iPSCs/Alginate, iPSCs/Alginate/chondrocytes and iPSCs/TGF- β 1/Alginate ($P < 0.05$), while it was less than that in the positive control group (chondrocytes) ($P > 0.05$). The expression of vascular endothelial growth factor (VEGF), a marker of degenerative chondrocytes, was high in iPSCs/ Alginate and iPSCs/TGF β 1/Alginate, while it was nearly zero for iPSCs/Alginate/ chondrocytes and iPSCs/TGF β 1/Alginate/ chondrocytes.

***In vivo* chondrogenesis of reprogrammed human OCs**

Immunohistochemistry at 6 weeks revealed that iPSCs in the following groups including iPSCs/Alginate/chondrocytes, iPSCs/TGF- β 1/Alginate and iPSCs/TGF- β 1/Alginate/ chondrocytes were obviously surrounded by collagen II-positive extracellular matrix, and iPSCs stained strongly for collagen II in the iPSCs/TGF- β 1/ Alginate/chondrocytes group compared to the other groups (Fig. 5).

Discussion

The innate avascularity, low cellularity and low cell turnover of AC tissue represent major hurdles for natural regeneration following an injury; therefore, the development of treatments allowing successful regeneration of cartilage lesions to avoid further degeneration of focal articular defects represents a major challenge (Giovannini et al., 2010). Many studies have investigated the ability of a variety of cues to replace or repair diseased or damaged cartilage tissue (Lin et al., 2008; Goepfert et al., 2010; Ahmed et al., 2011). Recent developments in regenerative medicine have been driven by discovery of iPSCs, which based on their capacity to proliferate in an undifferentiated state, their potential to differentiate into a variety of different lineages, and their possibility of being personalised therapies, provide an intriguing alternative to the use of stem cells in tissue engineering (Kusuma and Gerecht, 2010). In this study, we have successfully reprogrammed human OCs into fully functional iPSCs by lentivirus-mediated transduction of the four transcription factors OCT4, SOX2, KLF4, and c-MYC. The human iPSCs from OCs displayed hESC-like morphology, expressed hESC marker genes and possessed the ability to differentiate into all three germ layers *in vitro* and *in vivo*. However, it remains unclear whether OC-derived iPSCs may be used as a cell source for cartilage tissue engineering.

At present, two effective methods are used to induce the differentiation of stem cells into chondrocytes: the application of growth factors and co-culture. The traditional strategy has been to add specific growth factors directly to the media of stem cells cultured in a 3D environment. Previous research suggested that hESC-derived EBs in the 3D hydrogels might differentiate toward the chondrogenic lineage in a chemically defined chondrogenic medium supplemented with TGF- β 1 (Hwang et al., 2006). Recently, hESCs cultured in medium with TGF- β 1 produced a cartilage-specific matrix and showed evidence of chondrogenic potential without EB formation (Nakagawa et al., 2009). More recently, Gong et al., (2010) demonstrated that micromass cultures of pluripotent hESCs undergo direct, rapid, progressive, and substantially uniform chondrogenic differentiation in medium containing BMP-2 plus TGF- β 1. All these findings indicate that TGF- β 1 has a significant effect on the chondrogenic differentiation of human and mouse ESCs. However, because TGF- β 1 protein is unstable due to its short half-life *in vitro*, gene delivery may be necessary to provide protein expression for long periods at effective concentrations (Chen et al., 2011). In our study, lentiviral-mediated TGF- β 1 was used to induce the differentiation of iPSCs into chondrocytes. RT-PCR results showed expression of cartilage-related genes, including collagen II, aggrecan, and COMP, in the TGF- β 1-transduced OC-derived iPSCs, suggesting that they acquired a chondrogenic potential. However, VEGF expression was also detected in the TGF- β 1-transduced OC-derived iPSCs, suggesting that the TGF- β 1-transduced OC-derived iPSCs have a degenerative tendency. Patient/disease-specific iPSCs showed some of the disease features recapitulated in tissue culture after directed differentiation of iPSCs (Saha and Jaenisch, 2009; Ye et al., 2009; Liu et al., 2010), and it is possible that a similar process occurred in this study. VEGF is found at the zone of hypertrophic cartilage in the growth plate and plays an integral role in endochondral ossification and longitudinal growth of the skeleton (Gerber et al., 1999; Pufe et al., 2004). VEGF is largely downregulated in adult AC, but this angiogenic factor is expressed by diseased chondrocytes, such as those in

OA (Pufe et al., 2001). Therefore, inhibition of VEGF expression will most likely contribute to the differentiation of OC-derived iPSCs into chondrocytes.

Co-culture is the other method used for the differentiation of stem cells into chondrocytes. The primary advantage of co-culture is that the autocrine and paracrine factors secreted by one cell type readily interact with the other type of cell without direct cell–cell contact, thus mimicking, to a certain extent, the *in vivo* cartilage environment in which the transplanted cells are surrounded by native chondrocytes residing within the tissue (Mo et al., 2009; Aung et al., 2011). Moreover, it allows for cellular crosstalk, instruction, and stabilisation of cellular phenotype by direct physical contact between the surface receptors of different cells, which may lead to more efficient transduction of molecular signals coordinating chondrogenic differentiation (Heng et al., 2004). Co-culturing human MSCs with human ACs has been shown to enhance the mechanical properties and cartilage-specific ECM content of tissue-engineered cartilage formed from MSCs seeded in hyaluronic acid (HA) hydrogels and to decrease expression of collagen type X by MSCs, an important marker of MSC hypertrophy (Bian et al., 2011). Co-culturing equine MSCs with mature ACs improved expression of cartilage-specific genes and produced a more homogeneous production of ECM within the newly formed co-cultured cartilage (Lettry et al., 2010). Furthermore, the chondrogenic potential of hESCs could be enhanced in a high-density micromass pellet co-cultured with irradiated human chondrocytes, and co-culture resulted in a significantly decreased osteogenic potential (Bigdeli et al., 2009). In our study, differentiation of OC-derived iPSCs was induced by co-culturing with ACs, as demonstrated by the expression of cartilage-specific genes. Although the level of cartilage-specific genes in the co-culture system was lower than that in the transfected OC-derived iPSCs, VEGF expression was not detected, possibly due to endogenous inhibition of angiogenesis by factors, including metastatin, present in adult cartilage tissue (Moses et al., 1999; Liu et al., 2001).

One possible approach to overcome the individual shortcoming of the gene-transfer and co-culture systems in the differentiation of OC-derived iPSCs into chondrocytes would be to mix human chondrocytes with TGF- β 1-transduced OC-derived iPSCs in a co-culture system. In our co-culture system, co-culturing the TGF- β 1-transduced OC-derived iPSCs with ACs promoted the differentiation of OC-derived iPSCs into chondrocytes by enhancing the cartilage-specific gene expression, inhibiting VEGF expression and preventing degeneration of the chondrocytes.

To further validate the chondrogenic capacity of reprogrammed human OCs, *in vivo* study was performed. In ectopic cartilage formation, direct evidence of chondrogenic differentiation was collected by immunohistochemical study of collagen II. Collagen II, which is characteristic of a cartilage matrix, was present at higher level in the iPSCs/TGF β 1/Alginate/chondrocytes group, as compared to that in the iPSCs/Alginate/chondrocytes group and the iPSCs/TGF β 1/Alginate group. Histological examination of iPSCs revealed the obvious cartilage tissue formation generated in the co-culture of TGF- β 1-transfected OC-derived iPSCs with ACs in alginate matrix at 6 weeks.

Conclusion

Taken together, our data indicate that iPSCs can be generated from OCs using the defined factors and that *in vitro* co-culture of TGF- β 1-transfected OC-derived iPSCs with ACs in alginate matrix results in significantly improved chondrogenesis of iPSCs as shown by the gene expression profiles and the histological staining. Additionally, *in vivo* study also revealed the obvious cartilage tissue formed in the co-culture of TGF- β 1-transfected OC-derived iPSCs with ACs in alginate matrix. This combinational strategy will promote the use of iPSC-derived tissue in tissue engineering.

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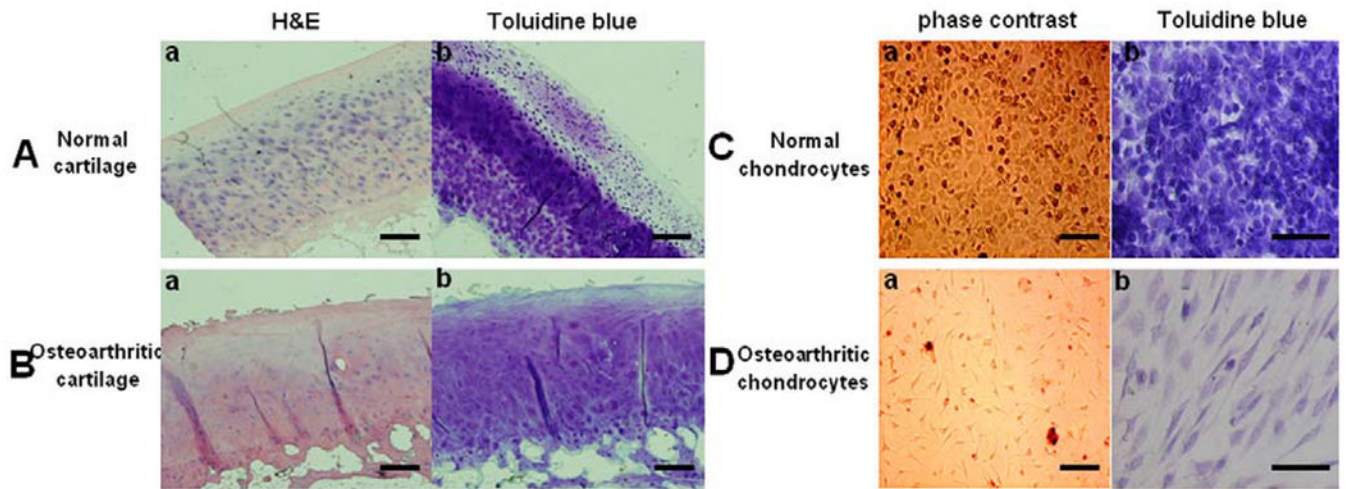


Fig. 1. Normal and osteoarthritic cartilage-derived cells. **(A)** Representative image of tissue from healthy AC. **(B)** Representative image of tissue from osteoarthritic cartilage. **(C)** Representative image of normal chondrocytes. **(D)** Representative image of OCs. Scale bars: 500 μm (**A** & **B**), 100 μm (**C** & **D**).

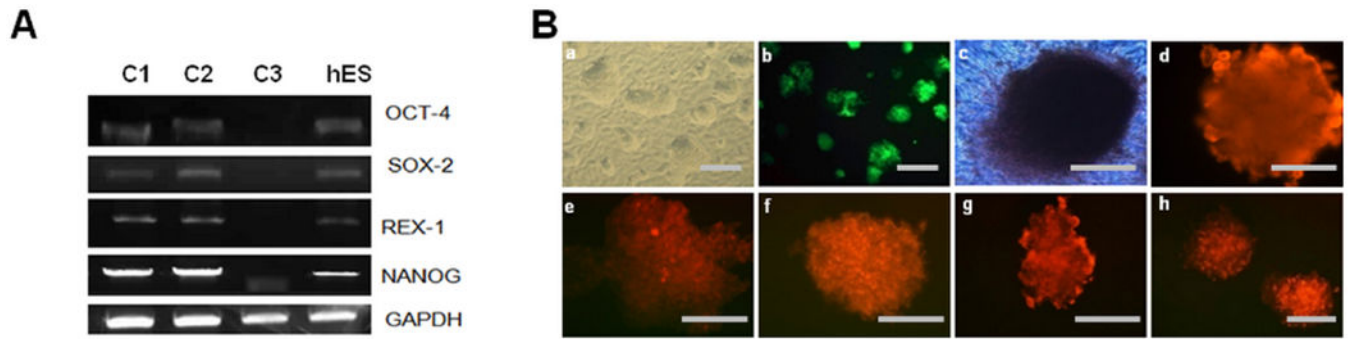
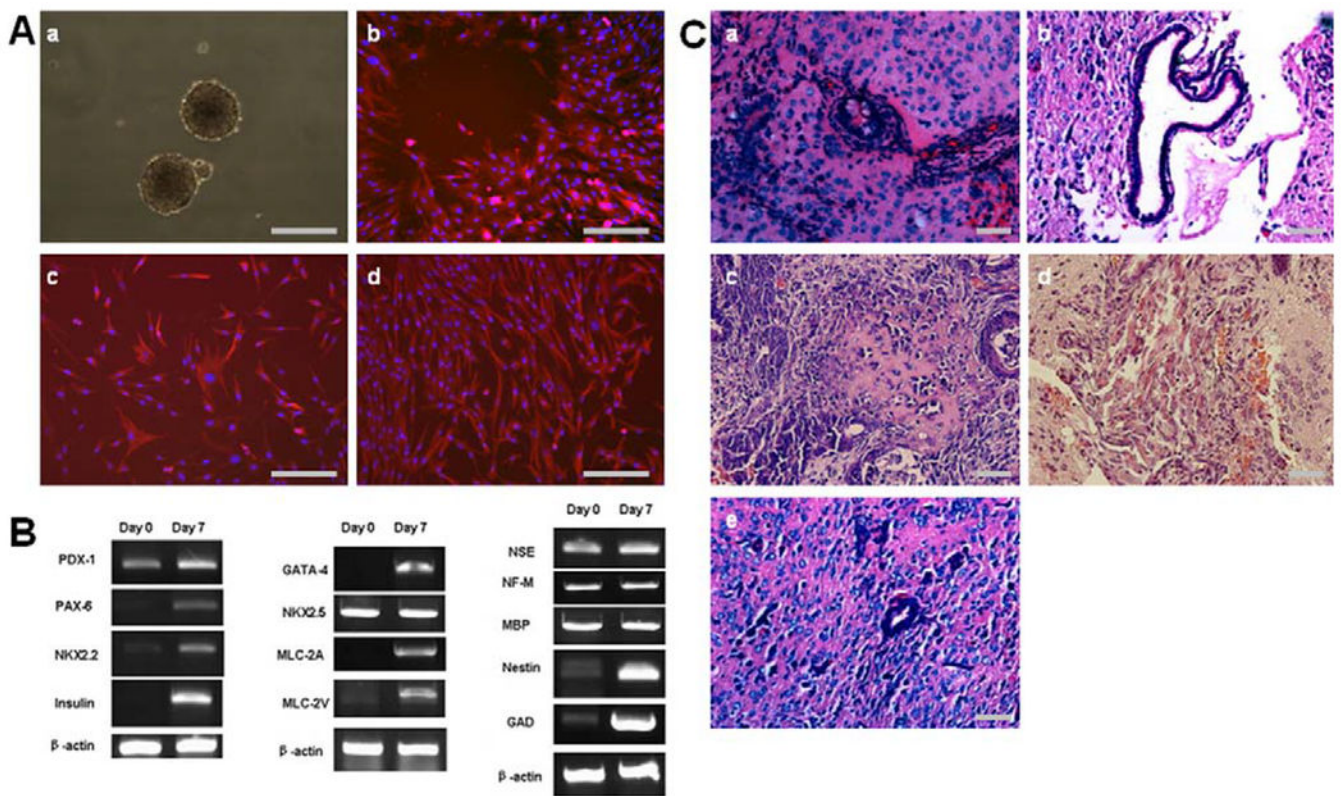


Fig. 2. Generation of iPSCs from human OCs. **(A)** RT-PCR revealed that human OC-derived iPSCs (C1 and C2) expressed hESC marker genes, while the other iPSCs (C3) did not. **(B)** The iPSC colonies were observed 2 weeks after transduction (phase contrast image **(a)** and fluorescence image **(b)**), and the expression of pluripotent markers of human OC-derived iPSCs was analysed by ALP **(c)** and immunofluorescence staining against NANOG **(d)**, SSEA-1 **(e)**, SSEA-4 **(f)**, TAR-60 **(g)**, and TAR-81 **(h)**. Scale bars: 200 μm.

**Fig. 3.**

Differentiation of iPSCs from human OCs *in vitro* and *in vivo*. **(A)** EBs were formed from human OC-derived iPSCs at day 2 **(a)**, and differentiation markers were detected by immunofluorescent staining of AFP **(b)**, nestin **(c)** and desmin **(d)**. **(B)** Direct differentiation of human OC-derived iPSCs into cell lineages of all three germ layers *in vitro*. Differentiated cells positive for PDX-1, PAX-6, NKX2.2, and insulin (endoderm); GATA-4, NKX2.5, MLC-2A, and MLC-2V (mesoderm); NSE, NF-M, MBP, GAD, and nestin (ectoderm) as detected by RT-PCR. **(C)** Teratomas formed 4–6 weeks after human OC-derived iPSCs (clone 1, C1) were injected subcutaneously into four locations on nonobese diabetes/severe-combined immunodeficient (NOD/SCID) mice. Various tissues of the three germ layers were detected by H&E staining, including gut-like epithelium **(a)**, respiratory-like epithelium **(b)**, bone **(c)**, muscle **(d)**, and neural tissue **(e)**. Scale bars: 200 μ m.

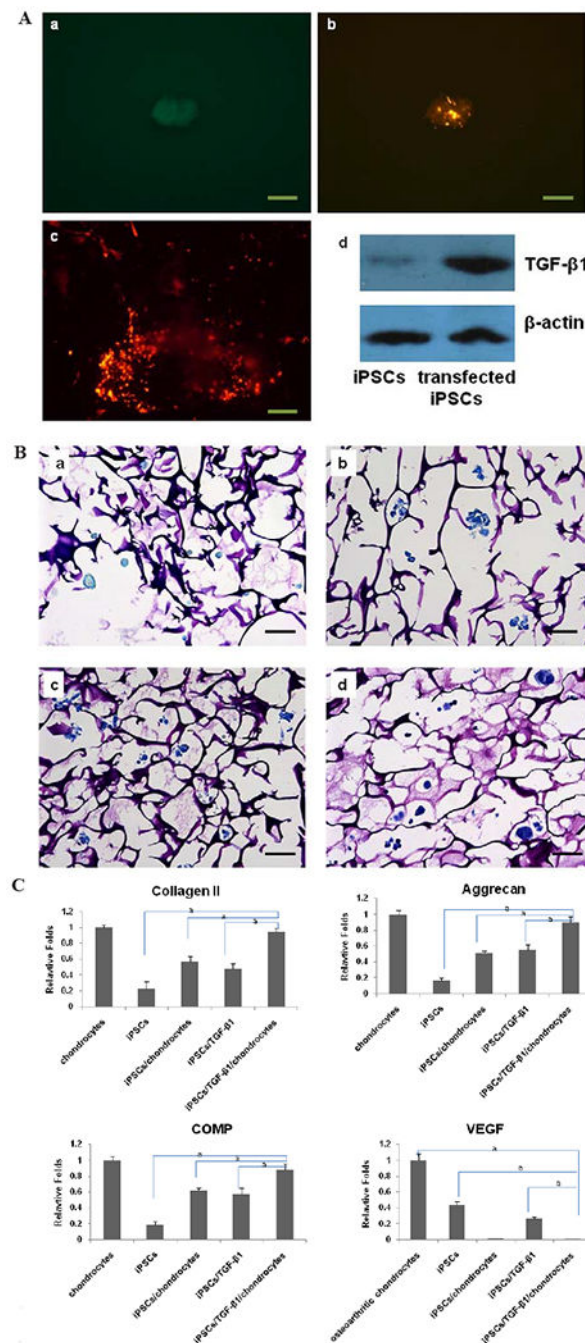


Fig. 4. *In vitro* chondrogenic differentiation of OC-derived iPSCs. (A) Representative images of OC-derived iPSCs with lentivirus-transduced TGF- β 1 at day 5 (a and b) and 14 (c) shown by fluorescence, and a western blot against TGF- β 1 in the nontransfected and transfected iPSCs (d). (B) Toluidine blue staining of iPSCs / Alginate (a), iPSCs / Alginate / chondrocytes (b), iPSCs / TGF- β 1 / Alginate (c) and iPSCs / TGF- β 1 / Alginate / chondrocytes (d). (C) Real-time PCR results showing gene expression for the cartilage markers collagen II, aggrecan and COMP as well as the degenerative cartilage marker

VEGF. ^(a) indicates that expression levels are statistically significant different from iPSCs/TGF- β 1/chondrocytes in alginate matrix. Scale bars: 200 μ m (**A**), 100 μ m (**B**).

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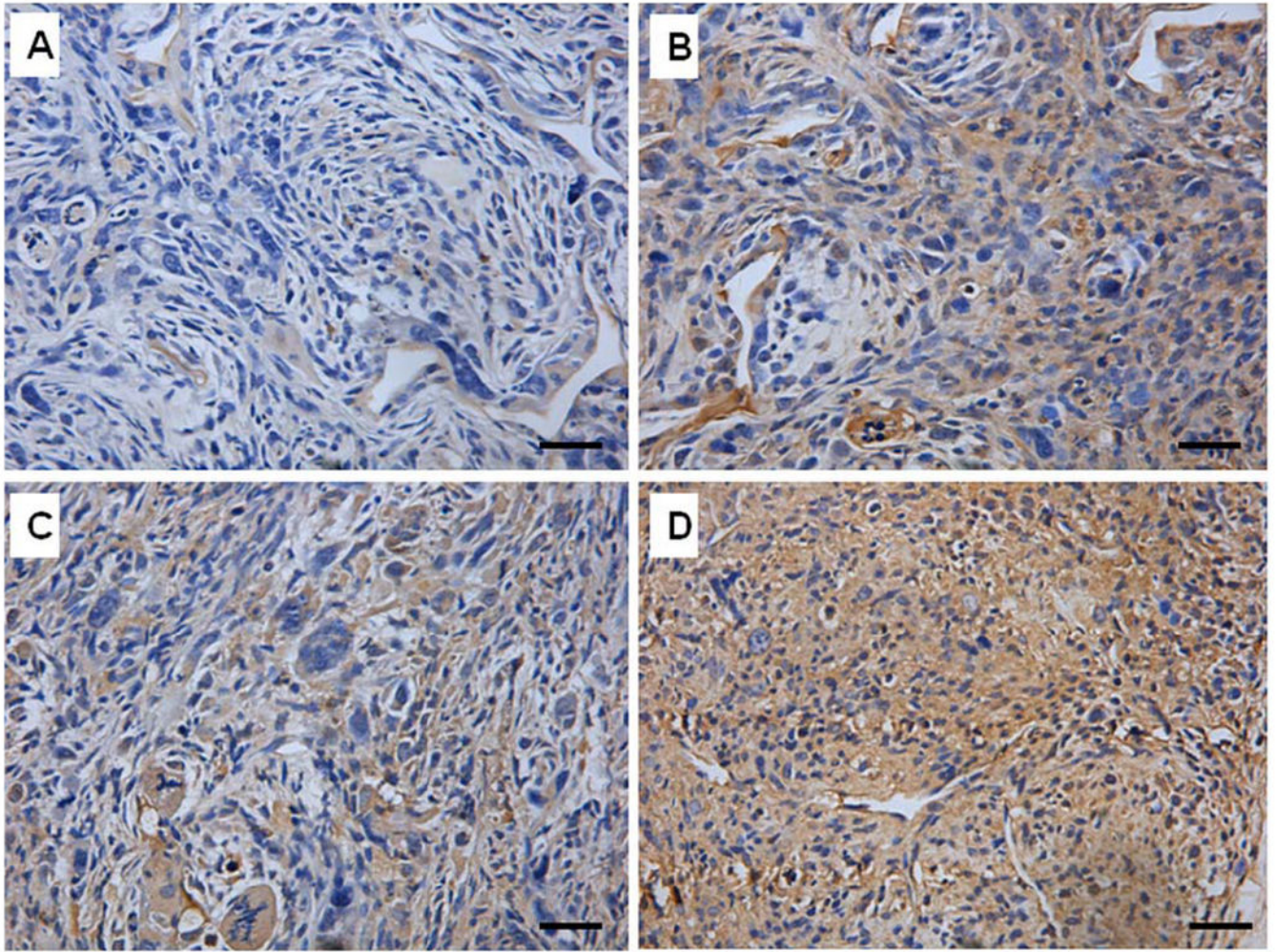


Fig. 5. Immunohistochemical analysis of ectopic cartilage formation of OC-derived iPSCs. Collagen II staining of iPSCs/Alginate (**A**), iPSCs/Alginate/chondrocytes (**B**), iPSCs/TGF- β 1/Alginate (**C**) and iPSCs/TGF- β 1/Alginate/chondrocytes (**D**). Scale bars: 100 μ m.

Table 1.

Applied antibodies

Name	Species	Dilution	Vendor
Anti-human SSEA-1	mouse	1:100	R&D
Anti-human SSEA-4	mouse	1:100	R&D
Anti-human TRA-1-60	mouse	1:500	Chemicon
Anti-human TRA-1-81	mouse	1:500	Chemicon
Anti-human NANOG	goat	1:100	R&D
AFP	mouse	1:500	Dako
Desmin	mouse	1:100	Neomarkers
Nestin	mouse	1:100	Chemicon

Table 2.

List of primers used for PCR

Gene	Forward	Reverse
β -actin	CGCACCTGGCATTGTCAAT	TTCTCCTTGATGTCACGCAC
OCT4	GAGGAGTCCCAGGACATGAA	GTGGTCTGGCTGAACACCTT
SOX2	GCCGAGTGGAAACTTTTGTGTC	GTTCATGTCGGCGTAACCTGT
REX-1	GCGTACGCGAAATTAAGTCCAGA	CAGCATCTAAACAGCTCGCAGAAT
NANOG	TCCCCAAAGCTTTTGTG	GGCCAGTTGTTTTTCTGCCC
GAPDH	GTGTTCTTACCCTCCCAATGTGT	ATTGTCATACCAGGAAATGAGCTT
Pdx-1	TTAGACCGAAGGGGAAAACC	TTAGGGAGCCTTCCAAATGTG
PAX-6	CCGGCAGAAAGATTGTAGAGC	CTAGCCAGGTTGCCAAGAAC
Nkx-2.2	GAAGA ACCCTTCTACGACA	TCGCCGCTTTCGGCTTCTTG
insulin	GCCTTTGTGAACCAACACCTG	GTTGCACTAGTTCTCCAGCTG
glucagon	GAGGGCTTGCTCTCTTCA	GTGAATGTGCCCTGTGAATG
NSE	CCCCTGATCTTCCCGATACAT	CCGATCTGGTTGACCTTGAGCA
NF-M	GAGCGCAAAGACTACTGAAGA	CAGCGATTTCTATATCCAGAGCC
MBP	TTAGCTGAATTCGGGTGTGG	GAGGAAGTGAATGAGCCGGTTA
Nestin	CAGCTGGCGCACCTCAAGATG	AGGGAAGTTGGGCTCAGGACTGG
GAD	GCGCCATATCCAACAGTGACAG	GCCAGCAGTTGCATTGACATAA
GATA	CCAAGCAGGACTCTTGGAAC	CAGCGTGTAAAGGCATCTGA
Nkx2.5	TATAAGCCTACCCCGCCTAT	TATAAGCCTACCCCGCCTAT
MLC-2A	ACAGAGTTTATTGAGTGCCCC	AAGGTGAAGTGTCCCAGAGG
MLC-2V	TATTGGAAACATGGCCTCTGGAT	GGTGTGAAAGGCTGATTACGTT
Collagen II	GAAACCATCAATGGTGGCTTCC	CGATAACAGTCTTGCCCCACTT
Aggreclin	GCCTTGAGCAGTTCACCTTC	CTCTTCTACGGGGACAGCAG
COMP	CAGGACGACTTTGATGCAGA	AAGCTGGAGCTGTCCCTGGTA
VEGF	CGTGTACGTTGGTGCCCGCT	GCTGTAGGAAGCTCATCTCT
β -actin	CGCACCTGGCATTGTCAAT	TTCTCCTTGATGTCACGCAC