Changes in Surface Markers of Human Mesenchymal Stem Cells During the Chondrogenic Differentiation and Dedifferentiation Processes In Vitro

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Objective. To investigate surface markers showing specific changes during the chondrogenic differentiation and dedifferentiation of human mesenchymal stem cells (MSCs).

Methods. Human MSCs from adult bone marrow were subjected to chondrogenic differentiation in 3-dimensional (3-D) alginate culture with or without transforming growth factor-β3 (TGFβ3) for 2 weeks, followed by dedifferentiation in monolayer for 1 week. Surface antigens were selected from those previously reported to show changes in expression during dedifferentiation of human articular chondrocytes (HACs).

Results. Flow cytometry was used to identify 3 groups of surface antigens with differential expression patterns that were quite different from those previously reported on HACs. Two groups of antigens were expressed at high levels on human MSCs. The expression of the first group of antigens (CD44, CD58, CD81, CD90, CD105, and CD166) was decreased reversibly by the 3-D alginate culture and irreversibly in the presence of TGFβ3, except for CD81, which showed reversible changes regardless of TGFβ3. The expression of the second group of antigens (CD49c, CD49e, and CD151) was decreased during chondrogenic differentiation only in the presence of TGFβ3. During all experimental stages, the expression of the third group of antigens (CD14, CD26, CD49f, CD54, CD106, CD119, and CD140a) was maintained at low levels (expressed on <30% of cells), although with some fluctuations.

Conclusion. We speculate that the second group of surface antigens could be negative markers for chondrogenic differentiation of human MSCs.

Mesenchymal stem cells (MSCs) have been widely studied as an alternative cell source for cartilage tissue engineering by virtue of their ability to differentiate into chondrogenic lineage and their phenotypic stability maintained for multiple passages (1–5). Most of the studies until now have focused on finding efficient chondrogenic conditions, such as a 3-dimensional (3-D) environment, growth factors, and/or mechanical forms of stimulation (6–10). Evaluation of the chondrogenic differentiation of MSCs has mainly depended on the expression of chondrocyte phenotypes such as SOX9, type II collagen, and proteoglycans. However, only a little is known about the molecular mechanisms or specific changes in gene expression patterns during the process (10–12), and limited information is available about changes in surface markers (only for CD90 and CD105) during the differentiation of MSCs (13).

Extensive analyses have identified surface markers of MSCs from rats, mice, and humans (14–17). Human MSCs are suggested to be positive for CD44, CD90, and CD105 and negative for CD14, CD34, and CD45 (17). However, their expression pattern was shown to change depending on the culture conditions or specific types of stimulation (13,18,19). Recently, Diaz-Romero et al reported on the expression patterns of 52 surface antigens on human articular chondrocytes (HACs) during the dedifferentiation process in monolayer culture for 2 weeks (20). They identified a subset of surface markers expressed on HACs and suggested that...
HACs had reverted to a primitive stem cell phenotype after dedifferentiation based on the induced expression of some human MSC markers, such as CD10, CD90, CD105, and CD166. The expression of some surface antigens, such as CD14, CD49a, CD54, and CD106, was also shown to be reduced along with the dedifferentiation process in their experiment. It is not clear, however, if these antigens can be regarded as specific markers for active chondrocytes or for the chondrogenic differentiation of stem cells.

In the present study, we investigated changes in the expression of surface antigens on human MSCs during the chondrogenic differentiation and dedifferentiation processes in order to enhance the understanding of phenotypic changes during these processes and to identify specific markers for differentiated chondrocytes. Human MSCs from the bone marrow were subjected to the following 3 conditions: 1) expansion in monolayer culture until passage 3, 2) chondrogenic differentiation in 3-D alginate culture for 2 weeks, and 3) dedifferentiation in monolayer culture for 1 week (Figure 1A). The expression pattern of selected surface markers reported by Díaz-Romero et al (20) was examined during these processes.

### MATERIALS AND METHODS

**Isolation and culture of human bone marrow MSCs.** With informed consent, bone marrow samples were obtained from trabecular bones of 3 hematologically normal patients undergoing routine total hip replacement surgery. The patients had a mean age of 64 years (range 55–73 years). Primary culture of bone marrow stromal cells was performed as previously described (21). Briefly, marrow cells were isolated from trabecular bone marrow samples in Dulbecco’s phosphate buffered saline (PBS) (Gibco BRL, Carlsbad, CA) and harvested by centrifugation at 500g for 5 minutes at room temperature. The cell pellet was resuspended in 10 ml α-minimum essential medium (α-MEM) supplemented with 10% fetal bovine serum (FBS) and passed through nylon mesh (90-μm pore size; Lockertex, Warrington, UK). Aliquots of cell suspensions were diluted with 0.4% (weight/volume) trypan blue, and the number and viability of cells were determined. Cells (1.5 × 10^7) were plated on a 150-mm culture plate and incubated at 37°C in a 5% CO₂ atmosphere. After 6 days, nonadherent hematopoietic cells were removed, and the MSCs on the culture plate were replenished with fresh medium supplemented with 10% FBS. The medium was changed twice per week thereafter. The cells were expanded in a monolayer culture until passage 3 with an interval of ~1–2 weeks for each passage, after which they were moved to a 3-D culture for chondrogenic differentiation.

**Chondrogenic differentiation of human MSCs in alginate layer culture.** The alginate layer culture of human MSCs was performed as described previously (9,22). Human MSCs were suspended in 2% alginate (Sigma, St. Louis, MO) solution at a density of 2 × 10^6/ml. The cell/alginate mixture was added slowly in a 12-well Transwell insert and spread evenly over the surface. The Transwell was then immersed in 1 ml of sterile 102 mM CaCl₂ for 10 minutes. The human MSC/alginate layer was washed twice with 0.15M NaCl for 10 minutes each and once with serum-free chondrogenic defined medium containing high-glucose Dulbecco’s modified Eagle’s medium (DMEM), ITS supplement (1.0 mg/ml insulin from bovine pancreas, 0.55 mg/ml human transferrin, and 0.5 μg/ml sodium selenite), 50 μg/ml L-ascorbic acid 2-phosphate, 100 μM dexamethasone, 40 μg/ml proline, 1.25 mg/ml bovine serum albumin, and 100 μg/ml sodium pyruvate (all from Sigma) (23). The human MSC/alginate layers were then moved to 35-mm culture dishes and overlaid with serum-free chondrogenic defined medium. The samples were cultured at 37°C in a 5% CO₂ atmosphere for 2 weeks for chondrogenic differentiation, with or without treatment with transforming growth factor β3 (TGFβ3). Samples were treated with 10 ng/ml TGFβ3 (R&D Systems, Minneapolis, MN) at the time of media change.

**Dedifferentiation of differentiated cells in a monolayer culture.** After chondrogenic differentiation for 2 weeks, the alginate layer cultures were washed twice with 0.15M NaCl and incubated in 3 volumes of 55 mM sodium citrate solution at 37°C for 15 minutes to dissolve the layer. The cells recovered were harvested by centrifugation at 1,500g for 5 minutes and were replated on a monolayer culture in high-glucose DMEM supplemented with 10% FBS for 7 days.

**Reverse transcriptase–polymerase chain reaction (RT-PCR).** Total RNA was extracted from cells using TRIzol reagent (Invitrogen, Carlsbad, CA). For the alginate culture,
cells were first recovered from the alginate layer as described above. A total of 1 μg RNA was used for the first-strand complementary DNA (cDNA) synthesis using the Superscript First Strand Synthesis System (Roche, Indianapolis, IN), and 2 μg of the synthesized cDNA was used for PCR. Sequences of specific primers were as follows: SOX9, 5ʹ-CACACAGCTCACTGCCTTG-3ʹ (sense) and 5ʹ-TTCGTTATTTTTAGGATCTCTCG-3ʹ (antisense); type II collagen, 5ʹ-GATAATGCACCTTGGACAT-3ʹ (sense) and 5ʹ-CCCCACAATTAAAGAAAG-3ʹ (antisense); and aggrecan, 5ʹ-GAAAGGTGTGTTGTTCACC-3ʹ (sense) and 5ʹ-GTCTACGGGCTTGTTG-3ʹ (antisense). GAPDH (5ʹ-GGACATGAGTCCTTCCAGAT-3ʹ [sense] and 5ʹ-GGTGAAGGTCGGAGTCAACGG-3ʹ [antisense]) was used as an internal control. To exclude the possibility of genomic DNA contamination, RT-PCR was also carried out using total RNA without RT.

Western blot analysis. Cell extracts were prepared using a lysis buffer (40 mM Tris HCl [pH 8.0], 120 mM NaCl, 0.5% Nonidet P40, 100 μg/ml phenylmethylsulfonyl fluoride, and 2 μg/ml each of aprotinin, pepstatin, and leupeptin). The total amount of proteins was determined using the bicinechinonic acid assay (Sigma). The proteins were separated on an 8% sodium dodecyl sulfate polycrylamide gel and transferred onto nitrocellulose membranes (Protran; Schleicher & Schuell, Keene, NH). After blocking, the blots were incubated overnight at 4°C with primary antibodies to type II collagen (Chemicon, Temecula, CA). The blots were then incubated with horseradish peroxidase–conjugated goat anti-mouse IgG for 1 hour at room temperature. Immunoreactive signals were detected using an enhanced chemiluminescence kit (Amer sham, Piscatway, NJ).

Flow cytometry. Cells recovered from monolayer or alginate culture were washed twice in PBS. Single-cell suspensions in PBS (1 × 10⁶ cells/20 μl) were stained at 4°C for 40 minutes with antibodies to CD14, CD26, CD44, CD49c, CD49e, CD49f, CD54, CD58, CD81, CD90, CD105, CD106, CD119, CD140a, CD151, or CD166 according to the manufacturers’ instructions. Isotype IgG was used as a negative control. All antibodies were purchased from BD PharMingen (San Jose, CA) except for CD105, which was purchased from Ancell (Bayport, MN). Detailed information about the antibodies we used is available on our Web site (www.cellntissue.com). The cells were then washed 3 times with 0.5 ml PBS and resuspended in 0.5 ml 2% paraformaldehyde in PBS for fixing. The cells were subjected to flow cytometry using a FACScan instrument and CellQuest software (Becton Dickinson Immunocytometry Systems, San Jose, CA).

RESULTS

Chondrogenic differentiation and dedifferentiation of human MSCs. The overall experimental plan of the study is illustrated in Figure 1A. The experiments were performed independently using MSCs from 3 donors in the whole study. MSCs from each donor showed a similar capacity for cell proliferation and chondrogenic differentiation. In the proliferation stage, MSCs from human bone marrow were expanded for 3 passages in order to yield a sufficient amount of cells with well-maintained stem cell phenotypes. Chondrogenic differentiation of human MSCs was then induced for 2 weeks using a 3-D alginate culture system in chondrogenic defined medium, with or without TGFβ3. Finally, the dedifferentiation stage was carried out by culturing the differentiated cells on a monolayer for 1 week in proliferation medium supplemented with 10% FBS.

When examined by RT-PCR analysis, human MSCs in an initial monolayer culture were negative for the expression of chondrogenic markers such as type II collagen, aggrecan, and SOX9 (Figure 1B). The expression of these chondrogenic markers was increased by 3-D alginate culture of human MSCs. It was significantly higher than that induced by treatment with TGFβ3. We speculate that 3-D alginate culture in defined medium without TGFβ3 was not sufficient to fully develop hyaline cartilage–like constructs with lacunae formation, as shown in our previous study using rabbit MSCs (9). After reculturing the differentiated cells on a monolayer for 1 week, the expression of chondrogenic marker genes was dramatically reduced in all experimental groups, showing that the cells had dedifferentiated and lost their chondrogenic phenotypes. The chondrogenic differentiation and dedifferentiation of human MSCs were also confirmed by Western blotting analysis for type II collagen (Figure 1C).

Effect of TGFβ3 treatment on the expression of selected sets of surface antigens. We analyzed the effect of TGFβ3 treatment on the expression of selected surface antigens that had shown changes in expression on HACs during the dedifferentiation process in the study by Diaz-Romero et al (20). Surface antigens were categorized into 3 groups as described by Diaz-Romero et al: 1) tetraspanins, ectoenzymes, and other miscellaneous surface antigens (CD26, CD81, CD90, and CD151); 2) cell adhesion molecules including integrins (CD44, CD49c, CD49e, CD49f, CD54, CD58, CD81, CD90, CD105, CD106, CD119, CD140a, CD151, or CD166 according to the manufacturers’ instructions. Isotype IgG was used as a negative control. All antibodies were purchased from BD PharMingen (San Jose, CA) except for CD105, which was purchased from Ancell (Bayport, MN). Detailed information about the antibodies we used is available on our Web site (www.cellntissue.com). The cells were then washed 3 times with 0.5 ml PBS and resuspended in 0.5 ml 2% paraformaldehyde in PBS for fixing. The cells were subjected to flow cytometry using a FACScan instrument and CellQuest software (Becton Dickinson Immunocytometry Systems, San Jose, CA).
not significantly changed without TGFβ3 but was reduced reversibly by TGFβ3 treatment during the 3-D alginate culture. The expression of CD26 (peptidase IV) was not changed significantly under any condition.

Among the integrins tested, human MSCs were positive for CD49c (α3 integrin) and CD49e (α5 integrin), but negative for CD49f (α6 integrin) (Figure 3). The expression of these integrins was not further changed by the 3-D alginate culture alone, and only the expression of CD49c was decreased reversibly by TGFβ3 treatment. Among the other cell adhesion molecules tested, human MSCs were positive for CD44 (hyaluronan receptor), CD58 (lymphocyte function–associated antigen 3), and CD166 (activated leukocyte cell adhesion molecule) and negative for CD54 (intercellular adhesion molecule 1) and CD106 (vascular cell adhesion molecule 1). The expression of CD44, CD58, and CD166 was decreased reversibly by the 3-D alginate culture alone and irreversibly in the presence of TGFβ3. The expression of CD54 and CD106 was not observed under any condition.

In the case of cell surface receptors, human MSCs were positive only for CD105 (endoglin) and negative for the other surface antigens (Figure 4). The expression of CD105 was decreased reversibly by the 3-D alginate culture but irreversibly in the presence of TGFβ3. The expression of CD14 (lipopolysaccharide receptor), CD119 (interferon-γ receptor), and CD140a (platelet-derived growth factor receptor α) was not detected at significant levels under any condition.

The results of flow cytometric analyses are summarized in Table 1. The surface antigens were classified into 3 groups based on their changes in expression in 3-D alginate culture in the presence or absence of
TGFβ3 treatment. The surface antigens in group 1 were all expressed at high levels in human MSCs in monolayer culture; their expression was decreased reversibly by the 3-D alginate culture and irreversibly in the presence of TGFβ3, except for CD81, which showed reversible changes regardless of TGFβ3. The surface antigens in group 2 were also well expressed in human MSCs in monolayer culture, and their expression decreased reversibly only in the presence of TGFβ3. The surface antigens in group 3 were expressed at low levels in human MSCs and showed no significant increase in their expression levels during the differentiation and dedifferentiation processes (mostly <30% overall).

Early changes in the expression of group 1 surface antigens. The expression of group 1 surface antigens other than CD81 was examined before 3-D alginate culture and on days 1, 3, 7, and 14 after 3-D alginate culture to determine whether the decrease in expression levels seen at 2 weeks was a rapid change or a gradual one. As shown in Figure 5, their expression levels clearly started to decrease from day 1, but the decrease was gradual with time after 3-D alginate cul-

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**Table 1.** Summary of the results of flow cytometry

<table>
<thead>
<tr>
<th>Group, surface antigen</th>
<th>Group 1</th>
<th>Group 2</th>
<th>Group 3</th>
</tr>
</thead>
<tbody>
<tr>
<td></td>
<td>Monolayer culture (3 passages)</td>
<td>Alinate culture (2 weeks)</td>
<td>Monolayer culture (7 days)</td>
</tr>
<tr>
<td>CD44</td>
<td>98.52 ± 3.24</td>
<td>48.14 ± 8.42</td>
<td>98.72 ± 2.43</td>
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<td>CD58</td>
<td>72.51 ± 23.52</td>
<td>12.42 ± 1.23</td>
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<td>CD9f</td>
<td>99.75 ± 0.27</td>
<td>43.58 ± 8.23</td>
<td>95.84 ± 0.85</td>
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<tr>
<td>CD90</td>
<td>99.97 ± 0.03</td>
<td>18.67 ± 2.54</td>
<td>99.56 ± 0.59</td>
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<tr>
<td>CD105</td>
<td>97.26 ± 4.19</td>
<td>13.56 ± 9.81</td>
<td>90.79 ± 2.78</td>
</tr>
<tr>
<td>CD106</td>
<td>99.68 ± 0.21</td>
<td>26.41 ± 8.54</td>
<td>98.79 ± 2.48</td>
</tr>
<tr>
<td>CD14</td>
<td>99.57 ± 0.31</td>
<td>97.25 ± 3.62</td>
<td>74.33 ± 4.36</td>
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<td>CD49c</td>
<td>99.92 ± 0.01</td>
<td>97.26 ± 0.30</td>
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<tr>
<td>CD151</td>
<td>99.85 ± 0.12</td>
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<td>94.36 ± 3.65</td>
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<td>CD26</td>
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<tr>
<td>CD24f</td>
<td>8.82 ± 1.82</td>
<td>10.28 ± 1.62</td>
<td>2.75 ± 1.47</td>
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<td>CD54</td>
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<td>12.21 ± 7.54</td>
<td>35.47 ± 8.16</td>
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<td>CD106</td>
<td>8.76 ± 1.28</td>
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<td>9.14 ± 0.26</td>
</tr>
<tr>
<td>CD119</td>
<td>13.50 ± 10.68</td>
<td>2.36 ± 0.53</td>
<td>8.66 ± 0.62</td>
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<tr>
<td>CD140a</td>
<td>11.98 ± 1.80</td>
<td>8.61 ± 1.08</td>
<td>7.39 ± 0.17</td>
</tr>
</tbody>
</table>

* Values are the mean ± SD percentage of cells expressing each surface antigen and are derived from 3 independent experiments using mesenchymal stem cells (MSCs) from individual donors. Results were also compared with those reported by Diaz-Romero et al (20). The surface antigens examined were classified into groups 1, 2, and 3 based on their expression patterns during chondrogenic differentiation and dedifferentiation of human MSCs (see Results).
† No data were available for the SD.
‡ CD81 showed a somewhat different expression pattern from the other group 1 antigens, in that transforming growth factor β3 (TGFβ3) did not affect its expression pattern at the dedifferentiation stage.

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**Figure 5.** The early expression pattern of group 1 surface markers other than CD81 during the chondrogenic differentiation of human MSCs. Under the experimental plan shown in Figure 1, the expression of CD44, CD90, CD105, CD166, and CD58 was examined before 3-D alginate culture (I) and on days 1, 3, 7, and 14 (II-1 day, II-3 days, II-7 days, and II-2W, respectively) after 3-D alginate culture by flow cytometry. See Figure 1 for definitions.
tude. This result suggests that the decrease in expression levels was not caused by rapid changes in the culture environment itself and might involve progressive changes in cell physiology. The expression of these surface antigens was also decreased in basal medium (α-MEM, 10% FBS), which suggests again that changes in their expression patterns were caused by changes in the culture environment but were not specific to chondrogenic differentiation (additional data are available at http://www.cellntissue.com/data/bbs/0905070_614216.jpg).

DISCUSSION

Using flow cytometry, we systematically investigated changes in the expression of 16 surface antigens on human MSCs that depended on chondrogenic differentiation and dedifferentiation processes. The surface antigens tested in the present study were chosen based on the previously reported study by Diaz-Romero et al (20), who identified groups of surface antigens specifically expressed in dedifferentiated human chondrocytes. As summarized in Table 1, flow cytometry was used to identify 3 groups of surface antigens showing differential expression patterns during the experimental stages. We could not find any specific marker whose expression was induced by the chondrogenic differentiation of human MSCs, but we did find subsets of surface antigens whose expression was reduced during the process. No specific expression pattern was observed according to the categories of surface antigens described by Diaz-Romero et al (20), suggesting that specific groups of antigens, such as cell adhesion molecules, cell surface receptors, and the like, have no apparent roles in chondrogenic differentiation and dedifferentiation. Specific functions of the surface antigens, particularly in cartilage, have been well summarized by Diaz-Romero et al (20). The expression pattern of the surface markers will be discussed in detail group by group and compared with the results reported by Diaz-Romero et al (20).

The group 1 antigens (CD44, CD58, CD81, CD90, CD105, and CD166) were all well-known surface markers of human MSCs except for CD81 (17,24) and were also expressed at high levels on expanding human MSCs in the present study. Their expression levels decreased during the chondrogenic differentiation process, with or without TGFβ3, consistent with a previous study showing a similar result for CD44, CD90, and CD105 in micromass culture of human MSCs (13). We speculate that the group 1 surface antigens might be markers of human MSCs for changes in the microenvironment, but not specific negative markers in active chondrocytes. First, their expression level started to decrease right after the 3-D alginate culture without TGFβ3 treatment and became significantly low on days 7 and 14, when no significant chondrogenic differentiation was observed. Second, the irreversible decrease in their expression induced by TGFβ3 treatment was inconsistent with their high level of expression in dedifferentiated HACs (20). Further, some of the markers such as CD44, CD58, and CD105 were also shown to be expressed in active HACs (20,25).

The group 2 surface antigens (CD49c, CD49e, and CD151) were also expressed at high levels in human MSCs. Their expression levels showed no significant change at all stages in the absence of TGFβ3, but decreased reversibly during the differentiation stage only in the presence of TGFβ3. This result suggests that they could be negative markers, at least for the chondrogenic differentiation of human MSCs. Further analyses are necessary to address this possibility. However, they do not appear to be negative markers for chondrocytes because they were shown to be expressed at high levels in active HACs (20,25). The group 3 surface antigens were expressed at low levels in human MSCs, and their expression levels fluctuated without a significant increase during the differentiation and dedifferentiation processes. Also, the low expression levels of CD54 and CD106 are inconsistent with their high levels of expression both in active and in dedifferentiated HACs (20).

Our results suggest that expression patterns of the surface markers overall were quite different between HACs and human MSCs during differentiation and/or dedifferentiation processes. We speculate that the chondrogenic differentiation condition of human MSCs in vitro is artificial, especially given continuous treatment with TGFβ3, and that it does not completely recapitulate the dynamic changes seen during chondrogenic development in vivo. There might also be innate differences between MSCs differentiating into chondrocytes in vivo and human MSCs isolated from adult bone marrow in vitro. Therefore, expression patterns of surface antigens in this study may not definitely represent the phenomena in vivo, but these antigens could be specific markers for chondrogenic differentiation of human MSCs in vitro.

Human MSCs isolated from adult bone marrow in this study were positive for CD44, CD49c, CD49e, CD81, CD90, CD105, CD151, and CD166 and were negative for CD14, CD26, CD49f, CD54, CD106, CD119, and CD140a. Most of them are already known
in human MSCs (17,24,26), but the expression pattern of CD49c was introduced for the first time in the present study. The expression of CD81 both on human MSCs and on chondrocytes has been reported previously, but its function has not yet been identified (27,28). Grogan et al showed that CD44, CD151, and CD49c were expressed at significantly higher levels on chondrocytes with higher chondrogenic capacity (25). However, they were all highly expressed on human MSCs in the present study and in other studies, and it is questionable whether they could be specific markers for active chondrocytes (27). CD58 was expressed at moderate levels on human MSCs in the present study. Independent groups have reported its expression either on normal HACs (20) or, in contrast, only in arthritic joints (29). The expression levels of CD54 and CD106 were shown to be variable depending on the donor ages and passage numbers (26).

Previously, clonal populations of dedifferentiated adult HACs were reported to gain stem cell properties, thus becoming able to differentiate into the chondrogenic, osteogenic, and adipogenic lineages (30). Diaz-Romero et al also suggested that dedifferentiated HACs had reverted to a primitive phenotype, at least in terms of surface markers, because some MSC markers such as CD10, CD90, CD105, and CD166 were up-regulated after dedifferentiation (20). However, we are not sure that human MSCs, once they have differentiated into the chondrogenic lineage with TGFβ3 treatment, could regain their stem cell capacity after dedifferentiation, because typical human MSC markers such as CD44, CD90, and CD105 were lost irreversibly during differentiation. The irreversible loss of typical human MSC markers could be an unexpected outcome of long-term treatment with TGFβ3 in vitro. It will be an interesting project to examine the stem cell phenotype, after dedifferentiation, of human MSCs that have differentiated with TGFβ3 treatment.

AUTHOR CONTRIBUTIONS

Drs. Min and Park had full access to all of the data in the study and take responsibility for the integrity of the data and the accuracy of the data analysis.

Study design. Min, Park.

Acquisition of data. Lee.

Analysis and interpretation of data. Lee, Choi.


Statistical analysis. Lee.

REFERENCES


