

Unfortunately, removal of this important shock absorber leads to accelerated osteoarthritis⁹⁻¹². Thus, new techniques designed to restore meniscal structure and function following injury are needed.

Mesenchymal stem cells are multipotent cells present in mature individuals and readily accessible from peripheral connective tissue sites such as bone marrow^{13,14}, periosteum¹⁵, adipose¹⁶, and the synovial lining of major joints¹⁷. These cells, which are capable of differentiating into osteoblasts, chondrocytes, adipocytes, and myocytes, represent an attractive potential means of regenerating damaged connective tissues including intra-articular structures of the knee, such as the meniscus¹⁸⁻²⁰.

Recent literature has suggested that synovial tissue-derived mesenchymal stem cells may have the potential to aid in healing and regeneration of cartilage injuries, such as those involving the meniscus^{18,20-24}. Synovial mesenchymal stem cells represent an attractive cell source because they can be harvested in a minimally invasive manner from synovial tissue and are easily expanded in

culture^{18,20-22}. In addition, multiple investigators have found that synovial mesenchymal stem cells possess a particularly high capacity for chondrogenic differentiation and proliferation compared with mesenchymal stem cells obtained from other tissues, such as bone marrow or periosteum^{18,20,22}.

Synovial mesenchymal stem cells are also capable of adhering to damaged intra-articular structures such as the meniscus and participating in the repair process in rat models²³⁻²⁶. In rats, however, the animals' innate regenerative capacity limits the ability of the investigator to compare the effect of the cells on tissue regeneration in experimental versus control groups²⁴. In contrast to smaller rodents, the limited inherent regenerative capacity of the rabbit meniscus makes the rabbit a more favorable animal model in which to evaluate meniscal regeneration techniques. Moon et al. found that, following medial meniscectomy, the rabbit meniscus only partially regenerated with primarily fibrous tissue²⁷. More recently, rabbit models have been used to study the effects

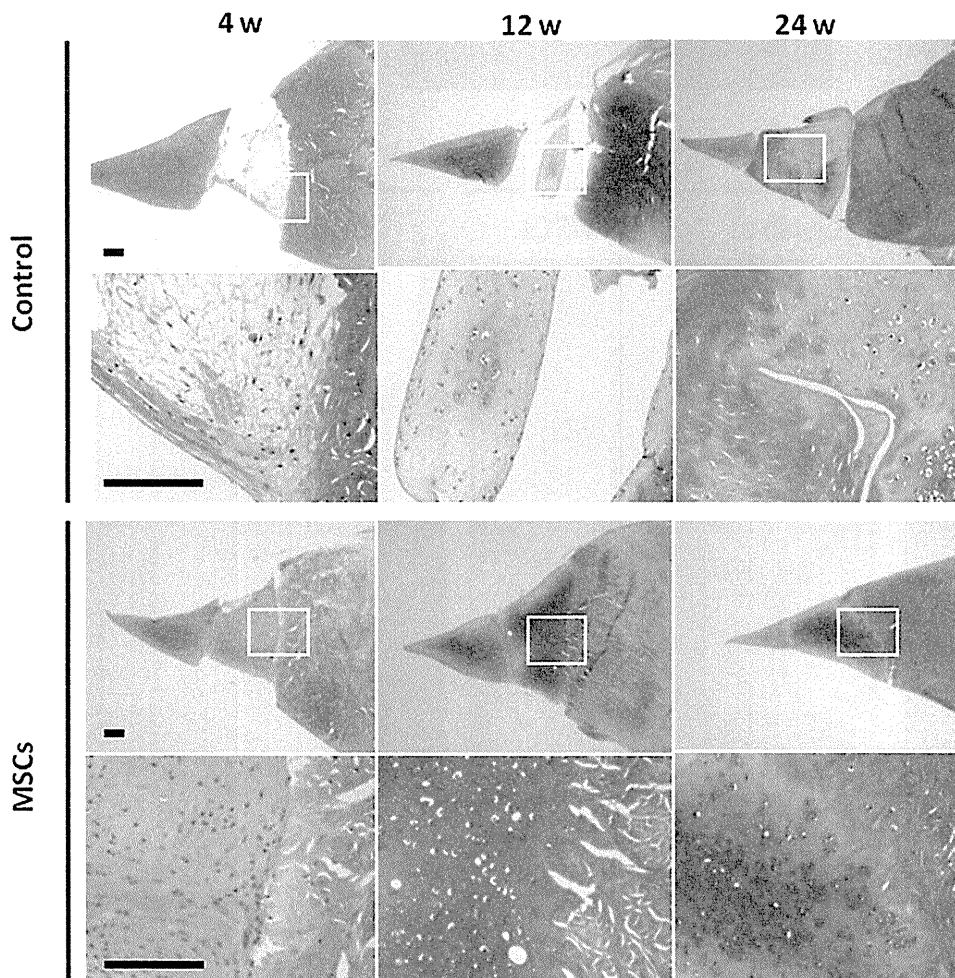


Fig. 5-A

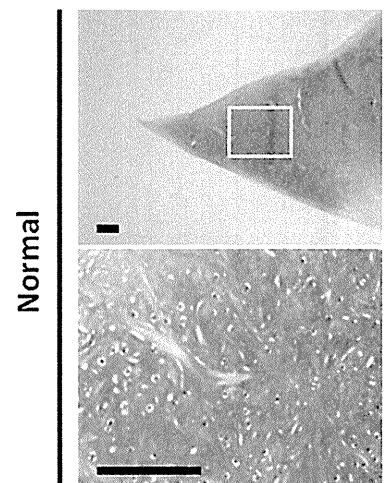


Fig. 5-B

Figs. 5-A through 5-D Synovial mesenchymal stem cells (MSCs) promote the meniscal regeneration (histological observation). **Fig. 5-A** Low and high-power images of representative sections of regenerated meniscus stained with safranin O at four, twelve, and twenty-four weeks after implantation of synovial mesenchymal stem cells. The inset shows the area seen at higher magnification in the photomicrograph below. Scale bars represent 200 μ m. **Fig. 5-B** Low and high-power representative images of the normal meniscus stained with safranin O. Scale bars represent 200 μ m.

of a variety of surgical interventions aimed at regenerating more normal meniscal tissue with use of scaffolds composed of collagen, polyglycolic acid, or gelatin with or without a potential biological regenerative stimulus²⁸⁻³⁰.

The purpose of this study was to evaluate the effect of synovial mesenchymal stem cell supplementation on meniscal regeneration in a rabbit model of partial meniscectomy.

Materials and Methods

Animals

The experimental protocols were approved by the Scott & White Institutional Animal Care and Use Committee and Texas A&M University Institutional Biosafety Committee. Two mature New Zealand White rabbits were used as cell donors for two experimental groups as follows: a single mature wild-type rabbit served as the synovial donor for Group A (fifteen rabbits), and a single transgenic rabbit bred to ubiquitously express green fluorescent protein (GFP; Kitayama Labes, Nagano, Japan)^{31,32} served as the synovial donor for Group B (four rabbits).

Tissue Harvesting and Mesenchymal Stem Cell Preparation

Through a medial parapatellar surgical approach, the right knee of one mature wild-type New Zealand White rabbit was accessed and synovium from the medial, lateral, and suprapatellar regions of the joint was resected. Bone marrow was harvested from the femoral intramedullary canal for comparison of the proliferation capacity of synovial and bone-marrow mesenchymal stem cells.

Isolated cells were cultured for three passages, and a fluorescent lipophilic tracer, CM-DiI (chloromethylbenzamido DiI; Invitrogen, Carlsbad,

California), was added to the cultured synovial tissue-derived cells (see Appendix). These CM-DiI-labeled cells were then used for implantation in experimental knees of Group-A rabbits to evaluate the effect of synovial mesenchymal stem cell supplementation on the quality and quantity of meniscal regeneration, as detailed below in the section on meniscectomy and cell implantation.

A separate subset of the experiment was performed with use of GFP-positive synovial tissue-derived cells obtained from the knee synovium of a transgenic New Zealand White rabbit bred to ubiquitously express GFP (Kitayama Labes)^{31,32}. These cells were processed as described above without the addition of CM-DiI. These GFP-positive synovial tissue-derived cells were implanted in the experimental knees of Group-B rabbits to track not only implanted cells but also their progeny, as detailed below in the section on meniscectomy and cell implantation.

Colony-Forming Assay

To compare proliferation potentials, synovial tissue-derived cells and bone marrow-derived cells from passage 3 were plated in 60-cm² dishes at densities of 100 or 1000 cells per dish, cultured in complete medium for fourteen days, and stained with 0.5% crystal violet in methanol for five minutes. The number of colonies was then counted. Colonies that were <2 mm in diameter and colonies that were faintly stained were ignored.

In Vitro Differentiation Assay

Isolated synovial tissue-derived cells were cultured under conditions conducive to adipogenesis, calcification, and chondrogenesis to assess for multipotentiality (see Appendix).

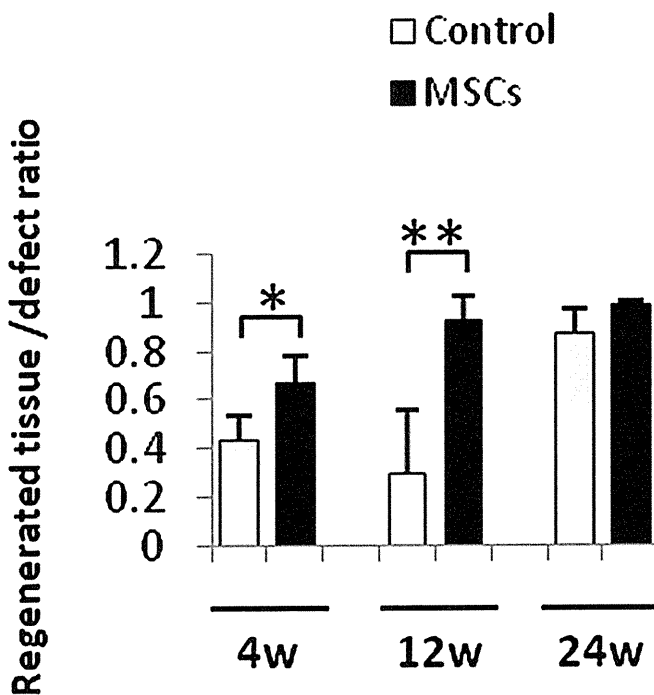


Fig. 5-C

Fig. 5-C Regenerated tissue-to-defect ratios are displayed as the mean and the standard deviation for synovial mesenchymal stem cell (MSCs) and control groups at each end point. *The difference between the groups was significant ($p < 0.05$). **The difference between the groups was significant ($p < 0.01$).

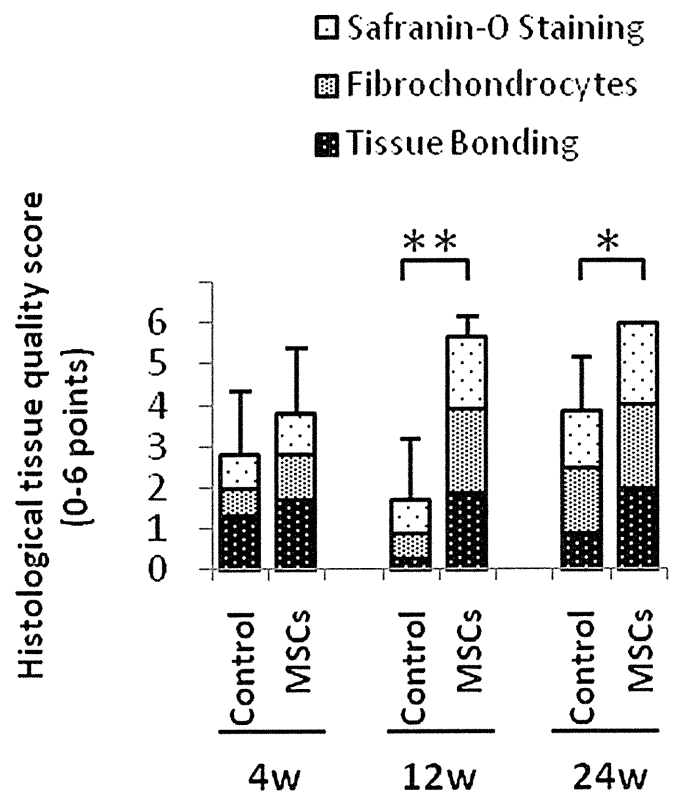


Fig. 5-D

Fig. 5-D Results of the histological scoring system for regenerated meniscus. The scores are displayed as the mean and the standard deviation. *The difference between the groups was significant ($p < 0.05$). **The difference between the groups was significant ($p < 0.01$).

Meniscectomy and Implantation of Synovial Mesenchymal Stem Cells

Nineteen additional mature New Zealand White rabbits were divided into two groups, consisting of fifteen rabbits in Group A and four in Group B.

Group A

With the animal under general anesthesia, each knee was approached through a medial parapatellar arthrotomy, maximally flexed, and a reproducible 1.5-mm-diameter full-thickness cylindrical defect was produced in the avascular inner two-thirds³³ of the anterior portion of the medial meniscus with use of a biopsy punch (Miltex, York, Pennsylvania).

With the tibial joint surface facing upward, 2×10^6 CM-Dil-labeled synovial tissue-derived mesenchymal stem cells in 50 μ L of phosphate-buffered saline solution (PBS)³⁴ were placed directly into the meniscal defect in each of the right knees with use of a 27-gauge needle. Knees were then held stationary for ten minutes during wound closure. In left knees, the same volume of plain PBS was implanted as a control, and the knees were once again held stationary for ten minutes.

Capsule and skin were closed in layers with absorbable suture. Rabbits were allowed to move freely in their cages. Medial menisci from both knees were harvested at four, twelve, and twenty-four-week end points.

Group B

A similar procedure was performed on the remaining four rabbits, with the goal of tracking the fate of implanted mesenchymal stem cells and their progeny. For this purpose GFP-positive synovial mesenchymal stem cells were implanted in

the right knees in the same manner as those in Group A. The same volume of PBS was placed in the left knees once again. Medial menisci from the right and left knees were then harvested at one-day, one-week, four-week, and twelve-week end points.

Histological Analysis

Meniscal specimens from Group A were fixed in 4% paraformaldehyde, decalcified, and embedded in paraffin. Specimens were then sectioned into slices 5 μ m thick in the radial plane. The quantity and quality of regenerated meniscal tissue were then evaluated as follows:

Tissue Quantity Analysis

The quantity of tissue regeneration was evaluated with use of random tissue sections obtained from the central portion of the meniscal defect as shown in Figure 1 and described in detail in the Appendix. The area of the original defect (D) and the area occupied by regenerated tissue inside the defect (R) were calculated with use of Photoshop CS3 software (Adobe Systems, San Jose, California). Tissue regeneration is expressed as the ratio of regenerated tissue area to the entire defect area (regeneration ratio = R/D).

Tissue Quality Scoring Analysis

A quantitative scoring system evaluating three dimensions of meniscal regeneration was used (see Appendix)³⁰. Sections to be scored were stained with safranin O. Histological scoring was performed by two investigators blinded to treatment category.

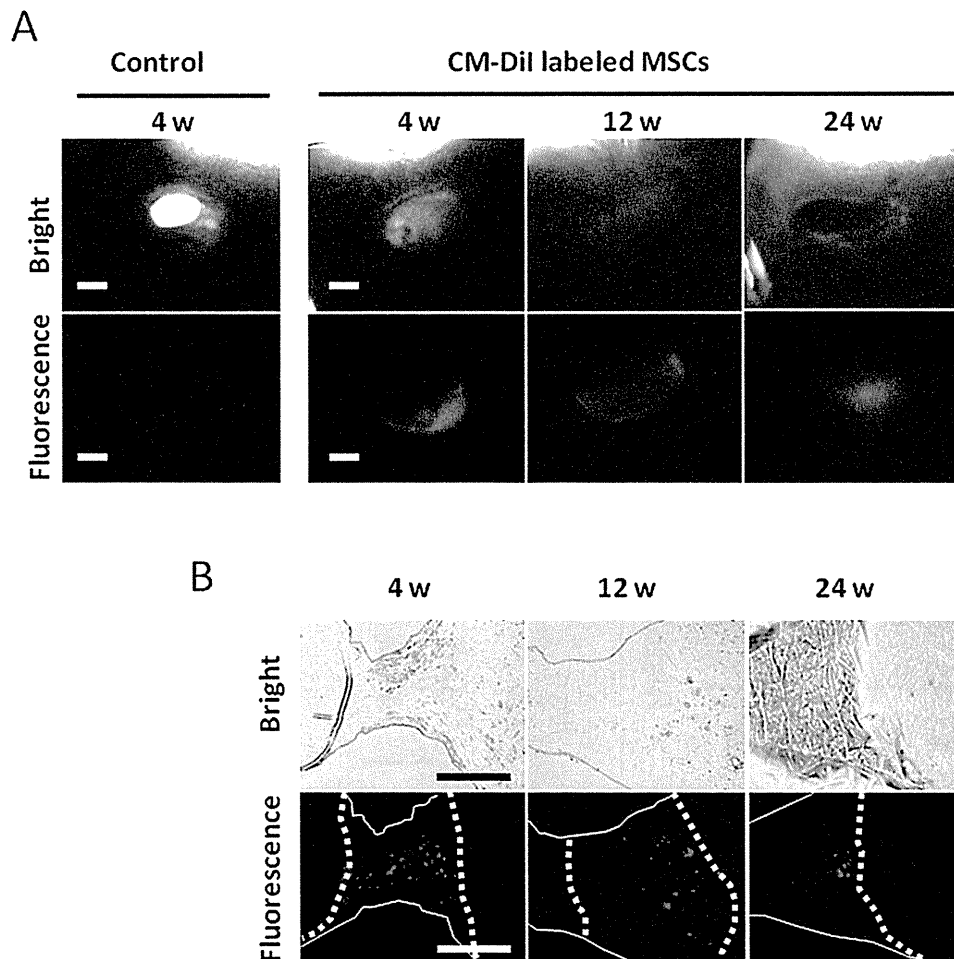


Fig. 6

Figs. 6-A and 6-B CM-Dil-labeled synovial mesenchymal stem cells (MSCs) adhere to sites of meniscal injury and remain at twenty-four weeks. Representative macroscopic appearance (**Fig. 6-A**) and histological sections (**Fig. 6-B**) of the meniscal defect after implantation of CM-Dil-labeled synovial mesenchymal stem cells under bright light (top) and fluorescence (bottom). In the histological sections (**Fig. 6-B**), the white solid line indicates the outer edge of the meniscus, and the white dotted line indicates the border between native meniscus and regenerated tissue. Scale bar represents 400 μ m.

Fluorescent Microscopy

Photographs of the gross and microscopic appearance of each Group-A meniscal specimen were made under fluorescence to demonstrate the presence or absence, as well as relative density, of CM-DiI-labeled synovial mesenchymal stem cells within the defects and surrounding intact meniscal tissue.

To determine the fate of the implanted GFP-positive synovial mesenchymal stem cells and their progeny in Group-B rabbits, menisci from both knees were harvested, fixed in 4% paraformaldehyde, and transferred to 20% sucrose solution. Specimens were flash-frozen, cut in a cryostat, and observed under fluorescent microscopy.

Immunohistochemistry for Type-I and Type-II Collagen

Frozen sections from the menisci of Group-B rabbits were also used to detect type-I and type-II collagen synthesis with use of standard immunohisto-

chemistry techniques (see Appendix). Background nuclei were counterstained with 4', 6-diamidino-2-phenylindole dihydrochloride (DAPI).

Statistical Methods

An a priori power calculation was performed with use of mean and standard deviation assumptions based on previously published data³⁰. A sample size of five rabbits was found to be sufficient to detect a difference of 3 points in the tissue quality score at the twelve-week end point, with a power of 89% at a significance level of 0.05. The Mann-Whitney U test was used to compare synovial mesenchymal stem cells and control groups at each period. P values of <0.05 were considered significant.

Source of Funding

Funding for this study was provided by an internal institutional grant from the Scott & White Research Grants Program as well as a grant from the National Institutes of Health (NIH/NCRR grant P40 RR 17447).

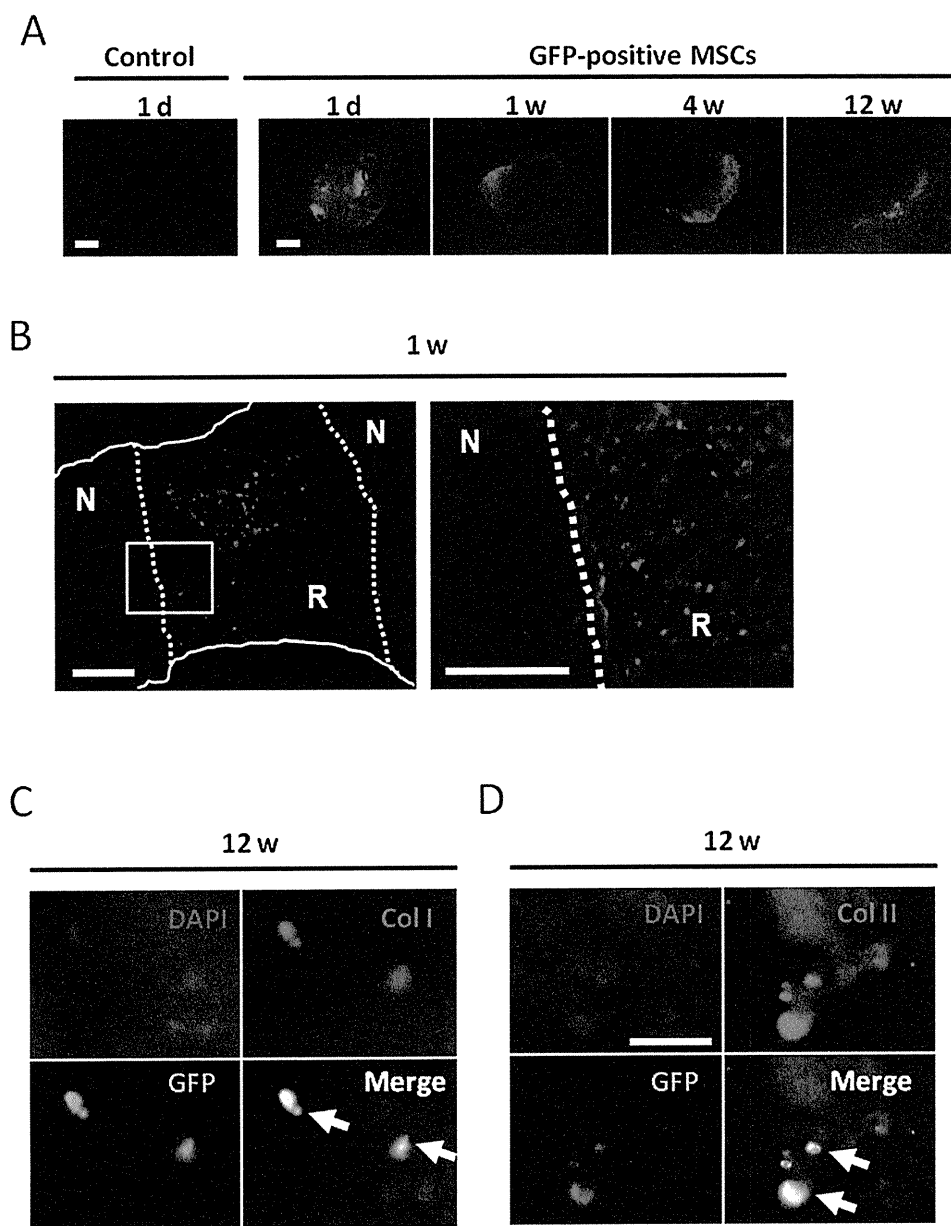


Fig. 7

Figs. 7-A and 7-B Green fluorescent protein (GFP)-positive synovial mesenchymal stem cells (MSCs) adhere to sites of meniscal injury in experimental knees, where they produce type-I and type-II collagen. Representative macroscopic appearance (**Fig. 7-A**) and histological sections (**Fig. 7-B**) of the meniscal defect one day to twelve weeks after the implantation of GFP-positive synovial mesenchymal stem cells under fluorescence. In the histological sections (**Fig. 7-B**), the white solid line indicates the outer edge of the meniscus. The white dotted line indicates the border between native meniscus (N) and regenerated tissue (R). The inset shows the area seen at higher magnification in the photomicrograph on the right. The scale bars represent 200 μm . **Figs. 7-C and 7-D** Fluorescent images of the regenerated meniscus at twelve weeks after implantation of GFP-positive synovial mesenchymal stem cells. The sections were immunostained with anti-type-I collagen (**Fig. 7-C**) and anti-type-II collagen (**Fig. 7-D**). Nuclei were counterstained with DAPI (4', 6-diamidino-2-phenylindole dihydrochloride; blue) to demonstrate the cellular background (GFP-positive cells as well as cells not expressing GFP). GFP-positive cells are shown in green (bottom left), and type-I collagen-producing cells (**Fig. 7-C**) and type-II collagen-producing cells (**Fig. 7-D**) are shown in red (top right). Arrows in merge (bottom right) indicate double-positive cells of GFP and type-I collagen (**Fig. 7-C**) or type-II collagen (**Fig. 7-D**). Scale bar represents 25 μm . Col I = type-I collagen, and Col II = type-II collagen.

TABLE I Regenerated Tissue-to-Defect Ratio

	Ratio*	P Value
4 weeks		
Control	0.44 ± 0.10	0.028†
MSC	0.67 ± 0.11	
12 weeks		
Control	0.29 ± 0.27	0.009‡
MSC	0.93 ± 0.10	
24 weeks		
Control	0.87 ± 0.10	0.077
MSC	0.98 ± 0.03	

*Regenerated tissue-to-defect ratios (range, 0 to 1) are given as the mean (and standard deviation) for synovial mesenchymal stem cell (MSC) and control groups at each end point. †The difference between the groups was significant ($p < 0.05$). ‡The difference between the groups was significant ($p < 0.01$).

Results

Synovial Mesenchymal Stem Cells Have High Proliferation Capacity and Multipotentiality

Abundant multipotent cells were isolated from the synovial tissue of the initial donor rabbit. Isolates produced spindle-shaped cells that adhered to plastic culture dishes under standard conditions, and appeared morphologically similar to the cultured bone-marrow mesenchymal stem cells (Fig. 2-A). While both synovial and bone-marrow mesenchymal stem cells were capable of forming single-cell colonies, the synovial mesenchymal stem cells demonstrated greater colony-forming potential when plated at densities of 100 cells per 60 cm² and 1000 cells per 60 cm² (Figs. 2-B and 2-C).

Isolated synovial mesenchymal stem cells were capable of differentiation into adipocytes and chondrocytes, and were calcified under the proper conditions (Figs. 2-D, 2-E, and 2-F). In adipogenic medium, colonies stained well with oil red O (Fig. 2-D). In calcification medium, colonies produced abundant calcium (Fig. 2-E). In chondrogenic medium, smooth glistening pellets, which stained diffusely with safranin O, were produced (Fig. 2-F). Thus, the isolated synovial mesenchymal stem cells met the requirements for multipotent mesenchymal stromal cells as set forth by Dominici et al.³⁵.

Locally Implanted Synovial Mesenchymal Stem Cells Promote Meniscal Regeneration

A representative photograph of the in vivo 1.5-mm-diameter full-thickness meniscal biopsy of the avascular portion of the anterior horn of the medial meniscus is shown in Figure 3-A. Application of either synovial mesenchymal stem cells suspended in PBS (right knees) or PBS alone (left knees) is demonstrated in Figure 3-B.

A gross representation of regenerated meniscal tissue within the 1.5-mm cylindrical defects is shown in Figure 4. In the control knees, grossly visible, full-thickness defects remained in

four of five specimens at four and twelve weeks (asterisks in Figure 4). In knees supplemented with synovial mesenchymal stem cells, however, such defects remained in only one specimen at four weeks and none at twelve weeks. The macroscopic difference between mesenchymal stem cell and control groups is less dramatic at twenty-four weeks, by which time four of five defects in control knees and five of five defects in knees supplemented with mesenchymal stem cells were completely covered with at least partial-thickness tissue on gross inspection.

Representative sections of regenerated meniscus stained with safranin O at four, twelve, and twenty-four weeks after implantation of synovial mesenchymal stem cells, as well as sections of normal meniscus stained with safranin O, are shown in Figures 5-A and 5-B. The quantity of regenerated meniscal tissue was greater when measured histologically in knees receiving synovial mesenchymal stem cells, reaching significance at four and twelve weeks ($p < 0.05$) (Fig. 5-C, Table I). By twelve weeks, an average of >90% of the cylindrical defect had filled in with regenerated tissue in knees supplemented with synovial mesenchymal stem cells compared with <30% in controls.

The quality of meniscal regeneration was also superior in knees supplemented with synovial mesenchymal stem cells at all end points, achieving significance at twelve and twenty-four weeks (Fig. 5-D, Table II). By twenty-four weeks, all menisci from knees treated with synovial mesenchymal stem cells scored the highest possible score in each of the three quality domains measured—staining with safranin O, existence of fibrochondrocytes, and extent of bonding between regenerated tissue and the borders of the meniscal defect.

Synovial Mesenchymal Stem Cells Adhere to Sites of Meniscal Injury and Differentiate into Type-I and II Collagen-Producing Cells

Gross and histologic fluorescence imaging revealed the presence of CM-DiI-labeled cells in high concentration within the meniscal

TABLE II Histological Tissue Quality Score

	Score*	P Value
4 weeks		
Control	2.8 ± 1.5	0.29
MSC	3.8 ± 1.6	
12 weeks		
Control	1.7 ± 1.5	0.008†
MSC	5.7 ± 0.4	
24 weeks		
Control	3.9 ± 1.3	0.021‡
MSC	6.0 ± 0	

*Results of histological scoring system (range, 0 to 6) for regenerated meniscus. The values are given as the mean and the standard deviation. †The difference between the mesenchymal stem cell (MSC) and control groups was significant ($p < 0.01$). ‡The difference between the groups was significant ($p < 0.05$).

defects of Group-A rabbits at four, twelve, and twenty-four weeks following implantation (Figs. 6-A and 6-B). Although the number of CM-DiI-labeled cells decreased with time, a small number of CM-DiI-labeled cells were observed in the regenerated tissue even at twenty-four weeks. Significant fluorescence was not detected elsewhere in the meniscus, suggesting that implanted mesenchymal stem cells preferentially adhered to the site of meniscal injury. No CM-DiI-labeled cells were detected by fluorescence in control specimens.

In Group-B rabbits, GFP-positive synovial mesenchymal stem cells were found within the meniscal defect as early as one day after injury and remained in high concentrations as late as twelve weeks after injury (Figs. 7-A and 7-B). No GFP-positive cells were detected in the contralateral, control menisci, thus supporting the assumption that neither synovial mesenchymal stem cells implanted in the experimental knee nor their progeny directly affect regeneration at the distant injury site in the contralateral knee.

The number of GFP-positive cells within the defect decreased with time from one day to twelve weeks. Even at twelve weeks, however, a population of GFP-positive cells remained in the regenerated tissue and expressed both type-I collagen (Fig. 7-C) and type-II collagen (Fig. 7-D), possibly aiding in the regenerative process.

Discussion

It has been hypothesized for over thirty years that a factor within synovium may enhance meniscal regeneration and healing, but the underlying mechanism remains unclear. Kim and Moon were among the first to document this association when, in 1979, they showed that meniscectomized rabbit knees subjected to complete synovectomy failed to regenerate meniscal tissue to the same extent as knees with intact synovium³⁶. More recently, investigators have demonstrated improved healing rates in meniscal tears supplemented with synovial flaps or grafts in animal models^{37,38} and synovial rasping in human clinical studies^{39,40}.

The discovery of mesenchymal stem cells in mature synovial tissue with exceptional potential for proliferation and chondrogenic differentiation suggests that these cells may be the source of regenerative stimulus within synovium^{17,18,20,22}. The results of this study provide more convincing evidence that synovial mesenchymal stem cells are capable of stimulating meniscal repair and regeneration.

While the quantity of regenerated tissue was significantly greater in knees supplemented with synovial mesenchymal stem cells at four and twelve-week end points, the defects filled relatively well with tissue by twenty-four weeks in control knees. This may be seen as a shortcoming of the experimental model; however, one must consider not only quantity but also quality of the regenerated tissue. Significantly inferior tissue-quality scores seen in control knees at the twenty-four-week end point suggest that these defects ultimately filled with hypocellular, poorly bonded, fibrous scar tissue as opposed to meniscal fibrocartilage. In contrast, meniscal defects treated with synovial mesenchymal stem cells regenerated with well-bonded tissue closely resembling normal meniscus in cellularity and matrix composition.

The results seen in control knees at twenty-four weeks may be analogous to filling of articular cartilage defects with fibrous tissue, as opposed to true regeneration with hyaline cartilage, after failed articular cartilage restoration procedures⁴¹.

Implanted mesenchymal stem cells found within the meniscal defects resemble native meniscal fibrochondrocytes not only in histological appearance but also in the types of collagen they produce. Meniscal fibrochondrocytes, articular chondrocytes, and fibroblasts, as well as the matrices these cells synthesize, can be distinguished by collagen typing⁴². The primary collagen in meniscal fibrocartilage is type-I collagen, but a substantial amount of type-II collagen is present as well^{42,43}. In contrast, hyaline cartilage is composed of primarily type-II collagen, and fibrous scar tissue is composed of largely type-I collagen^{42,44}. The implanted synovial mesenchymal stem cells in this experiment produced both type-I collagen and type-II collagen, suggesting differentiation into fibrochondrocytes and synthesis of a fibrocartilage matrix.

Much of the recent work surrounding meniscal regeneration has focused on the use of an active biological component added to a tissue scaffold or carrier compound, which functions to contain the biological component within the injured area^{30-32,45}. In contrast to prior studies, the improvements in meniscal regeneration in the current model were achieved without the use of additives designed to contain the implanted cells within the defect. Both fluorescently (CM-DiI) labeled and GFP-positive synovial mesenchymal stem cells preferentially adhered to meniscal defects, where they remained for up to twenty-four weeks. On the basis of the work by Koga et al., this adhesion appears to occur via local surface adhesion molecules, including ICAM-1 (intercellular adhesion molecule-1) and VCAM-1 (vascular cell adhesion molecule-1)⁴⁶. In addition, the attachment takes place rapidly, with >60% of synovial mesenchymal stem cells adhering to injured cartilage within ten minutes of application in an ex vivo human cartilage defect model⁴⁶. Thus, the cells themselves seem to have the inherent ability to quickly adhere to the defect and facilitate the regenerative process.

In an effort to determine the optimal source of mesenchymal stem cells for purposes of intra-articular regeneration, prior studies have compared the in vivo regenerative potential of mesenchymal stem cells obtained from a variety of tissue sources^{23,34}. Koga et al. found that mesenchymal stem cells derived from bone marrow and synovium have greater in vivo chondrogenic regenerative potential than mesenchymal stem cells derived from adipose and muscle³⁴. The chondrogenic and meniscal regenerative potential of mesenchymal stem cells obtained from synovium and bone marrow, however, appears to be more similar^{23,34}. Our rationale for selecting synovial mesenchymal stem cells was based largely on in vitro evidence of superior capacity of synovial mesenchymal stem cells for rapid proliferation and synthesis of chondrogenic matrix in rat²⁰, rabbit³⁴, and human studies¹⁸. The current experiment confirmed the superior in vitro proliferation capacity of rabbit synovial mesenchymal stem cells compared with rabbit bone marrow-derived mesenchymal stem cells. This experiment did not, however, directly compare the regenerative potential of

synovial mesenchymal stem cells and bone marrow-derived mesenchymal stem cells or other additional cell types *in vivo*.

Other approaches for addressing meniscal deficiency and irreparable meniscal tears in humans are under investigation, but an ideal solution has yet to be discovered. Meniscal allograft transplantation, first described in 1989 by Milachowski et al.⁴⁷, has shown promising results in select patients⁴⁸, but is complicated by concerns regarding processing and storage⁴⁹, subclinical host immune response⁵⁰, and a low tolerance for graft-recipient size mismatch⁵¹. The collagen meniscal implant (Collagen Meniscus Implant; ReGen Biologics, Hackensack, New Jersey), a tissue-engineered biologic scaffold fabricated from bovine type-I collagen, has also demonstrated encouraging results compared with partial meniscectomy⁴⁵; however, defect filling is incomplete at second-look arthroscopy⁴⁵ and concerns exist regarding the risk of immune reaction to bovine collagen⁵².

Furthermore, the above treatment options are designed to treat complete or partial meniscal defects, but neither provides a means to stimulate healing of avascular zone tears. *In vitro* meniscal tissue-culture experiments offer evidence that proregenerative cytokines such as transforming growth factor- β (TGF- β), insulin-like growth factor-1 (IGF-1), platelet-derived growth factor (PDGF), and vascular endothelial growth factor (VEGF) may potentially provide such a healing stimulus by increasing meniscal cell proliferation and collagen synthesis^{53,54}. *In vivo* evidence in support of the efficacy of these growth factors in meniscal healing is currently lacking, however. For example, the addition of VEGF coating to meniscal suture failed to increase angiogenesis or improve healing compared with conventional suture in avascular meniscal tears in sheep⁵⁵.

In the present study, we chose to evaluate the regenerative potential of synovial mesenchymal stem cells without scaffolds or additional growth factors. The ability of synovial mesenchymal stem cells to adhere independently to the site of meniscal injury, differentiate into fibrochondrocytes, and synthesize a new matrix that closely resembles native meniscal fibrocartilage without a scaffold or extrinsic cytokines seems to negate the need for such additional stimulus. In addition, this approach avoids the potential for complications associated with disease transmission and immune reaction and, in this model, results in virtually complete regeneration of the meniscal defect. It is possible, however, that meniscal regeneration stimulated by supplementation with synovial mesenchymal stem cells may be further enhanced by the addition of a tissue scaffold when used for large meniscal defects in subjects with little inherent regenerative capacity.


Limitations of this study include those that are inherent to the animal model and those that result from the experimental design. Although the capacity for meniscal regeneration is lower in rabbits than rats, it is likely greater than that in humans^{27,33}. In light of this, caution is encouraged when extrapolating the results of this study to humans.

With regard to the experimental design, the results of this study portray a simplified picture of the events occurring at a molecular level, as the present study did not evaluate the influence of synovial mesenchymal stem cells on the production

or local concentration of proregenerative cytokines such as TGF- β , IGF-1, or PDGF, which promote proteoglycan synthesis and increase cellularity within the meniscus^{53,54}. It is likely that the regenerative process involves a complex interplay between the intact meniscus adjacent to the defect, the implanted synovial mesenchymal stem cells, and alterations in local cytokine concentrations, and this was not fully elucidated in the present study. Finally, although the degree of tissue bonding was considered as a component of the tissue quality score, this experiment did not include a precise quantitative evaluation of the integration between regenerated and native meniscal tissue. Nor did it evaluate the biomechanical characteristics of the regenerated tissue, which are critical to meniscal function. Such an evaluation would be more appropriate in a model with a discrete linear tear or larger meniscal defect.

In conclusion, synovial mesenchymal stem cells implanted into the rabbit knee are capable of adhering to sites of meniscal injury, differentiating into type-I and type-II collagen-producing cells with appearance similar to native meniscal fibrochondrocytes, and enhancing both the quality and quantity of regenerated meniscal tissue. In addition to providing a potential explanation for the association between synovial stimulation and meniscal healing, these results may generate further stimulus for exploring the utility of synovial mesenchymal stem cells in the treatment of meniscal injury in large animal models or humans, with potential applications ranging from supplementing suture repair of avascular zone meniscal tears, to stimulating regeneration of large meniscal defects.

Appendix

 A table showing the histological tissue quality scoring system and an expanded Materials and Methods section describing the tissue harvesting and mesenchymal stem cell preparation, *in vitro* differentiation assays, and tissue quality analysis are available with the online version of this article as a data supplement at jbj.org. ■

NOTE: The authors thank Dina Gaupp, Center for Gene Therapy, Tulane Health Sciences Center, for assistance with histological sectioning.

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Update

This article was updated on May 16, 2012, because of a previous error. The legend for Figures 7-A and 7-B that had previously read “Representative macroscopic appearance (Fig. 7-A) and histological sections (Fig. 7-B) of the meniscal defect one day to twelve weeks after the implantation of GFP-positive green fluorescent protein under fluorescence” now reads “Representative macroscopic appearance (Fig. 7-A) and histological sections (Fig. 7-B) of the meniscal defect one day to twelve weeks after the implantation of GFP-positive synovial mesenchymal stem cells under fluorescence.”

Intra-articular injection of human mesenchymal stem cells (MSCs) promote rat meniscal regeneration by being activated to express Indian hedgehog that enhances expression of type II collagen

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ARTICLE INFO

Article history:

Received 24 November 2011

Accepted 20 June 2012

Keywords:

MSCs

Meniscus

Indian hedgehog

Parathyroid hormone-like hormone

Bone morphogenetic protein 2

SUMMARY

Objective: Meniscal regeneration was previously shown to be enhanced by injection of mesenchymal stem/stromal cells (MSCs) but the mode of action of the MSCs was not established. The aim of this study was to define how injection of MSCs enhances meniscal regeneration.

Design: A hemi-meniscectomy model in rats was used. Rat-MSCs (rMSCs) or human-MSCs (hMSCs) were injected into the right knee joint after the surgery, and PBS was injected into the left. The groups were compared macroscopically and histologically at 2, 4, and 8 weeks. The changes in transcription in both human and rat genes were assayed by species-specific microarrays and real-time RT-PCRs.

Results: Although the number of hMSCs decreased with time, hMSCs enhanced meniscal regeneration in a manner similar to rMSCs. hMSCs injection increased expression of rat type II collagen (rat-Col II), and inhibited osteoarthritis progression. The small fraction of hMSCs was activated to express high levels of a series of genes including Indian hedgehog (Ihh), parathyroid hormone-like hormone (PTH1H), and bone morphogenetic protein 2 (BMP2). The presence of hMSCs triggered the subsequent expression of rat-Col II. An antagonist of hedgehog signaling inhibited the expression of rat-Col II and an agonist increased expression of rat-Col II in the absence of hMSCs.

Conclusions: Despite rapid reduction in cell numbers, intra-articular injected hMSCs were activated to express Ihh, PTH1H, and BMP2 and contributed to meniscal regeneration. The hedgehog signaling was essential in enhancing the expression of rat-Col II, but several other factors provided by the hMSCs probably contributed to the repair.

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Introduction

Injuries to the fibrocartilage of the knee meniscus illustrate the limited ability of human cartilage to repair itself¹. For symptomatic injuries to the meniscus, a meniscectomy is often performed; a procedure which rapidly leads to osteoarthritis². Indications for surgical meniscal repair are limited³, and the results are not always satisfactory. One potential strategy to enhance meniscal regeneration is to use mesenchymal stem/progenitor cells that can differentiate into cartilage and other skeletal cells^{4,5}. A number of reports

demonstrated that healing of the surgically injured meniscus was enhanced by injection of the adult stem/progenitor cells from bone marrow referred to as mesenchymal stem or stromal cells (MSCs)^{6–8}. The observations were initially interpreted on the assumption that the MSCs repaired the tissue by engrafting and differentiating into chondrocytes. However, the number of cells that engrafted was apparently too low to account for the extensive repair frequently observed⁹. Similar observations were made with the use of MSCs in other models for human diseases in which repair of tissues was observed without significant long term engraftment of the cells^{9,10}. Therefore, the recent focus has been the repair of tissues by MSCs through paracrine factors the cells synthesize in culture^{9,11} and the additional therapeutic factors they are activated to express as a result of cross-talk with injured cells and tissues¹⁰.

In the present study, we hypothesized that intra-articular injection of human-MSCs (hMSCs) would enhance meniscal regeneration in rats after hemi-meniscectomy. We then capitalized

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on the species differences to simultaneously track changes in the hMSCs and the endogenous rat cells as the tissue was repaired in order to define how intra-articular injection of MSCs enhances meniscal regeneration.

Materials and methods

Animals

The rat studies were approved by the Institutional Animal Care and Use Committee of Texas A&M Health Science Center. Twelve-week-old male Lewis rats (LEW/Crl; Charles River Laboratories; Wilmington, MA) were used in all experiments with $n = 5$ in each group for quantification of area and histology, and $n = 4$ in each group for RT-PCRs.

Cell preparations

Passage 3 cells were used for all assays. Frozen passage 1 bone marrow-derived human-MSCs (hMSCs) were obtained from the Center for the Preparation and Distribution of Adult Stem Cells (<http://medicine.tamhsc.edu/irm/msc-distribution.html>), and were expanded as described previously¹². Bone marrow-derived rat-MSCs (rMSCs) were isolated from 12-week-old Lewis rats and expanded as described previously¹³.

To label the cells for some experiments, CM-Dil (Invitrogen; Carlsbad, CA) was added at $5 \mu\text{L}/1 \times 10^6$ cells/ml in phosphate buffered saline (PBS). The cells were incubated for 20 min at 37°C in 5% humidified CO₂, centrifuged at $480 \times g$ for 5 min, and washed twice with PBS.

Meniscectomy and cell injection

Under isoflurane anesthesia a hemi-meniscectomy was performed on both knees as described previously⁷. Immediately after surgery, 2×10^6 CM-Dil labeled hMSCs or rMSCs in 50 μL PBS were injected intra-articularly into the right knee joint. The left knee joint received 50 μL PBS as the control. The rats were allowed to walk freely in the cage.

For experiments with cyclopamine, which is an inhibitor of hedgehog signaling, 2×10^6 hMSCs were injected postoperatively on day 0, and cyclopamine (50 μL of 20 μM in 0.1% DMSO/PBS; Sigma–Aldrich) was injected intra-articularly on postoperative days 0, 1, and 3. Controls were injected with vehicle alone at the same time points. For experiments with Smoothed agonist (SAG), we injected SAG solution (50 μL of 100 nM or 500 nM in PBS; Santa Cruz) intra-articularly on postoperative day 0. We injected cyclopamine, SAG, or vehicle alone (control) bilaterally into knees of each rat ($n = 4$ knees in each group).

Quantification of meniscus area

The whole medial meniscus ($n = 5$ each time point) was collected at 2, 4, and 8 weeks after injection. The menisci were photographed and the area measured with Scion image software.

Histology

After photography for area measurements, menisci were fixed in 4% paraformaldehyde and embedded in paraffin. Five μm horizontal sections were cut and stained with Toluidine blue or subjected to immunohistochemistry.

To assess osteoarthritic changes, proximal tibias were dissected at 2, 4, and 8 weeks after injection and fixed in 4% paraformaldehyde. They were then decalcified (Decalcifying Solution;

Richard-Allan Scientific; Kalamazoo, MI) at 4°C for 14 days and embedded in paraffin. The sections stained with Safranin-O and fast green to visualize the cartilage. Two examiners blinded to treatments scored the sections using the Osteoarthritis Research Society International (OARSI) osteoarthritic cartilage histopathology grading system¹⁴. The scoring system ranges from 0 to 24, with 0 representing normal and 24 representing the most severe osteoarthritis.

Immunohistochemistry

Rat type II collagen was visualized using diaminobenzidine (DAB) staining of a mouse monoclonal antibody (clone II-4C11) from MP Biomedicals. For immunofluorescence, primary antibodies to Indian hedgehog (Ihh) (ab52919), bone morphogenetic protein 2 (BMP2) (ab6285), parathyroid hormone-like hormone (PTH-LH) (ab55631), and parathyroid hormone receptor 1 (PTH-R1) (ab75150) were obtained from Abcam. Species-specific AlexFluor-488 conjugated secondary antibodies were from Invitrogen.

Real-time RT-PCR analysis for selected mRNAs

RNA was isolated from regenerated menisci using RNA Bee (Tel-Test; Friendswood, TX) according to manufacturer's instructions and cleaned by RNeasy Mini Kit (QIAGEN; Valencia, CA). Total RNA was used in one-step real-time RT-PCR (ABI 7900 Sequence Detector, Applied Biosystems; Foster City, CA) using QuantiTect Probe RT-PCR Kit (QIAGEN). Reactions were incubated at 50°C for 30 min, 95°C for 15 min, then 40 cycles at 95°C for 15 s followed by 60°C for 1 min.

For normalization of gene expression, human-specific GAPDH primers and probe (TaqMan Gene Expression Assays ID, Hs99999905_m1), rat-specific GAPDH primers and probe (TaqMan Gene Expression Assays ID, Rn01775763_g1), or 18S rRNA primers and probe (TaqMan Gene Expression Assays, ID Hs03003631_g1) were used as internal controls (all from Applied Biosystems; Supplemental Table 3). All primers were tested for species-specificity; the values for critical threshold were indistinguishable from background with RNA from the alternative species.

Statistical analysis

For comparison of two groups, the Mann–Whitney *U* test was used. For multiple comparisons of three groups or more than three groups, one-way ANOVA followed by Bonferroni's *post hoc* test was used. *P* values less than 0.05 were considered to be statistically significant. Data were analyzed using StatView software.

Additional methods

Detailed methods for cell culture, surgical procedures, immunohistochemistry and microarrays are described in the Supplemental Methods.

Results

Intra-articular injection of hMSCs promoted meniscal regeneration and inhibited development of osteoarthritis

To determine whether intra-articular injection of hMSCs would enhance regeneration of rat meniscus, we performed hemi-meniscectomy⁷ in both knees of wild-type Lewis rats. We then closed the knee joint and immediately injected CM-Dil labeled hMSCs (2×10^6 cells in 50 μL PBS) into the right knee joint and an equal volume of PBS into the left knee. For positive controls for the

xenotransplantation, we repeated the experiments with the same number of rMSCs. At 2 and 4 weeks, injection of the hMSCs enhanced regeneration as indicated by the morphology of the meniscus [Fig. 1(A)] and the size of regenerated area [Fig. 1(B)]. At 8 weeks the beneficial effects of the hMSCs were apparent in

histological sections stained with Toluidine blue for metachromasia or labeled with antibodies to type II collagen [Fig. 1(C)]. At 8 weeks the absolute area of regeneration was not significantly different between PBS and hMSC menisci because of the innate ability of rat meniscus to regenerate itself [Fig. 1(B)]. Of special note was that

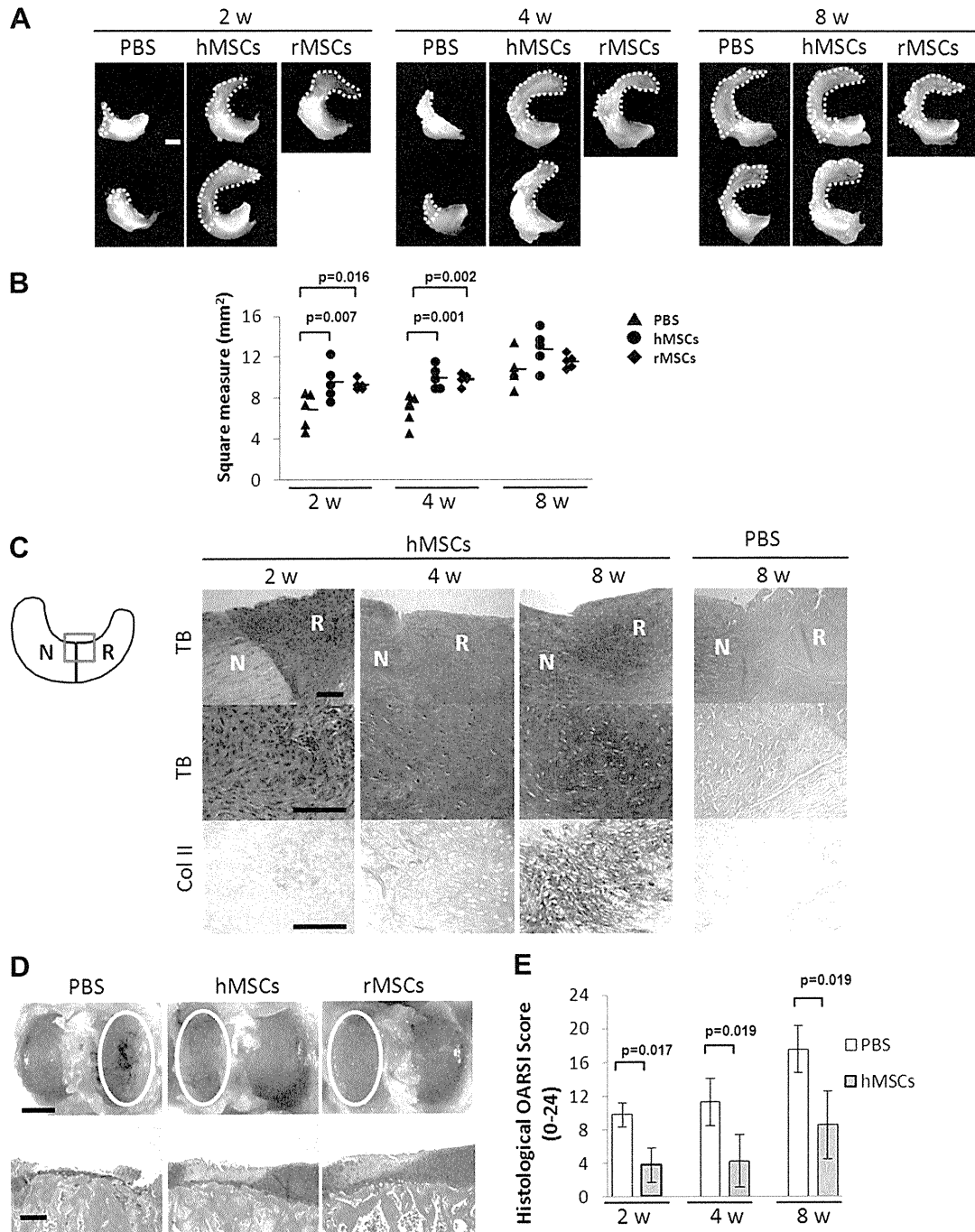


Fig. 1. Intra-articular injection of hMSC promoted regeneration of rat meniscus and inhibited development of osteoarthritis after the meniscectomy. (A) Representative macroscopic findings of the meniscus 2, 4, and 8 weeks after the injection of PBS (left), hMSCs (middle), or rMSCs (right). The white dotted line indicates the regenerated tissue. Scale bar, 1 mm. (B) Area of the total meniscus injected with PBS, hMSCs, or rMSCs at 2, 4, and 8 weeks. The horizontal lines are mean values; $n = 5$ for each group (one-way ANOVA followed by Bonferroni post-tests. P values less than 0.017 were considered to be statistically significant). (C) Representative sections of the meniscus stained with Toluidine blue (top and middle), and immunostained for type II collagen (bottom) after PBS or hMSCs injection. The staining in the PBS-treated sample was less with Toluidine blue and the antibody for type II collagen. The schema of the meniscus on the left is shown for orientation. Symbols: N, native meniscus; R, regenerated meniscus; TB, Toluidine blue; Col II, type II collagen. Scale bar, 100 μ m. (D) Representative gross photographs (top) and sections (bottom) of the joint surface of the tibia at 8 weeks. The cartilage was stained with India ink to identify fibrillation and erosion. The white circle indicates the medial tibial plateau. The tibia was sectioned coronally and stained with safranin-O and fast green to identify cartilage (red). Scale bars, 2 mm (top) or 200 μ m (bottom). (E) Quantification of histological analysis using the OARSI cartilage osteoarthritis histopathology grading system. Values are mean with lower and upper limit of 95% CI; $n = 5$ for each group (Mann–Whitney U test).

there was no significant difference between the ability of xenotransplantation of hMSCs and syngeneic transplantation of rMSCs to enhance regeneration [Fig. 1(B)].

We next evaluated articular cartilage damage in adjacent femoral and tibial articular surfaces. Both articular surfaces had osteoarthritic change but they were more severe on the medial tibial plateau than on the medial femoral condyle (data not shown). Therefore, we quantified the osteoarthritic change in the tibial plateau. At 8 weeks after the meniscectomy, severe degenerative changes were observed in the medial tibial plateau in PBS knees, but hMSCs inhibited the osteoarthritis progression after meniscectomy [Fig. 1(D and E)].

Detection of the injected hMSCs

CM-Dil + hMSCs were readily detected in histological sections of the regenerated meniscus but it was apparent that they decreased rapidly in number with time (Supplemental Fig. 1). The labeled cells seen at 2 and 4 weeks had the morphology of fibroblasts but the small number of CM-Dil-labeled cells seen at 8 weeks had the more spherical morphology of chondrocytes (Supplemental Fig. 1). We confirmed the persistence of the hMSCs by labeling the sections with human-specific nuclear antibody (Supplemental Fig. 2). For a quantitative measure of the engrafted human cells^{15,16}, we prepared standard curves by adding predetermined numbers of hMSCs to the surgically injured menisci just before homogenization, extracted RNA, and assayed it with a quantitative RT-PCR assay specific for human GAPDH [Supplemental Fig. 3(A–C)]. Approximately 2% of hMSCs injected into the synovial space were recovered in the regenerating tissue at day 1, and 1% of hMSCs were recovered at day 3. No hMSCs were detected at day 7 with the assay with the RT-PCR for human GAPDH [Supplemental Fig. 3(B)]. The standard curve [Supplemental Fig. 3(A)] suggested that the assay was sensitive to the detection of 200–1000 hMSCs. However, hMSCs were still detected by immunohistochemistry at 8 weeks (below). The discrepancy is probably explained by the standard curve over-estimating the sensitivity of the assay, since it was generated by simply adding hMSCs to the tissue before extraction of the RNA; efficient extraction of RNA from a cartilaginous tissue such as the meniscus that is rich in proteoglycans usually requires digestion of the matrix and isolation of the cells¹⁷. Therefore the real-time RT-PCR assay reflected the rapid decrease in hMSCs in the first few days after administration but probably under-estimated the cells remaining in the tissue at later times.

The injected hMSCs were activated to express *Ihh*, *PTH1LH*, and *BMP2*

The use of a xenotransplantation made it possible to follow simultaneously changes in both the donor hMSCs and the endogenous rat cells in the regenerating meniscus. A total of 2×10^6 hMSCs were injected into the hemi-meniscectomized knee joint of rat, and RNA was extracted from the meniscus 3 days after the injection, a time at which assays for human GAPDH mRNA indicated there were adequate amounts of human mRNA for analyses. As we observed previously^{15,18}, an initial impression of the potential cross-talk between the hMSCs and host rodent cells can be obtained by assaying the same RNA on species-specific microarrays.

Therefore, the same samples of RNA were assayed on human microarrays and on rat microarrays, and the data were filtered for cross-hybridization (see Supplemental Methods). The data (Supplemental Table 1) indicated that the human transcriptome was changed with up-regulation of 93 human transcripts. Also, the data suggested that a series of human genes were down-regulated, but the number could not be accurately defined because signals for

many human genes were eliminated from the analysis by filtering for cross-hybridization in the samples which contained a low ratio of human RNA to rat RNA.

The 40 most highly up-regulated human transcripts (Supplemental Table 1) were subjectively examined for candidate genes of interest, and human-specific real-time RT-PCR assays were used to confirm the microarray data [Fig. 2(A)]. Because administration of hMSCs enhanced regeneration of the meniscus, we elected to focus on the genes related to growth factors, such as PDGF, VEGF, FGF, TGF- β , BMP2 and PTHLH. As observed previously¹⁹, the RT-PCR assays indicated that the increase in expression of some of the genes was larger than the values obtained from the microarrays. On day 3 the increase in PTHLH was 443-fold and the increase in BMP2 was 194-fold. The microarray data did not reveal any changes in *Ihh*, but *Ihh* was observed to up-regulate BMP2 and PTHLH during the endochondral ossification^{20,21}. Therefore we assayed for *Ihh* expression. Surprisingly, human *Ihh* was also increased 22-fold at day 3 compared to control hMSCs. Expression of human *Ihh* not detected on day 7, apparently because only a small number of hMSCs remained in the tissue [Supplemental Fig. 3(B)].

We next examined the time sequence of changes in the human transcriptome [Fig. 2(B)]. The gene expression of human PTHLH increased 516-fold as early as day 1. The level per human cell (normalized by Δ Ct for specific for human GAPDH) increased further between day 3 and day 7 (data not shown), but because the RNAs we have isolated included both human and rat RNAs and the number of human cells was rapidly decreasing, the total level of transcript were normalized again by 18S rRNA, which contains sequences that are 100% conserved among all the eukaryotes. The total level of transcript of PTHLH (normalized by Δ Ct for eukaryotic 18S rRNA) was lower on day 7 [Fig. 2(B)]. BMP2 also increased 663-fold at day 1, and further increased at day 7 if expressed on a per human cell basis but not as total tissue content (data not shown).

Immunohistochemistry confirmed that the injected hMSCs that adhered to the injured site expressed *Ihh*, PTHLH, and BMP2 [Fig. 2(C–E)].

hMSCs up-regulated expression of chondrogenic genes in rat cells

The data from the rat microarrays indicated that there was up-regulation of expression of 91 rat genes and down regulation of 42 rat genes by a factor of 2-fold or more. The most highly up-regulated 40 rat genes are shown in Supplemental Table 2.

Assays by rat-specific real-time RT-PCR were carried out for transcripts of genes expressed during chondrogenesis. Injection of the hMSCs increased the levels of Sox9 mRNA at day 3 and day 7 (Fig. 3). As expected from the assays by immunohistochemistry [see Fig. 1(C)], injection of the hMSCs increased expression of the rat Col2a1 gene at day 7 and 14. The increases in several other rat genes were more variable. There was no consistent effect of the hMSCs on expression of rat *Ihh*, or the rat BMP receptor BMPRI1A. However, hMSC injection increased the expression of rat receptor for PTHLH (PTH-R1) and BMP2 on day 3, and PTHLH on day 14. The increased expression of PTHLH and PTH-R1 was evaluated by immunohistochemistry (Fig. 4). Interestingly, not only human (CM-Dil positive) cells but also rat (CM-Dil negative) cells surrounding hMSCs expressed PTHLH and PTH-R1 at day 7.

The role of human *Ihh*

We elected to test the hypothesis that activation of hMSCs to express *Ihh* was an essential step in enhancing the regeneration of the rat meniscus. To test the hypothesis, we used cyclopamine, a steroid alkaloid that inhibits *Ihh* signaling by serving as an antagonist to the downstream target Smoothened. We elected to

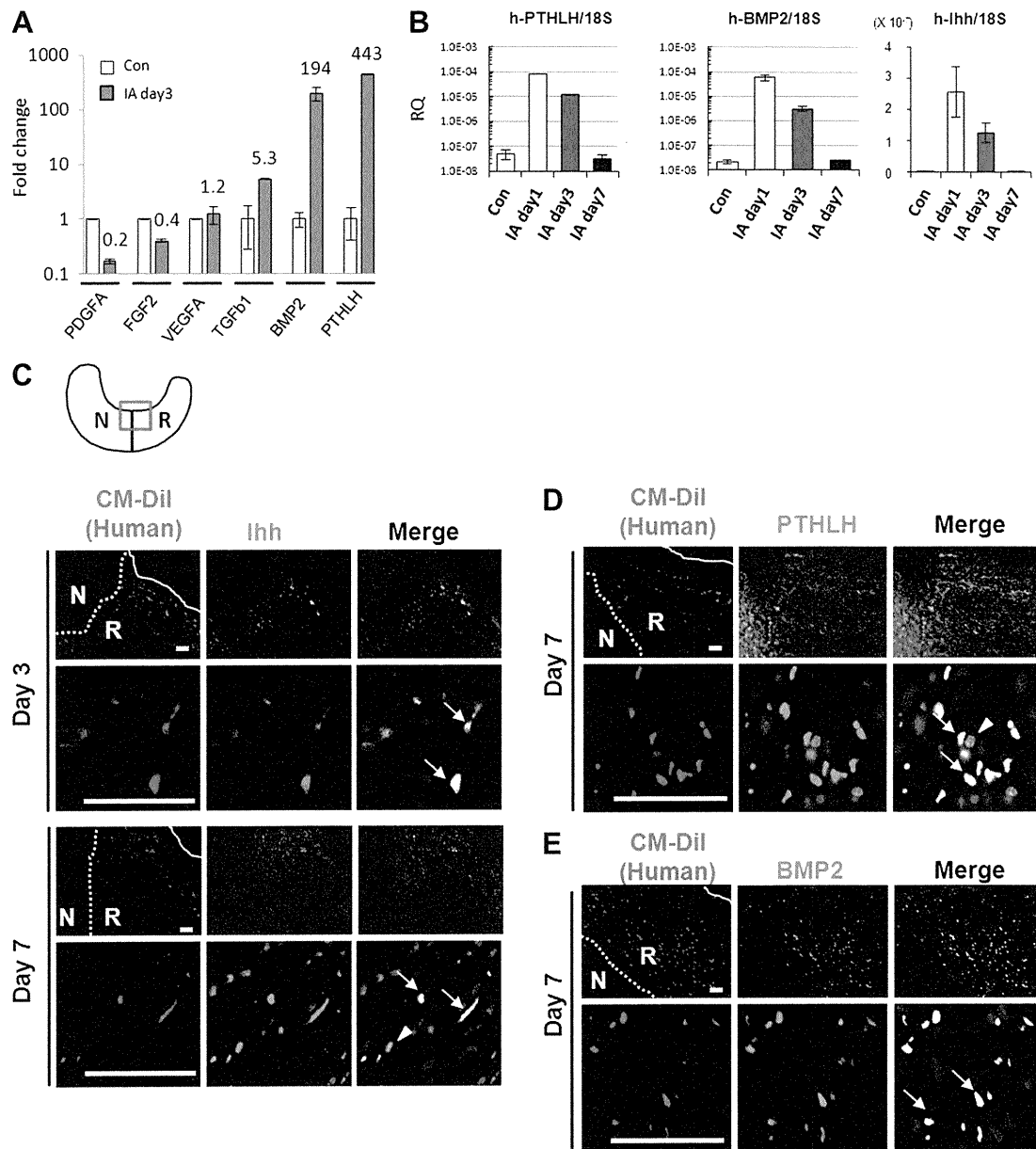


Fig. 2. Intra-articular injected hMSCs were activated to express *Ihh*, *PTHLH*, and *BMP2*. (A) Real-time RT-PCR for human-specific mRNA in regenerated meniscus 3 days after injection. Values are mean with lower and upper limit of 95% CI of fold increase over Con, normalized by $\Delta\Delta Ct$ for human-specific GAPDH; $n = 4$. Symbols: Con, 20,000 hMSCs added to injured meniscus from PBS injected knee before RNA extraction; IA day 3, regenerated meniscus 3 days after intra-articular (IA) injection of hMSCs. (B) Time sequence of human gene expression by real-time RT-PCR for human *PTHLH*, *BMP2*, and *Ihh* in regenerated meniscus. Values are mean with lower and upper limit of 95% CI of the relative quantities (RQ) normalized to 18S rRNA reflecting total levels of both human and rat transcripts. (C–E) Immunohistochemistry of anti-human/rat *Ihh* (C), *PTHLH* (D), or *BMP2* (E) 3 or 7 days after hMSC injection. Primary antibodies for *Ihh* and *PTHLH* reacting with both human and rat were used. Low magnification (top) and higher magnification (bottom) of regenerated meniscus. Right column: merged images (CM-Dil, red; *Ihh*/*PTHLH*/*BMP2*, green; DAPI, blue). The white dotted line indicates border between native and regenerated meniscus, and the white solid line indicates outer edge of regenerated meniscus. CM-Dil labeled hMSCs (red) expressed *Ihh*, *PTHLH*, and *BMP2* (green) (arrows in merge). Note: CM-Dil negative rat-derived cells surrounding hMSCs also expressed *Ihh* and *PTHLH* (arrowhead in merge) at day 7. The schema of the meniscus on the top left (C) is shown for orientation. Symbols: N, native meniscus; R, regenerated meniscus. Scale bars, 50 μ m.

inject cyclopamine on day 0, 1, and 3 because it had a half-life of only 4 h after ip or po administration²² and a similar dosage schedule was used previously in experiments on repair of a cranial defect²³. In the presence of cyclopamine, hMSCs did not enhance expression of rat *Col2a1* [Fig. 5(A)]. Therefore the results were consistent with the hypothesis that expression of *Ihh* was essential. As expected, administration of cyclopamine also decreased expression of *Patched* (*Ptc1*) and *Gli1*, downstream targets of hedgehog signaling [Fig. 5(B and C)]. However, cyclopamine did not inhibit morphological regeneration of the meniscus as fibrous tissue [Fig. 5(D and E)]. As a further test of the hypothesis we used

SAG, a Smoothed agonist of hedgehog signaling. We elected to inject SAG on day 0 because expression of human *Ihh* reached a peak on day 1 and then declined [Fig. 2(B)].

Injection of SAG after the meniscal surgery enhanced expression of rat *Ptc1*, *Gli1*, and *Col2a1* gene in the absence of hMSCs [Fig. 5(A–C)]. However, injection of SAG did not increase the area of repair [Fig. 5(D and E)]. The results suggested that expression of *Ihh* by the hMSCs was essential for enhancing the rat-*Col II* expression but probably not in itself sufficient for the enhanced regeneration observed with administration of hMSCs in the model (Supplemental Fig. 4).

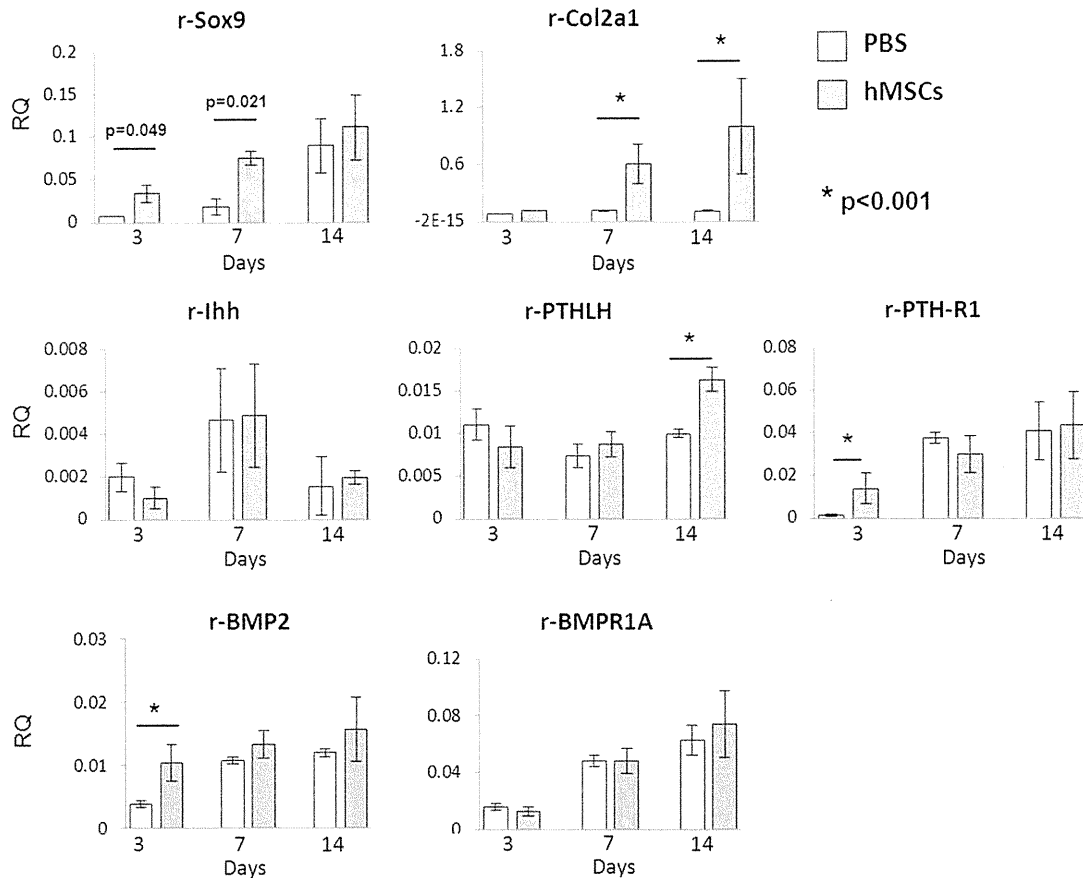


Fig. 3. hMSCs up-regulated expression of chondrogenic genes in rat cells. Real-time RT-PCR assay for rat-specific mRNA (Sox9, Col2a1, Ihh, PTHLH, PTH-R1, BMP2, and BMPR1A) in the regenerated meniscus 3, 7 and 14 days after PBS or hMSC injection. Values are mean with lower and upper limit of 95% CI of the RQs normalized to rat-specific GAPDH; $n = 4$ for each group. * $P < 0.001$ (Mann–Whitney U test).

Discussion

As reported previously^{6–8}, injection of MSCs enhanced the regeneration of surgically injured meniscus. Of special interest was the observation here that xenotransplantation of hMSCs was as effective as syngeneic rMSCs in repairing the rat meniscus⁷. The effectiveness of the hMSCs in the immunocompetent rats is consistent with previous reports that MSCs were immune privileged^{24–26} and that hMSCs had about the same half-life after injection into the hippocampus of wild-type mice as after injection into the hippocampus of immunodeficient mice¹⁸. The hMSCs promoted both the synthesis of rat type II collagen in the regenerating meniscus and apparently the resilience of the tissue since they reduced the tibial osteoarthritis that developed in control knees. The hMSCs employed here may have been particularly effective because they were prepared with a standardized protocol that generates cultures enriched for early progenitor cells^{12,27,28}.

We previously evaluated the fate of rMSCs by *in vivo* imaging and histology in the same rat hemi-menisectomy model⁷. *In vivo* imaging analysis demonstrated that more than 1000 of intra-articular injected rMSCs were detected in the knee joints for 28 days. The number of intra-articular injected LacZ+ rat cells were also decreased with time, but small number of LacZ+ cells were observed in the regenerated meniscus up to 12 weeks. In the present study, we did not directly compare the number of injected cells in rMSCs vs hMSCs; however apparently the rat cells survived somewhat longer than human cells⁷.

The use of xenotransplantation of hMSCs made it possible to use species-specific assays obtain an initial impression of cross-talk

between the hMSCs and the host cells in the meniscus^{15,18}. Real-time RT-PCR assays of mRNA for human-specific GAPDH demonstrated that only a small number of the hMSCs injected into the synovial space engrafted to the regenerating edge of the meniscus. The cells disappeared rapidly but immunohistochemistry of the tissue demonstrated that a small number were detectable at 8 weeks. Microarray data demonstrated that shortly after the hMSCs had engrafted, there were marked changes in their transcriptomes with up-regulation of 93 human genes. Therefore, exposure of the hMSCs to the *in vivo* microenvironment had a major effect on the human cells. Real-time RT-PCR assays to confirm some of the microarray data demonstrated several hundred fold increases in the levels of transcripts for the cartilage characteristic genes PTHLH and BMP2. These increases were accompanied by increases in the expression of human Ihh.

The time course for changes in the transcriptomes of the human cells and the rat cells in the regenerating edge of the meniscus suggested that the activation of the human cells to express Ihh, PTHLH and BMP2 promoted the subsequent expression of the Col2a1 gene by the rat cells (Fig. 6). The results indicated that the expression of Ihh was critical since an inhibitor of hedgehog signaling negated the effects of hMSCs in promoting expression of the rat Col2a1 gene. The inhibitor was not species-specific and therefore inhibited expression of both human and rat Ihh. Unfortunately, several experiments to use siRNAs to knock down the Ihh transcripts in the hMSCs under the conditions used previously¹⁵ were unsuccessful: the transcripts were knocked down with the negative controls with the scrambled siRNA controls to the same extent as with the specific siRNAs. However, the conclusion that

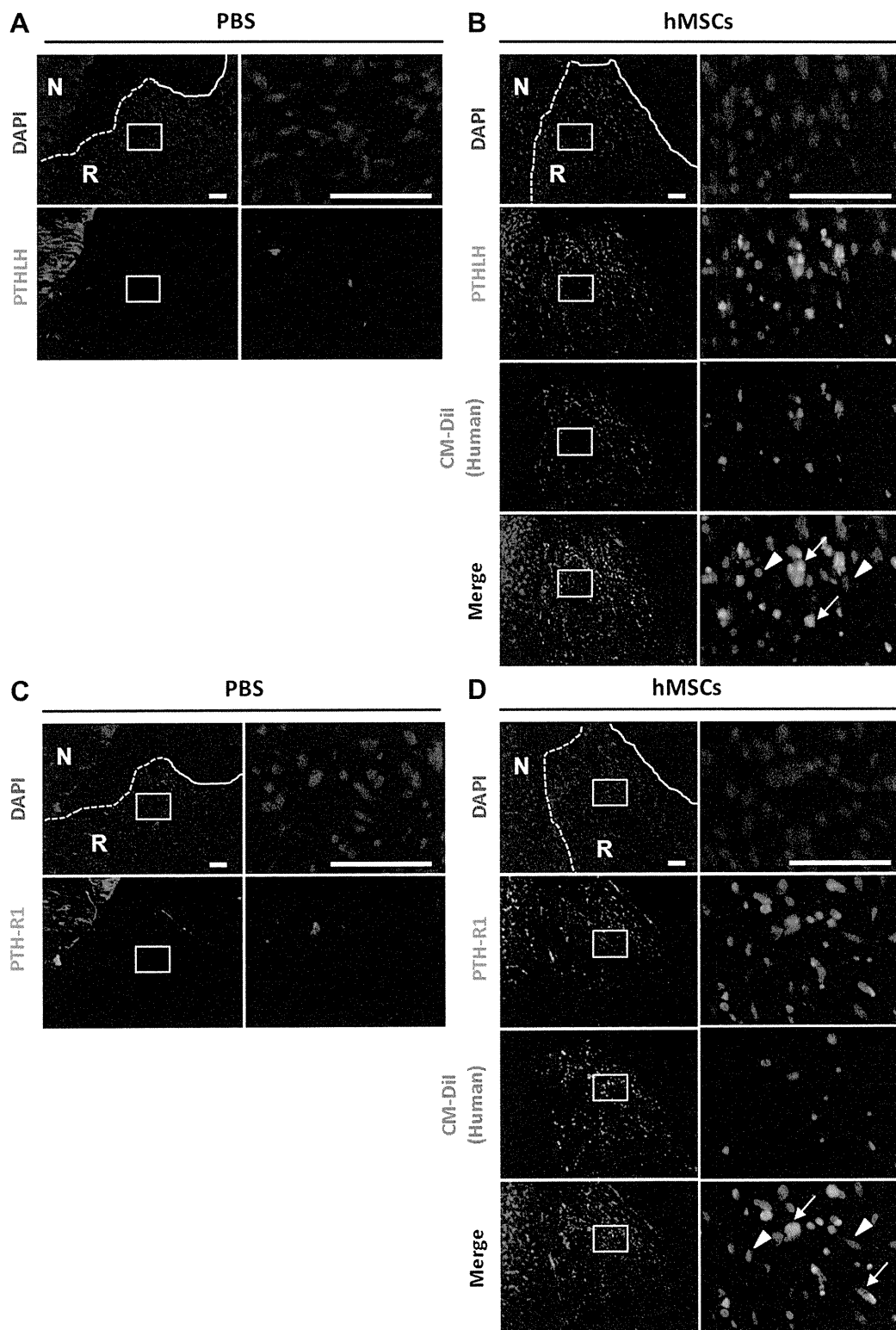


Fig. 4. A large number of PTHLH/PTH-R1 positive cells were found in the regenerated tissue after injection of hMSCs. (A–D) Immunohistochemistry of anti-human/rat PTHLH (A, B) or PTH-R1 (C, D) 7 days after the injection of PBS (A, C) or hMSCs (B, D). Low magnification images are shown in the left and higher magnification of the regenerated tissue is shown in the right. Staining of nuclei (DAPI, blue), CM-Dil (red), and PTHLH or PTH-R1 (green) is shown. The white dotted line indicates the border between the native meniscus and regenerated tissue, and the white solid line indicates the outer edge of the regenerated tissue. A large number of PTHLH or PTH-R1 positive cells (green) were observed in the regenerated meniscus after the injection of hMSCs (B, D). In contrast, few PTHLH or PTH-R1 positive cells were present in the PBS injection group (B, D). Note that not only CM-Dil positive human cells (arrows in merge) but also CM-Dil negative rat cells (arrowheads) expressed PTHLH or PTH-R1.

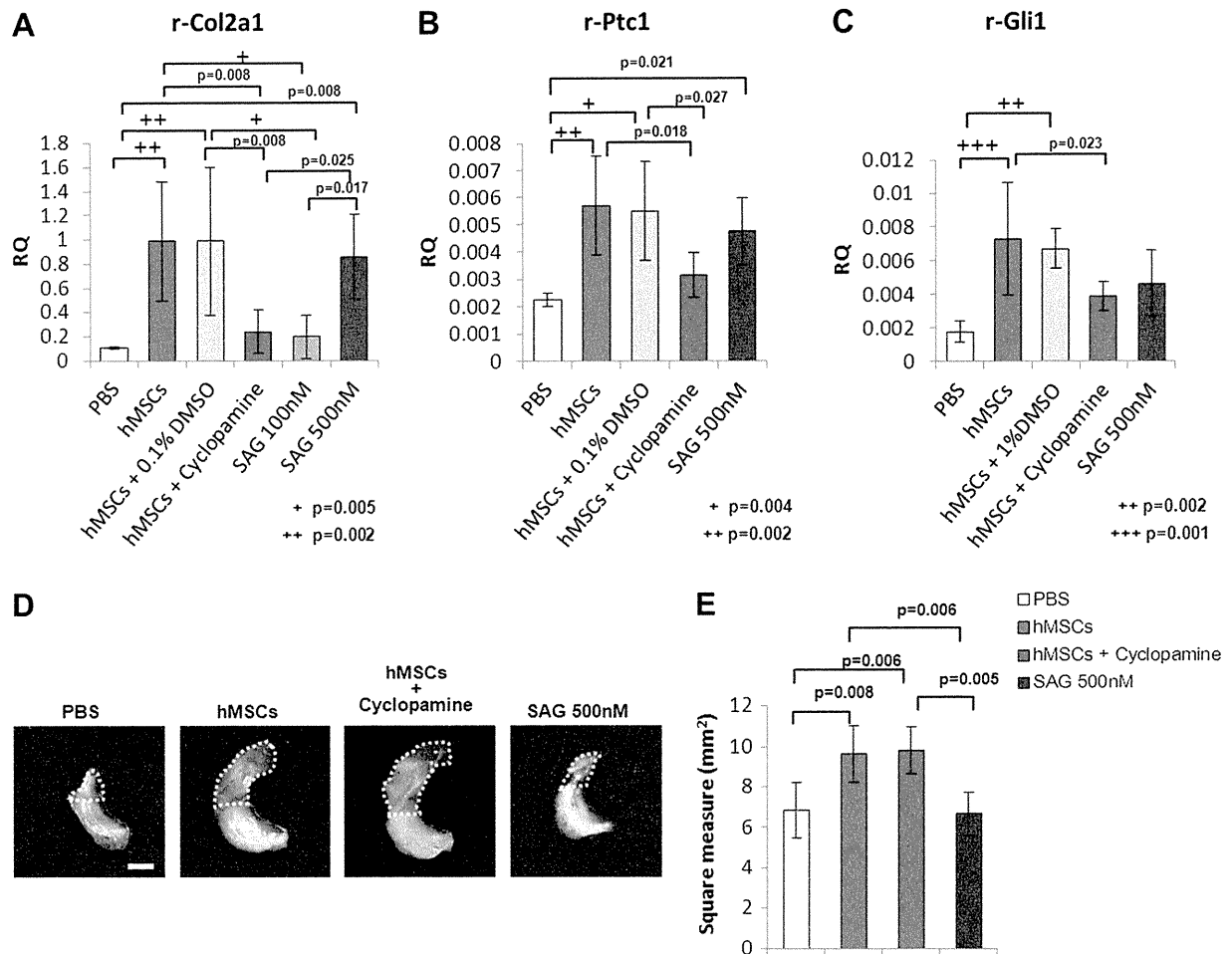


Fig. 5. Effects of an inhibitor and an agonist of hedgehog signaling. (A) Real-time RT-PCR assay for rat-specific *Col2a1* mRNA. RNA was recovered from the regenerated meniscus 2 weeks after the various treatments, including injection of PBS, hMSCs, hMSCs plus 0.1% DMSO (as a control of cyclopamine), hMSCs plus cyclopamine (50 μ L of a 20 μ M in 0.1% DMSO), 100 nM or 500 nM Smoothed agonist (SAG). Values are mean with lower and upper limit of 95% CI of the RQs normalized to rat-specific GAPDH; $n = 4$ for each group. $+P = 0.005$, $++P = 0.002$ (one-way ANOVA followed by Bonferroni post-tests. P values less than 0.0033 were considered to be statistically significant). (B, C) Real-time RT-PCR assay for rat-specific *Ptc1* (B) and *Gli1* (C), downstream targets of hedgehog signaling. RNA was recovered from the regenerated meniscus 3 days after the various treatments. Values are mean with lower and upper limit of 95% CI of the RQs normalized to rat-specific GAPDH; $n = 4$ for each group. $+P = 0.004$, $++P = 0.002$ in B, $++P = 0.002$, $+++P = 0.001$ in C (one-way ANOVA followed by Bonferroni post-tests. P values less than 0.005 were considered to be statistically significant). (D) Representative macroscopic photographs of the meniscus 2 weeks after the various treatments. Scale bar: 1 mm. (E) Area of the total meniscus 2 weeks after the various treatments. Values are mean with lower and upper limit of 95% CI; $n = 4$ for each group (one-way ANOVA followed by Bonferroni post-tests. P values less than 0.0083 were considered to be statistically significant).

expression of *Ihh* by the hMSCs played a critical role was supported by the observation that an agonist of hedgehog signaling prompted expression of the rat *Col2a1* gene in the absence of hMSCs.

The agonist of hedgehog signaling did not however fully duplicate the effects of the hMSCs because it did not enhance restoration of the morphology of the meniscus. Therefore the hMSCs made additional contributions to the regeneration process that probably involved the expression of human PTHLH, BMP2, and perhaps other factors.

The increased expression of the *Col2a1* produced by the agonist of hedgehog signaling raises the possibility that such agonists might provide a therapy for improving cartilage repair. However, the effects of hedgehog signaling are highly dependent on context²⁹. Lin *et al.*³⁰ demonstrated that hedgehog signaling was increased in human osteoarthritic samples and in mice with surgically induced osteoarthritis. Also, transgenic mice over-expressing hedgehog were more predisposed to osteoarthritis than wild-type mice. In addition, they observed that pharmacological and genetic inhibition of hedgehog signaling decreased the development of osteoarthritis. Experiments within cultured

chondrocytes were consistent with these observations. The reasons for the dramatic differences from the observations made here are not apparent. The data however suggest that the relatively brief increase in hedgehog signaling we observed after a single administration of hMSCs immediately after hemi-menisectomy enhanced a process of orderly chondrogenesis much as occurs during normal development of the skeleton. In contrast, the more chronic increases in hedgehog signaling that Lin *et al.*²⁶ examined in the setting of the extensive tissue destruction seen in osteoarthritis apparently drive a disordered hypertrophy of chondrocytes that exacerbates the pathological changes. One consequence of these observations is that agonists of hedgehog signaling may improve meniscal regeneration and chondrogenesis after acute injuries, but chronic administration of the same agents may increase the severity of osteoarthritis³⁰.

The results are consistent with previous observations with MSCs in that increases in expression of *Ihh* and PTHLH were seen during chondrogenic differentiation of MSCs *in vitro*^{31,32}. The results are also consistent with previous observations on the roles of *Ihh*, PTHLH and BMP2 in chondrogenesis. *Ihh* is one of the master

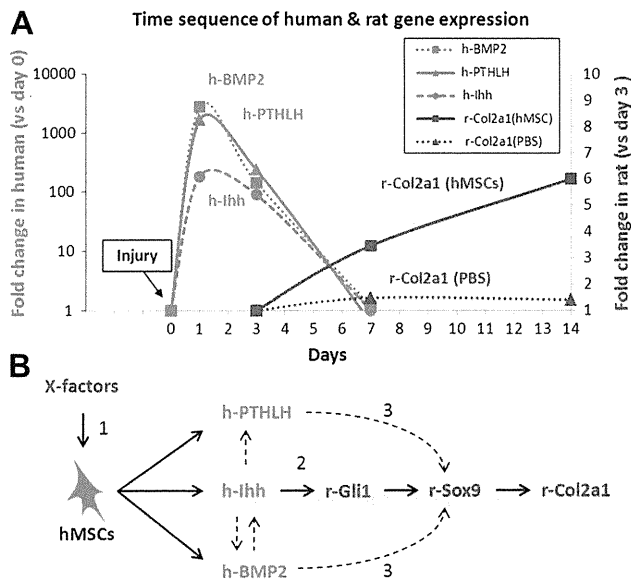


Fig. 6. Possible mechanism of rat meniscal regeneration by intra-articular injection of hMSCs. (A) Summary graph of time sequence of human and rat gene expression changes by species-specific real-time RT-PCR. Intra-articular injected hMSCs were activated to express human Ihh, PTHLH, and BMP2 (red) as early as day 1. Values for human genes are expressed as fold increase over values for controls (Con) of cultured hMSCs, normalized by $\Delta\Delta Ct$ for 18S to reflect total levels of both human and rat transcripts. Rat Col2a1 gene expression in the regenerated meniscus was increased after the injection of hMSCs (blue). Values for rat genes are expressed as fold increase over values for PBS day 3, normalized by $\Delta\Delta Ct$ for rat GAPDH. Values are means from RT-PCR data presented in Figs. 2(B) and 3. Symbols: Con, 20,000 hMSCs added to injured meniscus from PBS injected knee before RNA extraction. (B) Summary diagram of the possible mechanism for rat meniscal regeneration by hMSCs. (1) Intra-articular injected hMSCs were activated to express Ihh, PTHLH, and BMP2 when they were exposed to the *in vivo* microenvironment. (2) Human Ihh expressed by the hMSCs enhanced expression of rat Col2a1 gene probably through the downstream hedgehog target of rat Gli1 and then through Sox9. hMSCs did not enhance expression of rat Gli1 and Col2a1 in the presence of cyclopamine. In contrast, injection of Smoothed agonist enhanced expression of rat Gli1 and Col2a1. (3) PTHLH and BMP2 may also have enhanced expression of rat Col2a1 through rat Sox9. Symbols: X-factors, unknown factors which activate hMSCs to express Ihh, PTHLH, and BMP2.

regulators of both chondrocyte and osteoblast differentiation during endochondral bone formation. Ihh is expressed in mesenchymal cells at the early stage of pre-chondrocytes and the late stage of pre-hypertrophic chondrocytes³³. Ihh stimulates chondrocyte proliferation directly and, through stimulation of PTHLH synthesis, determines the distance from the end of the bone at which chondrocytes stop proliferating and undergo hypertrophic differentiation³⁴. The results are also consistent with previous observations that Gli, a downstream target of hedgehog signaling, can up-regulate Sox9 by binding to its promoter^{35,36}. During endochondral ossification, PTHLH is synthesized by chondrocytes and perichondrial cells at the ends of the developing bones. It stimulates the proliferation of chondrocytes and suppresses their terminal differentiation^{31,37,38}. PTHLH up-regulates Sox9 transcription³⁹, which has been shown to promote the differentiation of mesenchymal cells into chondroblasts⁴⁰. Thus, PTHLH may have acted upon the endogenous rat mesenchymal cells and induced their differentiation through the Sox9 pathway. In addition, PTHLH up-regulates the expression of Bcl2, an apoptosis inhibitor⁴¹, which also may have a role in regulating rat matrix production⁴². BMP signaling has been shown to play an important role in the development of bone and cartilage. BMP2 has been demonstrated to be a potent stimulator of chondrocyte metabolism and differentiation *in vitro*^{43–45} and *in vivo*^{46,47}. For example, BMP2 stimulated DNA synthesis of bovine meniscal cells *in vitro* in a dose-dependent manner⁴⁸. In addition, Ihh is a direct target gene of BMP2⁴⁹, and

Ihh promotes BMP expression. In the injured meniscus, BMP2 produced by hMSCs may have stimulated endogenous rat mesenchymal cells to proliferate and differentiate. It may also have activated Ihh signaling in human and rat cells.

In effect, the observations presented here demonstrated that the hMSCs that engrafted in the knee joint of the hemimiscectomized rat were activated by the microenvironment of the injured meniscus to express extremely high levels of three genes that play critical roles in the development of normal cartilage: Ihh, PTHLH and BMP2. The human cells rapidly decreased in number but their presence triggered the subsequent expression of type II collagen by the rat meniscal cells. The expression of human Ihh was essential for up-regulating rat genes for type II collagen, Patched, and Gli1, an observation that raised the possibility that agonist of Ihh signaling may be useful in treating acute injuries to the meniscus and perhaps articular cartilage. However, for clinical application, interspecies differences have to be considered. The inherent healing capacity of the human meniscus has been shown to be lacking in the inner third and is very limited in the middle third⁵⁰. We used a rat model, and rat meniscus has a greater spontaneous healing potential⁷. To demonstrate the effectiveness of agonist of Ihh signaling for meniscus regeneration, further experimental studies in larger animal models are needed. In addition, our data lack details about biomechanical properties and the contents of collagen and proteoglycans in the regenerated meniscus. Also, it is uncertain whether the regenerated menisci will prevent secondary osteoarthritic change in the long term. Therefore, further studies should be carried out.

Contributions

Study design: MH, IS, TM and DP. Study conduct: DP. Data collection: MH, HC, and JY. Data analysis: MH, HC, and JY. Data interpretation: MH, HC, RL, JY, IS, and DP. Drafting manuscript: MH, HC, and DP. Revising manuscript content: RR and DP. Approving final version of manuscript: MH, HC, RL, RR, JY, TM, IS, and DP. DP takes responsibility for the integrity of the data analysis.

Role of the funding source

Not applicable.

Conflict of interest

DJP is a co-founder of Temple Therapeutics LLC. The other authors state that they have no conflicts of interest.

Acknowledgments

We thank Laura Quinlivan for assistance with animal experiments and Dina Gaupp, Center for Gene Therapy, Tulane University Health Sciences Center, for assistance with histological sectioning. This study was supported in part by National Institutes of Health/National Center for Research Resources Grant P40 RR 17447.

Supplementary material

Supplementary data related to this article can be found online at <http://dx.doi.org/10.1016/j.joca.2012.06.002>

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