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The Importance of Connexin Hemichannels During Chondroprogenitor Cell Differentiation in Hydrogel Versus Microtissue Culture Models

Karsten Schrobback, PhD, Travis Jacob Klein, PhD, and Tim B.F. Woodfield, PhD^{1,2}

Appropriate selection of scaffold architecture is a key challenge in cartilage tissue engineering. Gap junctionmediated intercellular contacts play important roles in precartilage condensation of mesenchymal cells. However, scaffold architecture could potentially restrict cell-cell communication and differentiation. This is particularly important when choosing the appropriate culture platform as well as scaffold-based strategy for clinical translation, that is, hydrogel or microtissues, for investigating differentiation of chondroprogenitor cells in cartilage tissue engineering. We, therefore, studied the influence of gap junction-mediated cell-cell communication on chondrogenesis of bone marrow-derived mesenchymal stromal cells (BM-MSCs) and articular chondrocytes. Expanded human chondrocytes and BM-MSCs were either (re-) differentiated in micromass cell pellets or encapsulated as isolated cells in alginate hydrogels. Samples were treated with and without the gap junction inhibitor 18-α glycyrrhetinic acid (18αGCA). DNA and glycosaminoglycan (GAG) content and gene expression levels (collagen I/II/X, aggrecan, and connexin 43) were quantified at various time points. Protein localization was determined using immunofluorescence, and adenosine-5'-triphosphate (ATP) was measured in conditioned media. While GAG/DNA was higher in alginate compared with pellets for chondrocytes, there were no differences in chondrogenic gene expression between culture models. Gap junction blocking reduced collagen II and extracellular ATP in all chondrocyte cultures and in BM-MSC hydrogels. However, differentiation capacity was not abolished completely by 18aGCA. Connexin 43 levels were high throughout chondrocyte cultures and peaked only later during BM-MSC differentiation, consistent with the delayed response of BM-MSCs to 18αGCA. Alginate hydrogels and microtissues are equally suited culture platforms for the chondrogenic (re-)differentiation of expanded human articular chondrocytes and BM-MSCs. Therefore, reducing direct cell-cell contacts does not affect in vitro chondrogenesis. However, blocking gap junctions compromises cell differentiation, pointing to a prominent role for hemichannel function in this process. Therefore, scaffold design strategies that promote an increasing distance between single chondroprogenitor cells do not restrict their differentiation potential in tissue-engineered constructs.

Introduction

AMAGE TO ARTICULAR CARTILAGE due to injuries or diseases frequently leads to a loss of the tissue's structure and function. This causes pain and impaired mobility for affected people and places a significant financial burden on healthcare systems and economies worldwide. Despite major advances in cartilage tissue engineering aimed at growing the patient's own cells on scaffolding materials for subsequent retransplantation, the reliable, functional repair or regeneration of articular cartilage remains a challenge. Key challenges in cartilage tissue engineering include the appropriate selection of cell source and scaffold architecture.

Multipotent mesenchymal stromal cells from human bone marrow (BM-MSC) and chondrocytes isolated from articular cartilage biopsies are currently the main sources for chondroprogenitor cells. However, we are still lacking appropriate culture protocols to initiate stable chondrogenic phenotypes of these cell types *in vitro*. For example, articular chondrocytes, in particular, when harvested from clinically relevant elderly patients, rapidly dedifferentiate during *ex vivo* propagation and lose their chondrogenic phenotype.² Unfortunately, this process seems only partially reversible under *in vitro* conditions as seen by sustained high levels of collagen type I, a marker of inferior fibrocartilage formation.^{2,3} BM-MSCs, on the other hand, tend to differentiate in a process

¹Institute of Health and Biomedical Innovation, Queensland University of Technology, Kelvin Grove, Australia.

²Department of Orthopedic Surgery, Center for Bioengineering & Nanomedicine, University of Otago, Christchurch, New Zealand.

more reminiscent of endochondral ossification, including the onset of tissue hypertrophy and calcification.⁴ To overcome some of these issues and to support the ex vivo differentiation of chondroprogenitor cells and the formation of functional cartilage tissue, a wide range of approaches for cell culture have been developed that could broadly be classified into scaffold-/matrix-based and biomaterial-free techniques.^{5,6} Among the former, hydrogels are very popular matrices for cartilage tissue engineering, as they provide a three-dimensional (3D) culture environment favorable for chondrogenic differentiation. 7-9 The simplest and most widely used scaffoldfree method is micromass pellet or microtissue culture, in which cell pellets are formed by centrifugation, allowing extensive cell-cell communication at the start of the culture. 10,11 This may be important, in particular, in the early stages of cell differentiation, considering the importance of cell-cell contacts to initiate chondrogenesis during precartilage condensation of mesenchymal cells in skeletogenesis. 12 However, the architecture of hydrogels, where the majority of cells are spatially separated by the hydrogel matrix, seems to inherently restrict direct cell-cell contacts and could potentially affect chondrogenic differentiation.

Intercellular communication during early chondrogenesis occurs via cell adhesion molecules, such as N-cadherin and neural cell adhesion molecule, or gap junctions. 13,14 While the first two molecules disappear in the course of chondrogenesis, ¹³ gap junctions can still be found in mature cartilage. ¹⁵ Gap junctions are intercellular channels formed by connexins that allow for rapid exchange of ions, metabolites, and other small-molecule messengers. ¹⁶ They are expressed in virtually all tissues and are involved in various cellular functions, including cell growth, differentiation, and death. One half of a gap junction channel that is not connected to an adjacent cell can still be functional and is termed a hemichannel. The major member of the connexin protein family in human cartilage is connexin 43 (Cx43, expressed by the GJA1 gene). ¹⁷ Apart from the role of gap junctions in mesenchymal cell differentiation, connexin hemichannels also seem to be involved in chondrocyte mechanotransduction.¹⁸ However, the role of gap junctional communication during the different stages of chondroprogenitor differentiation and its overall contribution to cartilage tissue formation is still less well defined. This is important in considering the appropriate culture platform, that is, hydrogel based or microtissues, for the different cell types.

This study was, therefore, aimed at comparing the differentiation potential of human articular chondrocytes and BM-MSCs in hydrogel and micromass culture systems with a particular focus on the influence of gap junction-mediated cell–cell interactions on *in vitro* chondrogenesis. We employed sodium alginate hydrogels and micromass pellets as 3D differentiation models that either restricted or supported direct intercellular contacts. We also used a pharmaceutical inhibitor of gap junction activity to test the importance of this form of cell–cell communication during various stages of chondroprogenitor differentiation and the formation of cartilaginous tissue *in vitro*.

Materials and Methods

Cell isolation and expansion

Articular cartilage was collected with institutional ethics approval from consenting patients (four male donors, age 26–

50 years) undergoing either limb amputations or reconstructions of the anterior cruciate ligament. Cartilage specimens were frozen in Optimal Cutting Temperature compound (OCT; Sakura, Finetek, Tokyo, Japan) for histological analysis. Chondrocytes were isolated from full-thickness cartilage of femoral condyles or grooves as previously described. 19 Chondrocytes were propagated on tissue culture plastic (3000 cells/ cm²) in chondrocyte basal medium (low-D-glucose Dulbecco's Modified Eagle Medium with 4 mM L-alanyl-L-glutamine, 1 mM sodium pyruvate, 10 mMN-(2-hydroxyethyl)piperazine-N'-(2-ethanesulfonic acid) (HEPES), 0.1 mM nonessential amino acids, 50 U/mL penicillin/50 µg/mL streptomycin (Pen/ Strep; all from Life Technologies, Carlsbad, CA), 0.1 mM L-ascorbic acid 2-phosphate, and 0.4 mM L-proline (both Sigma Aldrich, St. Louis, MO)) supplemented with 10% fetal bovine serum (FBS; Hyclone, Logan, UT). Cells were used in subsequent experiments after passage 2 (approximately five to six population doublings).

Mesenchymal stromal cells were isolated from bone marrow aspirates obtained under appropriate ethical approval from the iliac crest of consenting patients (one female+two male donors, age 22–30 years) undergoing spinal fusion surgery. Total bone marrow cells were plated in α-MEM supplemented with 10% FBS, Pen/Strep, and 1 ng/mL fibroblast growth factor 2 (FGF-2; Millipore, Billerica, MA). BM-MSCs were initially seeded at 100 cells/cm², subsequently plated at 1000 cells/cm², and used between passages 3 and 4.

All cells were maintained at 37°C in a humidified 5% CO₂/95% air CO₂ incubator with medium refreshed twice per week. Cells were passaged when subconfluent with 0.25% trypsin with 1 mM ethylenediamine tetraacetic acid (trypsin/EDTA; Life Technologies) at 37°C for 5 min and counted by trypan blue (Life Technologies) exclusion in a hemocytometer.

Chondrogenic differentiation

Chondrogenic potential was evaluated using either micromass pellet or alginate hydrogel cultures. Pellets of both chondrocytes and BM-MSCs were formed by centrifugation.¹⁹ Briefly, cells resuspended in a chondrogenic media were transferred into V-bottom-shaped, 96-well microplates (AXY-GEN Scientific, Union City, CA) at 2×10^5 cells in 0.2 mL media per well and centrifuged at 600 g for 5 min. After 3 days, cells had condensed into micropellets that could be lifted from the bottom of the wells and transferred into 48-well plates containing 0.5 mL chondrogenic media. Cells were also encapsulated in 2% w/v sodium alginate (Pronova UP LVG, Novamatrix/FMC BioPolymers, Sandvika, Norway) at 10⁷ cells/mL.²⁰ Briefly, the cell alginate suspension was transferred into wells (height: 1.5 mm, diameter: 4 mm, volume: $\sim 20 \,\mu\text{L}$) of a stainless steel custom-made mold, placed between sterile filter papers, and soaked in 102 mM CaCl₂ for 10 min. Hydrogel constructs were then washed with serum-free basal medium twice for 5 min and transferred into 48-well plates containing 0.5 mL chondrogenic media. Cells were cultivated in serumfree high-glucose chondrocyte basal medium supplemented with 1.25 mg/mL bovine serum albumin, 10⁻⁷ M dexamethasone, (all from Sigma), 1% v/v ITS, and 10 ng/mL transforming growth factor type beta 1 (TGF-β1; both Life Technologies) and kept under reduced (5%) oxygen tension in a ProOx C-Chamber (Biospherix, Redfield, NY) inside a cell culture incubator. To reversibly block gap junction activity, $80\,\mu\text{M}$ of $18\text{-}\alpha$ glycyrrhetinic acid ($18\alpha\text{GCA}$; Sigma) was added to the chondrogenic media of selected constructs at various time intervals. This concentration for $18\alpha\text{GCA}$ is well less than what has been reported to be toxic to human fibroblasts in long-term experiments but potent enough to reduce intercellular junctional communication in albumin-containing cultures. Cultures were maintained till 9 days with media refreshed every 3 days and conditioned media was stored at -80°C for further analysis. Photographs of pellets were taken with a phase-contrast microscope (NIKON, Tokyo, Japan) at day 4 and 9 of culture, and pellet sizes were measured using ImageJ (National Institutes of Health, Bethesda, MD).

Quantification of glycosaminoglycans and DNA

Cells in constructs were harvested for glycosaminoglycan (GAG) and DNA quantification by incubation with 0.5 mg/mL proteinase K (Life Technologies) in a phosphate buffer (20 mM Na₂HPO₄, 30 mM NaH₂PO₄.H₂O, and 5 mM EDTA, pH 7.1) at 60°C overnight.

The GAG content of conditioned media as well as digested alginate samples and cell pellets was quantified using the 1,9-dimethylmethylene blue dye (DMMB) assay (pH 1.5).²² Absorbances at 525 and 595 nm were measured in a Benchmark Plus microplate spectrophotometer (Bio-Rad Laboratories, Hercules, CA), and the concentrations were calculated using the ratio of absorbances, compared with a quadratic standard curve prepared from chondroitin sulfate C (Sigma). Data from the first media change were excluded to allow for diffusion of unbound alginate from gels groups.

The Quant-iTTM PicoGreen[®] dsDNA assay (Life Technologies) was used to quantify DNA content in digested samples, with fluorescence (485 nm excitation, 520 nm emission) measured using a POLARstar OPTIMA fluorescence microplate reader (BMG Labtech, Offenburg, Germany).

Ribonucleic acid extraction and real-time reverse transcriptase-polymerase chain reaction

To measure absolute expression levels of selected genes of interest, real-time reverse transcriptase-polymerase chain reaction (qRT-PCR) was used. Ribonucleic acid (RNA) was isolated using the PureLink™ RNA Micro Kit (Life Technologies) according to the manufacturer's instructions, which included mechanic disruption of cell pellets with pestles and dissolution of alginate hydrogels in 10 mM HEPES, 50 mM sodium EDTA, and 150 mM NaCl for 10 min, followed by sample homogenization with a 20-gauge needle and the removal of genomic DNA by column treatment with DNase I (Life Technologies).

First-strand complementary DNA (cDNA) synthesis was performed using the SuperScriptTM III first-strand synthesis supermix for qRT-PCR (Life Technologies) with no more than 300 ng of total RNA in 20 μ L reactions following the manufacturer's protocol.

Primers were either used as previously published for *COL1A1*, *COL2A1*, *COL10A1*, *ACAN*, *RPL13A*, and $B2M^{23}$ or designed using PrimerBank²⁴ and Primer BLAST (NCBI, Bethesda, MD) for *GJA1* ($5' \rightarrow 3'$ F: GACCGGATAGTCAA GTTCGTAGC, R: GCAGGAGCTGTCCACGTAG), and TBP ($5' \rightarrow 3'$ F: GAGCCAAGAGTGAAGAACAGTC, R: CA TCACAGCTCCCCACCATATT). All PCR reactions were

done in duplicate in 10 μ L volumes in a 7500 PCR system (Applied Biosystems, Foster City, CA). Reactions contained 1× SYBR® Green PCR Master Mix (Applied Biosystems), 200 nM of each forward and reverse primers, and 0.3 μ L of undiluted cDNA and started with 10 min denaturation at 95°C followed by 40 cycles of 95°C for 15 s and 60°C for 30 s. The cycle threshold (Ct) value of each gene was normalized to the geometric mean of the housekeeping genes *B2M*, *RPL13A*, and *TBP* using the comparative Ct method (2 $^{-\Delta Ct}$).

Immunofluorescence staining analysis

To determine the expression of proteins of interest, samples from alginate constructs, pellets, and native cartilage were frozen in OCT (Sakura) and sectioned. Fresh-frozen sections were fixed in 100% cold acetone for 15 min and then hydrated in 50 mM BaCl₂/100 mM Tris HCl (pH 7.3) buffer for 1 h. Antigen retrieval was only included in the collagen I and II staining procedures by applying 0.1% w/v hyaluronidase (Sigma) in phosphate-buffered saline (PBS) for 20 min at 37°C. Sections were then probed overnight at 4°C in PBS with 2% donkey or goat serum (2% serum/PBS; Jackson ImmunoResearch, West Grove, PA) and one of the following antibodies: anti-collagen I (mouse IgG_{2a}, 1:300, I-8H5; MP Biomedicals, Solon, OH), anti-collagen II (mouse IgG₁, 1:200, II-II6B3; Developmental Studies Hybridoma Bank [DSHB]), anti-aggrecan (mouse IgG₁, 1:300, 969D4D11; Life Technologies), anti-connexin 43 (rabbit IgG, 1:200, C6219; Sigma), or isotype controls (rabbit IgG, ABCAM/Sapphire, Redfern, NSW; mouse IgG, Jackson; both 1:1000).

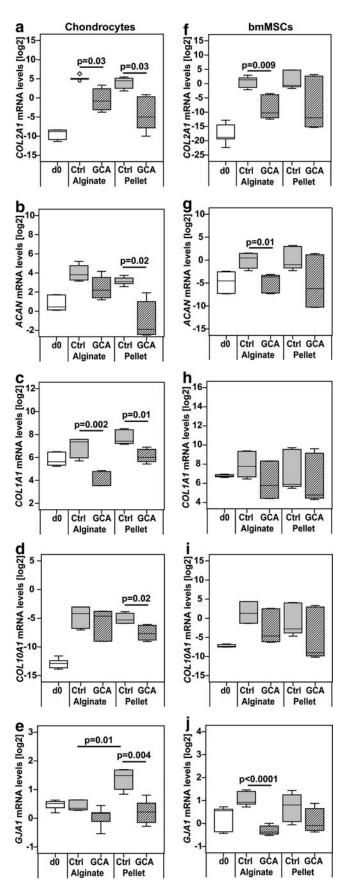
Subsequently, samples were incubated in 2% serum/PBS with a mix of either AlexaFluor®488-labeled goat antimouse (IgG₁-specific) and AlexaFluor®594-labeled goat anti-mouse (IgG_{2a}-specific) or AlexaFluor488-labeled donkey anti-mouse IgG and AlexaFluor594-labeled donkey anti-rabbit IgG (all minimal x-reaction with other species, 1:150, Jackson) and $5\,\mu\text{g/mL}$ 4',6-diamidino-2-phenylindole (DAPI; Sigma) for 1 h at room temperature. Images were captured with fixed exposure times on an Axio Imager.A1 microscope with epi-fluorescence attachment (Carl Zeiss, Jena, Germany).

Adenosine-5'-triphosphate assay

To quantify adenosine-5'-triphosphate (ATP) released into media, samples were taken at day 9 of chondrogenic culture, heated for 5 min at 95°C to inactivate ATPases, and stored at -80°C. The ATP Determination kit (Life Technologies) was used according to the manufacturer's instructions, and luminescence was measured in 10 cycles of 2 s detections using the POLARstar micro-plate reader, immediately after addition of the luciferase reaction mix to the media samples. ATP concentrations were calculated from averaged luminescence measurements compared with a quadratic standard curve of ATP provided with the assay kit and further normalized to the average DNA content of samples from each treatment group and donor at day 9 of culture.

Statistical analysis

Statistical analyses were performed using SPSS Statistics v18 (IBM, Somers, NY). To test for differences between the cell culture types as well as control and $18\alpha GCA$ -treated



samples in alginate or pellet models, data from gene expression, biochemical, and ATP assays were transformed using the binary logarithm to improve normality distribution of the data sets and tested by analysis of variance (ANOVA) using a general linear model including Dunnett's T3 post hoc test for multiple comparisons between groups. To test the effect of intermittent blocking of gap junctional communication on chondrogenic differentiation, gene expression data from day 3, 6, and 9 of pellet cultures were subjected to ANOVA using a general linear model, including Tukey's HSD post hoc test. Statistically significant differences were considered to be present at p < 0.05.

Results

Effect of cell–cell interactions and gap junction blocking on gene expression during chondrogenic differentiation

To study the importance of cell-cell interactions during differentiation of chondroprogenitor cells in vitro, expanded chondrocytes and BM-MSCs were cultured in hydrogel or micromass pellets, which either restricted or allowed extensive direct cell-cell interactions, respectively. In addition, gap junctional communication was blocked using the pharmaceutical compound 18\alpha GCA. After 9 days of differentiation culture, the expression of the chondrogenic marker collagen type II (COL2A1) was significantly lower in 18αGCA-treated chondrocytes in both culture types compared with untreated controls (p = 0.03; Fig. 1a). However, blocking gap junctions reduced the mRNA levels of aggrecan (ACAN), another chondrogenic marker, in chondrocyte pellets (p = 0.02) but not in alginate cultures (p=0.19; Fig. 1b). Similar to *COL2A1*, transcription of the dedifferentiation marker collagen type I, COL1A1, was hampered by 18α GCA in chondrocyte alginate (p=0.002) and pellet cultures (p = 0.01) in comparison to controls (Fig. 1c). Chondrocytes exhibited decreased transcription of collagen type X (COL10A1), a marker for tissue hypertrophy and calcification, only when cultured in pellets in the presence of a gap junction blocker (p = 0.02; Fig. 1d). The gene expression of the major gap junction protein (GJA1) connexin 43 was elevated in untreated chondrocyte pellets compared with hydrogel cultures (p = 0.01), but decreased only in pellets on 18α GCA treatment (p = 0.004; Fig. 1e). In

FIG. 1. Gene expression levels of (a, f) collagen II (COL2A1), (b, g) aggrecan (ACAN), (c, h) collagen I (COL1A1), (d, i) collagen X (COL10A1), and (e, j) connexin 43 (GJA1) in (a-e) human articular chondrocytes or (f-j) BM-MSCs after treatment with the gap junction blocker 18αGCA. Cells were differentiated in alginate hydrogels or pellet cultures in the absence (Ctrl) or presence of 18α GCA (GCA). The gene expression levels were measured at day 0 (d0) and day 9 of culture with real-time reverse transcriptasepolymerase chain reaction, then normalized to housekeeping genes using the comparative deltaCt method, and transformed into the binary logarithm (log2). For clarity, statistical significance was only shown for comparisons between 18αGCAtreated groups and controls as well as alginate Ctrl versus pellet Ctrl (n=6 from three donors of each cell type). 18α GCA, $18-\alpha$ glycyrrhetinic acid; BM-MSC, bone marrow-derived mesenchymal stromal cell.

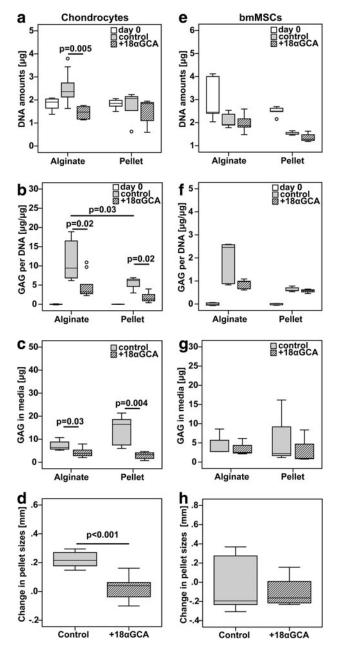


FIG. 2. Effect of gap junction blocking on DNA and GAG synthesis of (a-d) human articular chondrocytes and (e-h) BM-MSCs in alginate hydrogel and micromass pellet cultures. Cells of both types were cultured under identical chondrogenic conditions in the absence (control) or the presence of the gap junction blocker 18-α glycyrrhetinic acid ($+18\alpha$ GCA). The amounts of (a, e) DNA and (b, f) GAG per DNA were measured in constructs collected before (day 0) and at the end of the culture period. The cumulative amount of (c, g) secreted GAG was measured in conditioned media from day 6 and 9 of culture. The (d, h) change in pellet sizes was measured based on photographs taken with a phase-contrast microscope at day 4 and 9 of culture. For clarity, statistical significance was only shown for comparisons between 18\alpha GCA-treated groups and controls or alginate Ctrl versus pellet Ctrl (chondrocytes from three donors: n=7 pellets, n=9 alginate constructs, n=15pellets for size measurements; MSC from two donors: n=5pellets or alginate constructs, n=9 pellets from three donors for size measurements). GAG, glycosaminoglycan.

MSCs, gap junction blocking reduced mRNA levels of COL2A1 (p=0.009) and ACAN (p=0.01) in alginate cultures compared with their respective controls, and although similar trends were observed in pellets, they failed to reach significance (COL2A1: p=0.3; ACAN: p=0.39; Fig. 1f, g). Transcription levels of both COL1A1 and COL10A1 were not significantly different between 18αGCA-treated MSC alginate and pellet samples and control groups (Fig. 1h, i). In line with the chondrogenic markers, GJA1 expression was only lower in treated MSC alginate cultures (p < 0.0001) versus controls but unchanged in pellets in response to gap junction blocking (p = 0.52; Fig. 1j). The mRNA levels of the matrix molecules (COL2A1, ACAN, COL1A1, and COL10A1) were not significantly different between untreated alginate and pellet cultures in either chondrocytes or MSCs (Fig. 1).

Effect of cell–cell interactions and gap junction blocking on DNA and matrix production during chondrogenic differentiation

To study the importance of cell-cell interactions and gap junctional communication on matrix production of chondroprogenitor cells in vitro, we compared the DNA and GAG production of chondrocytes and BM-MSCs cultured for 9 days in hydrogel or micromass pellets in the presence or absence of 18αGCA. Blocking of gap junctions decreased DNA content (p=0.005), GAG per DNA (p=0.02), and the GAG in media (p=0.03) as well as the size (p<0.001) of chondrocyte pellets compared with untreated groups (Fig. 2a-d). Similarly, GAG per DNA in constructs as well as the GAG secreted into the media was also lower in chondrocyte alginate cultures treated with 18\alpha GCA versus controls (Fig. 2b, c). However, DNA content was not different between these two groups. The GAG per DNA accumulated by chondrocytes over 9 days was higher in untreated alginate cultures in comparison to pellets (p=0.03; Fig. 2b). In contrast to chondrocytes, DNA amounts, GAG per DNA, GAG in media, and pellet sizes were not different between BM-MSCs cultured in the presence and absence of the gap junction blocker in either alginate or pellet cultures (Fig. 2e-h). DNA and GAG synthesis of BM-MSCs was also comparable between alginate and pellet control cultures (Fig. 2e-g).

Effect of cell–cell interactions and gap junction blocking on protein expression

The deposition of extracellular matrix after 9 days in differentiation culture under varying levels of gap junctional communication was assessed by probing with antibodies specific for collagen type I and II. Exposure to the gap junction blocker reduced the staining intensity for both the chondrogenic collagen type II and the dedifferentiation marker collagen type I in all culture models with chondrocytes (Fig. 3a-d). Compared with control pellets, sections from 18aGCA-treated chondrocyte pellets revealed overall higher cell density and less matrix at day 9 of culture and unusual islands of intense collagen I staining (Fig. 3c, d). Similar to chondrocytes, 18αGCA-treated MSCs showed less immunoreactivity against collagen I compared with untreated controls (Fig. 3e-h). Collagen type II staining was weak in both culture types with BM-MSCs and only limited to single cells in the untreated controls (Fig. 3e-h). Inherent

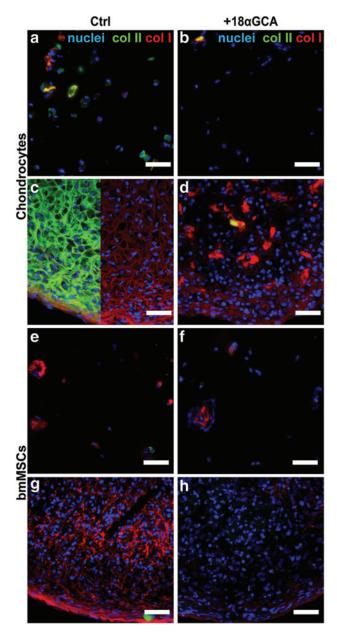


FIG. 3. Expression of collagen type II and type I from (a–d) human articular chondrocytes and (e–h) BM-MSCs in (a, b, e, f) alginate hydrogel and (c, d, g, h) micromass pellet cultures. Cells of both types were cultured under identical chondrogenic conditions in the (a, c, e, g) absence (Ctrl) or the (b, d, f, h) presence of the gap junction blocker 18-α glycyrrhetinic acid (+18αGCA). Representative samples were processed for fresh-frozen tissue immunohistology at day 9, simultaneously probed with antibodies against collagen type II (col II, green) or collagen type I (col I, red), and counterstained with DAPI (blue). In panel c, col II signal was omitted in the right half of the image for better visibility of col I staining. Scale bars = 50 μm. DAPI, 4',6-diamidino-2-phenylindole. Color images available online at www.liebertpub.com/tea

differences in cell seeding densities between alginate and pellets cultures preclude a direct comparison of their matrix qualities based on immunofluorescence staining.

Sections from the various groups at day 3 and 9 of culture were also probed with antibodies for the cartilaginous ma-

trix protein, aggrecan, and the gap junction protein connexin 43 (Fig. 4). All groups revealed immunoreactive regions for aggrecan that were confined to the extracellular matrix. At day 9, untreated alginate and pellet controls with chondrocytes and BM-MSCs showed slightly more intense immunoreactivity for aggrecan and connexin 43 compared with 18αGCA-treated groups (Fig. 4b, c, e, f, h, i, k, l). There were no clear differences in connexin 43 staining between day 3 and 9 of chondrocytes or BM-MSCs in alginate cultures (Fig. 4a, b, g, h). However, chondrocyte pellets showed more intense, punctate staining throughout all regions at the earlier compared with the later time point, which coincided with higher cell density and lower immunoreactivity for aggrecan at day 3 compared to day 9 (Fig. 4d, e). In contrast, with BM-MSCs, areas of more intense, clustered staining for connexin 43 appeared to be localized to the outer rim of pellet sections at both time points (Fig. 4j, k). Control sections from the superficial and middle/deep zone of human native cartilage revealed extracellular matrix positive for aggrecan as well as localized cellular staining for connexin 43, which was seemingly less intense compared with pellet cultures (Fig. 4m, n). Stainings with IgG isotype control antibodies from mouse and rabbit proved negligible unspecific background reactivity (Fig. 4o).

Effect of cell culture models and gap junction blocking on extracellular ATP levels

Connexin hemichannels have been implicated in the release of ATP into the extracellular environment. Therefore, we measured ATP levels in the conditioned media from chondrocytes and BM-MSCs cultured in alginate or pellets in the presence or absence of 18α GCA between days 6 and 9 and normalized these values to the average DNA content of samples from each treatment group and donor at day 9 of culture. Gap junction blocking decreased the amount of ATP in media normalized to DNA of chondrocyte alginate (p=0.007) and pellet cultures (p=0.002; Fig. 5a). The same effect was observed for BM-MSCs in alginate cultures (p=0.014) but not in pellets (p=0.89; Fig. 5b). ATP amounts in media per DNA were not different between alginate and pellet control groups of chondrocytes (p=0.18) nor BM-MSCs (p=0.7; Fig. 5a, b).

Discussion

Intercellular communication via gap junctions is an important driver of mesenchymal condensation in early chondrogenic differentiation. ¹² However, it is far from clear whether direct cell-to-cell contacts and the exchange of messenger molecules through gap junctions also have an essential role in the differentiation of chondroprogenitor cells in *ex vivo* engineered cartilage constructs. Furthermore, scaffold architecture and composition play a critical role not only in modulating chondrogenic differentiation and diffusion of nutrients, ^{25,26} but also in controlling cell distribution in 3D. ²⁷ However, the influence of 3D cell distribution and concomitant cell–cell communication on the differentiation capacity of chondroprogenitor cells offered by either high-density microtissue versus more discrete hydrogel culture platforms for cartilage tissue engineering has also not been well established.

Our data show that differentiating chondrocytes or BM-MSCs in micromass cell pellets brought no advantage compared with alginate hydrogel cultures in terms of the cell's

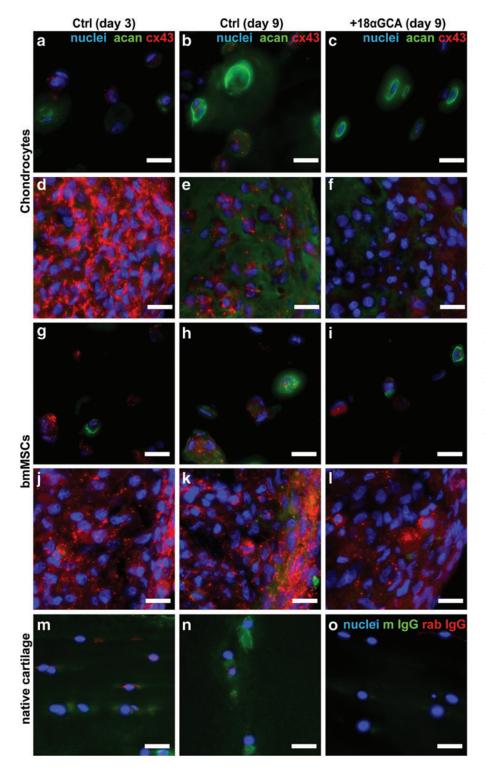


FIG. 4. Expression of aggrecan and connexin 43 from (a-f) human articular chondrocytes and (g-l) BM-MSCs in $(\mathbf{a}, \mathbf{b}, \mathbf{c}, \mathbf{g}, \mathbf{h}, \mathbf{i})$ alginate hydrogel and (d, e, f, j, k, l) micromass pellet cultures. Cells of both types were cultured for (a, d, **g**, **j**) 3 days or (**b**, **c**, **e**, **f**, **h**, **i**, **k**, **l**) 9 days under identical chondrogenic conditions in the (a, b, d, e, g, h, j, **k**) absence (Ctrl) or the (**c**, **f**, **i**, **l**) presence of the gap junction blocker 18-a glycyrrhetinic acid (+18αGCA). Representative samples were processed for freshfrozen tissue immunohistology, simultaneously probed with antibodies against aggrecan (acan, green) or connexin 43 (cx43, red), and counterstained with DAPI (blue). Sections from the (m, o) superficial or (n) the middle zone of native human articular cartilage either (m, n) stained for aggrecan/ connexin 43 or (o) probed with unspecific antibodies against mouse IgG (m IgG) and rabbit IgG (rab IgG) are shown for comparison. Scale bars = $20 \,\mu m$. Color images available online at www.liebertpub .com/tea

chondrogenic marker expression and GAG synthesis. Our findings are in agreement with other studies, in which neither human chondrocytes²⁸ nor human BM-MSCs²⁹ showed clear differences in chondrogenic marker expression when differentiated in either pellet or alginate cultures. This suggests that promoting direct intercellular contacts may not be critical for the differentiation of chondroprogenitor cells *in vitro*.

Despite this, we still found that blocking gap junctional intercellular communication using the pharmaceutical inhibitor $18\alpha GCA$ negatively affected chondrogenic differentiation, as evident in a reduction in GAG synthesis and in the expression of the important chondrocyte marker, collagen type II (Figs. 1 and 2). Interestingly, $18\alpha GCA$ treatment did not altogether abolish the expression of chondrogenic marker molecules and seemed to affect chondrocytes and BM-MSCs to a different extent depending on the cell culture model. Aggrecan mRNA levels of chondrocytes, for example, were only significantly reduced in pellets but not in

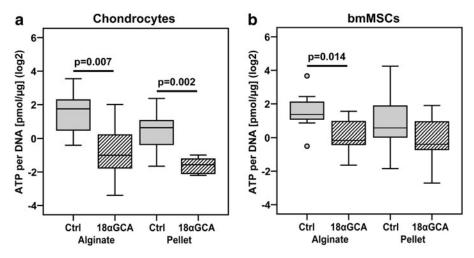


FIG. 5. Effect of gap junction blocking on ATP secretion of (a) human articular chondrocytes and (b) BM-MSCs in alginate hydrogel and micromass pellet cultures. Cells of both types were cultured under identical chondrogenic conditions in the absence (Ctrl) or the presence of the gap junction blocker $18-\alpha$ glycyrrhetinic acid ($+18\alpha$ GCA). The amount of secreted ATP was measured in conditioned media from day 9 of culture, normalized to the average DNA content of samples from each treatment group and donor at day 9 of culture, and transformed into the binary logarithm (log2) (n=8-12 from two donors of each cell type). ATP, adenosine-5'-triphosphate.

alginate, which may be explained by the higher connexin 43 expression in the former compared with the latter (Fig. 1). In contrast, while there was a strong impact on *COL2A1* and aggrecan expressions in BM-MSC alginate cultures, reduction in these markers was less severe in BM-MSC pellets. It should be noted that there were large donor-to-donor variations for BM-MSCs in terms of their chondrogenic potential, which may explain the high standard deviations in the mRNA and biochemical data sets and could be—at least in part—blamed for the difficulties in detecting significant effects for this cell type.

Another interesting difference between the two cell types was that 18αGCA treatment had the most profound effect in the first few days of chondrocyte differentiation in pellets; however, chondrogenesis of BM-MSCs appeared to be disturbed only past day 3 (Supplementary Fig. S1; Supplementary Data are available online at www.liebertpub.com/ tea). This has also been observed in mesenchymal cells from chicken limb buds, which only showed a discernable response to 18αGCA treatment 48 h after initiation of chondrogenic differentiation.³⁰ It could, therefore, be speculated that BM-MSCs, the more immature chondroprogenitor cell type as evidenced by relatively low COL2A1 and ACAN expression at the start of the differentiation, respond only to impediments of their gap junction function after the chondrogenic phenotype has been established. It is interesting to note in this regard that connexin 43 protein expression was, indeed, highest at day 3 of chondrocyte pellet culture whereas BM-MSC pellets appeared to exhibit higher connexin 43 levels at day 9 than day 3 (Fig. 4). In addition, our results and those of other groups indicate that connexin 43 mRNA and protein levels decrease in cells in response to 18αGCA at concentrations higher than 20 μM.³¹ This raises the question about the mechanism of action of the gap junction blocker 18\(\alpha\)GCA, which does not seem to be fully elucidated to date. We only observed minor effects on cell numbers in treated cultures. DNA levels in 18αGCA samples were not different from controls except for chondrocyte alginate cultures, in which case DNA quantities of the $18\alpha GCA$ group were still at similar levels as measured at day 0 of culture (Fig. 2). Some evidence points to the incorporation of $18\alpha GCA$ into the cell membrane, where it disrupts the formation of gap junction plaques and possibly induces a conformational change of connexin closing the gap junctions. Furthermore, it has been proposed that glycyrrhetinic acid interferes with the interaction between connexin 43 and phosphatases or kinases, altering connexin phosphorylation and leading to reversible closure of gap junction and reduced gap junction intercellular communication. 31,33 Therefore, the main mode of actions of $18\alpha GCA$ is most likely a combination of decrease in connexin expressions and interference with opening of gap junctions.

A role for gap junctional intercellular communication in cell differentiation has been established for a number of cell types, including epithelial and mesenchymal cells.30,34 Our study shows for the first time that chondrogenesis is even hampered by 18αGCA in alginate cultures with already limited cell-cell contacts. This suggests that the effect of gap junction-mediated crosstalk on chondrocyte differentiation is mainly exerted through the formation and activity of hemichannels. Hemichannels are aqueous membrane pores that allow for the controlled release of small messenger molecules (e.g., ATP, prostaglandin E₂, ions) for autocrine/ paracrine cellular communication in response to a range of stimuli.³⁵ In chondrocytes, connexin hemichannels seem to be involved in the release of intracellular ATP into the extracellular space after cyclic loading, thereby modulating purinergic and calcium signaling pathways. ^{18,36} In our study, we found, indeed, reduced ATP levels in the media of almost all groups treated with a gap junction blocker compared with their controls (Fig. 5). Only media samples from BM-MSC pellets showed no differences for this nucleotide, which may be explained by ATP release from dying or dead cells that were observed at relatively high levels in these

cultures (Fig. 2e). In summary, our data clearly point to a broader, important role for hemichannel activity and ATP release in chondrogenic differentiation even in the absence of mechanical stimuli. Indeed, a study on the mouse chondrogenic cell line ATDC5 revealed synchronized ATP oscillations during early differentiation that appeared gap junction dependent and positively affected prechondrogenic condensation.³⁷ However, the signals driving gap junctional ATP release in the absence of cyclic mechanical cues still remain elusive.

In this regard, it is tempting to speculate about the potential use of small-molecule messengers, such as ATP, as relatively inexpensive media supplements to improve quality of engineered cartilage constructs. However, it may not be as simple as continuously adding ATP to culture media. When human BM-MSCs and bovine chondrocytes were differentiated in agarose hydrogels, the presence of ATP did not enhance matrix contents and brought only minor improvements to mechanical properties of the constructs. Therefore, the role for extracellular ATP in chondrogenic differentiation is likely to be complex and requires further investigation.

Conclusions

Alginate hydrogel and microtissue culture systems are equally suited to study the *in vitro* differentiation of articular chondrocytes and bMSCs. Immediate cell-to-cell interactions offered by micromass culture do not play a significant role in the *in vitro* chondrogenesis of expanded human chondrocytes and human BM-MSCs compared with hydrogel culture. Therefore, restricting direct intercellular communication through scaffold architecture should not affect the chondrogenic potential of isolated cells in scaffoldbased cartilage repair strategies. However, for the first time to our knowledge, our study demonstrates the importance of connexin hemichannels in gap junction-mediated crosstalk between chondrocytes during differentiation that goes beyond their involvement in chondrocyte mechanotransduction, and appears to control the release of small molecules, such as ATP, for autocrine and paracrine signaling. Further studies need to address the specific role for connexin hemichannels in coordinating or regulating chondrogenic differentiation and identify the specific cues that drive hemichannel activity. This will give new insights into the complex processes of chondrogenesis and potentially reveal new important factors and strategies to improve current approaches in cartilage tissue engineering.

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Disclosure Statement

No competing financial interests exist.

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Address correspondence to:

Tim B.F. Woodfield, PhD

Department of Orthopaedic Surgery

Centre for Bioengineering & Nanomedicine

University of Otago

PO Box 4345

Christchurch 8140

New Zealand

E-mail: tim.woodfield@otago.ac.nz

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