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Developmentally Inspired Combined Mechanical and Biochemical Signaling Approach on Zonal Lineage Commitment of Mesenchymal Stem Cells in Articular Cartilage Regeneration

Tahereh Karimi, Danial Barati, Ozan Karaman, Sina Moeinzadeh, and Esmail Jabbari

Biomimetic Materials and Tissue Engineering Laboratory, Department of Chemical Engineering, University of South Carolina, Columbia, SC 29208, USA

Abstract

Articular cartilage is organized into multiple zones including superficial, middle and calcified zones with distinct cellular and extracellular components to impart lubrication, compressive strength, and rigidity for load transmission to bone, respectively. During native cartilage tissue development, changes in biochemical, mechanical, and cellular factors direct the formation of stratified structure of articular cartilage. The objective of this work was to investigate the effect of combined gradients in cell density, matrix stiffness, and zone-specific growth factors on the zonal organization of articular cartilage. Human mesenchymal stem cells (hMSCs) were encapsulated in acrylate-functionalized lactide-chain-extended polyethylene glycol (SPELA) gels simulating cell density and stiffness of the superficial, middle and calcified zones. The cell-encapsulated gels were cultivated in medium supplemented with growth factors specific to each zone and the expression of zone-specific markers was measured with incubation time. Encapsulation of 60×10^6 cells/mL hMSCs in a soft gel (80 kPa modulus) and cultivation with a combination of TGF- β 1 (3 ng/mL) and BMP-7 (100 ng/mL) led to the expression of markers for the superficial zone. Conversely, encapsulation of 15×10^6 cells/mL hMSCs in a stiff gel (320 MPa modulus) and cultivation with a combination of TGF- β 1 (30 ng/mL) and hydroxyapatite (3%) led to the expression of markers for the calcified zone. Further, encapsulation of 20×10^6 cells/mL hMSCs in a gel with 2.1 MPa modulus and cultivation with a combination of TGF- β 1 (30 ng/mL) and IGF-1 (100 ng/mL) led to up-regulation of the middle zone markers. Results demonstrate that a developmental approach with gradients in cell density, matrix stiffness, and zone-specific growth factors can potentially regenerate zonal structure of the articular cartilage.

Keywords

articular cartilage; zonal structure; human MSC; cell density gradient; stiffness gradient; growth factor gradient; chondrogenesis

Corresponding author: Esmail Jabbari, Ph.D., Associate Professor of Chemical and Biomedical Engineering, Swearingen Engineering Center, Rm 2C11 University of South Carolina, Columbia, SC 29208, Tel: (803) 777-8022, Fax: (803) 777-0973, jabbari@mailbox.sc.edu.

Introduction

Articular cartilage also called hyaline cartilage is a multi-functional tissue with a highly organized structure which provides a lubricating surface for the gliding joints and a load-bearing matrix attached to the underlying bone.^{1, 2} The postnatal articular cartilage due to its avascular nature and low metabolism lacks the ability for complete self-repair through native healing mechanisms.³ About 70% of people over the age of 65 suffer from degenerative joint disease or osteoarthritis (OA).⁴ The center for disease control and prevention estimates that nearly 27 million Americans suffer from joint pain and stiffness, loss of function and disability.⁵ Clinical approaches to articular cartilage repair include debridement, lavage and joint replacement in advanced forms of OA.³ Current treatment methods rarely restore the full function to the joint.³ The challenge in regeneration of articular cartilage is recreating the spatially variant structure of the native tissue with complex interactions between resident cells, extracellular matrix (ECM) and soluble growth factors. Several studies have reported construction of a homogenous matrix with bulk properties of the native cartilage tissue for implantation in a cartilage defect using a top-down approach in which chondrogenic cells were seeded into preformed porous scaffolds or embedded in injectable matrices loaded with signaling factors.^{4, 6-8} The generated tissues lacked the desired functions because top-down approaches do not mimic the intricate zonal organization of the native articular cartilage.

Articular cartilage is structurally and functionally composed of multiple distinct zones including superficial, middle, deep, and calcified zones^{1, 2, 9} with each zone exhibiting a defined ECM composition and organization.^{1, 3} Recent studies have attempted to mimic this spatial organization using the bottom-up approach with zone-specific chondrocytes isolated from animal tissues.^{10, 11} Ng and coworkers used zone-specific chondrocytes seeded on layered agarose hydrogels to recreate depth-dependent anisotropy of the articular cartilage.¹⁰ Sharma and collaborators used bilayer hydrogels to reorganize zone-specific chondrocytes into a stratified structure.¹² Cui and collaborators printed human chondrocytes into osteochondral plugs layer-by-layer to repair cartilage defects.¹³ Although chondrocytes are the most relevant cell source for regeneration of articular cartilage defects, there is limited quantity of autologous chondrocytes for transplantation. In addition, expansion of chondrocytes in monolayer culture causes dedifferentiation with decreased proteoglycan and collagen type II (Col II) synthesis.^{4, 14-16} An attractive alternative is the use of mesenchymal stem cells (MSCs) harvested from the patient's bone marrow or adipose tissue.⁴ MSCs can be expanded and differentiated into zone-specific chondrocytes by seeding them in scaffolds that mimic the mechanical and biochemical properties of the articular cartilage and cultivated in a medium supplemented with chondrogenic growth factors.¹⁷ Several approaches including multilayer hydrogels^{18, 19, 13} and multilayer electrospun fibers^{20, 21} have been applied to create scaffolds mimicking the composition and mechanical properties of the native articular cartilage. Steele et al. generated bilayer poly(ϵ -caprolactone) (PCL) scaffolds comprised of an aligned fiber layer mimicking the morphology and mechanics of the superficial zone laminated to a porous scaffold simulating the middle zone of articular cartilage.²¹

The articular cartilage zones are maintained by distinct cellular phenotypes, ECM compositions, and growth factors. The superficial zone is characterized by thin collagen fibrils parallel to the articular surface and the chondrocytes in this zone have high expression of Sox-9, an early chondrogenic differentiation marker, and superficial zone protein (SZP) for lubricating the joint during movement.²² The middle zone which takes up as much as 60% of the cartilage volume is composed of collagen fibrils thicker than those in the superficial zone with oblique alignment to the cartilage surface. The middle zone provides compressive strength and dimensional stability and it is characterized by pre-hypertrophic chondrocytes with high expression of glycosaminoglycans (GAG) and aggrecans (AGC).^{3, 18} The deep zone consists of large diameter fibrils oriented perpendicular to the articulating surface (similar to the calcified zone) with functions similar to the middle zone. Chondrocytes in the deep zone express highest amount of proteoglycans.²³ The calcified zone is an avascular tissue at the interface between uncalcified hyaline cartilage and subchondral bone and collagen fibers in this zone are oriented perpendicular to the articular surface. The calcified cartilage has a unique mineralized matrix composition, with chondrocytes expressing markers of hypertrophy including collagen type X (Col X) and alkaline phosphatase (ALP).^{3, 5, 18} To limit the scope of the study and the number of experimental groups, we focused on those zones that provided unique functions to the articular cartilage, namely superficial, middle, and calcified zones.

In mature articular cartilage tissue the average modulus increases significantly from 0.08 MPa in the superficial zone to 2.1 and 320 MPa in the middle and calcified zones, respectively.^{5, 24} Conversely, the cell density decreases significantly from 60×10^6 cells/mL in the superficial zone to 20×10^6 and 15×10^6 cells/mL in the middle and calcified zones, respectively.²⁵ Members of transforming growth factor- β (TGF- β) family are used extensively in tissue engineering of articular cartilage to induce chondrogenic differentiation of MSCs. In addition to TGF- β , several other growth factors are known to direct the formation of stratified structure of articular cartilage during native tissue development.^{1, 2, 4, 26} Cultivation of MSCs with TGF- β 1 (3 ng/mL) and BMP-7 (300 ng/mL) induced the synthesis of SZP protein, a marker for chondrocytes in the superficial zone²² whereas IGF-1 did not.^{27, 28} Similar results were obtained when bovine articular cartilage was treated with TGF- β 1 and BMP-7.²⁹ Cultivation of chondrocytes isolated from the middle zone with TGF- β 1 (30 ng/mL) and IGF-1 (100 ng/mL) significantly increased collagen synthesis.³⁰⁻³² TGF- β 1 also contributes to the maintenance of calcified cartilage zone as deletion of TGF- β 1 receptor gene from chondrocytes delayed endochondral ossification³³ and cultivation of bovine hypertrophic chondrocytes with a combination of TGF- β 1 (30 ng/mL) and 3% (w/v) hydroxyapatite (HA) increased matrix deposition and mineralization.^{33, 34}

Strategies that mimic zonal organization of the articular cartilage are more likely to create an engineered tissue with more effective clinical outcome. In this work, we propose a developmentally inspired approach to *in vitro* regeneration of articular cartilage tissue by recapitulating biochemical and biomechanical regulatory factors during cartilage development. In that regard, the superficial zone was simulated in this work by encapsulating 60×10^6 cells/mL human mesenchymal stem cells (hMSCs) in an 80 kPa gel

loaded with 3 ng/mL TGF- β 1 and 100 ng/mL BMP-7; the middle zone was simulated by encapsulating 20×10^6 cells/mL hMSCs in a 2.1 MPa gel loaded with 30 ng/mL TGF- β 1 and 100 ng/mL IGF-1; and the calcified zone was simulated by encapsulating 15×10^6 cells/mL hMSCs in a 320 MPa gel reinforced with nanofibers aligned perpendicular to the articular surface and loaded with 30 ng/mL TGF- β 1 and 3% HA. Although natural gels such as collagen,³⁵ alginate,³⁶ hyaluronic acid, and chitosan³⁷ have been used for cartilage tissue engineering, the stiffness and resorption rate of those matrices cannot be tuned to the specific requirement of each zone. Polyethylene glycol (PEG) gels are inert, non-immunogenic, and compatible with encapsulation of MSCs.^{38, 39} Recently, we reported that PEG macromers chain-extended with short hydroxy acid segments, like L-lactide or glycolide, generate micellar hydrogels with a wide range of stiffness from 1 to 2000 kPa and resorption times from a few days to a few months.⁴⁰ In this work, we used the lactide-chain-extended PEG gels functionalized with acrylate groups (SPELA) to experimentally simulate the synergistic effect of matrix stiffness, cell density, and supplementing the culture medium with growth factors corresponding to those in the superficial, middle and calcified zones on lineage commitment of the encapsulated hMSCs.

Experimental

Materials

Polyethylene glycol (PEG, nominal molecular weights 4.6 kDa), dichloromethane (DCM), N,N-Dimethylformamide (DMF), diisopropylcarbodiimide (DIC), 4-dimethylaminopyridine (DMAP), trifluoroacetic acid (TFA), triisopropylsilane (TIPS), diethyl ether, and hexane were purchased from Acros (Fairfield, OH). The Rink Amide NovaGel™ resin, all Fmoc-protected amino acids, and hydroxybenzotriazole (HOBt) were purchased from Novabiochem (EMD Biosciences, San Diego, CA). Calcium hydride, triethylamine (TEA), paraformaldehyde, 4,6-diamidino-2-phenylindole (DAPI), insulin, penicillin, streptomycin, L-Proline, ascorbic acid, sodium pyruvate, insulin transferrin selenium + ITS Premix, and β -glycerol phosphate were purchased from Sigma-Aldrich (St. Louis, MO). Acetomethoxy derivative of calcein (cAM) and ethidium homodimer (EthD) were purchased from Molecular Probes (Life Technologies, Grand Island, NY). Insulin growth factor-1 (IGF-1) and Transforming growth factor- β 1 (TGF- β 1) were purchased from Lonza (Allendale, NJ) and Bone morphogenetic protein-7 (BMP-7) was purchased from Novus (Littleton, CO). Bovine serum albumin (BSA) was purchased from Jackson ImmunoResearch (West Grove, PA). Dulbecco's phosphate-buffer saline (PBS), trypsin-EDTA, DMEM cell culture medium, fetal bovine serum (FBS), Alexa Fluor 594 Phalloidin, and Quant-it Pico-Green dsDNA reagent kit were purchased from Invitrogen (Carlsbad, CA). Spectro/Por dialysis tube (molecular weight cutoff 3.5 kDa) was purchased from Spectrum Laboratories (Rancho Dominguez, CA). DCM was purified by distillation over calcium hydride. All other solvents were reagent grade and were used as received without further purification.

Synthesis and gelation of SPELA hydrogels

Previous clinical studies indicate that regeneration of the superficial layer of articular cartilage occurs 3-4 weeks post-surgery⁴¹ while full thickness regeneration and complete load bearing occurs at 8-12 weeks post-surgery.^{41, 42} Therefore in this study SPELA gels

with resorption times of 4, 8 and 12 weeks were synthesized to simulate turnover rate of the superficial, middle, and calcified zones, respectively. The SPELA macromers were synthesized in two steps by condensation polymerization of PEG with L-lactide monomer followed by reaction with acryloyl chloride as we previously described.^{38, 43} The number of lactide monomers at each macromer chain-end was varied by changing lactide to PEG molar ratio. Compressive modulus of the gel was tuned to the value corresponding to one of cartilage zones by varying SPELA macromer concentration in aqueous solution as we previously described.^{38, 43} Based on the desired resorption rate and compressive modulus, the second and third columns of Table 1 specify SPELA compositions simulating the superficial, middle, and calcified zones.⁴⁰ The notation "SPELAn (m%)" represents m% SPELA macromer in the hydrogel precursor solution and lactide to PEG ratio of n. Acrylamide-terminated glycine-arginine-glycine-aspartic acid peptide (Ac-GRGD) was synthesized on Rink Amide NovaGel™ resin in the solid phase, purified by high-performance liquid chromatography (HPLC), and characterized by electrospray ionization (ESI) mass spectrometry as we previously described.⁴⁴ Ac-GRGD peptide in the amount of 2wt%, based on SPELA macromer weight, was added to the hydrogel precursor solution to facilitate cell adhesion. The precursor solution was crosslinked by ultraviolet (UV) polymerization with 4-(2-hydroxyethoxy) phenyl-(2-hydroxy-2-propyl) ketone (Irgacure 2959; CIBA, Tarrytown, NY) as photoinitiator to form a hydrogel. The precursor solution was prepared by adding the initiator solution (5 mg initiator in 1 mL PBS) to the solution of SPELA macromer in PBS. Next, the solution was degassed, transferred to a Teflon mold, covered with a transparent glass plate and fastened with clips. Then, the precursor solution was irradiated with a BLAK-RAY 100-W mercury long wavelength (365 nm) UV lamp (Model B100-AP; UVP, Upland, CA) for 200 seconds. Fiber-reinforced hydrogels were fabricated to form matrices with compressive modulus of 320 MPa in which aligned nanofiber microsheets were electrospun from a 10 wt% solution of poly(DL-lactide) (PLA, Durect, Birmingham, AL, 95 kDa MW) in 1, 1, 1, 3, 3, 3-hexafluoro-2-propanol (HFIP, VWR). The injection rate, electrical potential, and needle-to-collector distance was 1.0 mL/h, 20 kV, and 7.5 cm, respectively, and the fibers were collected on a rotating wheel at 1200 rpm as we previously described.^{45, 46} Next, the fiber microsheets were dip-coated in 35 wt% SPELA5 hydrogel precursor solution, wrapped around a 5-mm diameter stainless steel rod, and crosslinked as described above. After crosslinking, the disk-shape samples were cut from the gel using an 8 mm cork borer and the cut samples were equilibrated in PBS for 24 h at 37°C.

A rheometer (TA Instruments, New Castle, DE) was used to measure compressive modulus of the gel samples at a uniaxial displacement rate of 7.5 mm/s. Slope of the linear line fitted to the stress-strain curve at 5-10% strain was taken as compressive modulus (E).⁴⁷ To measure mass loss, the equilibrated gel samples were dried under vacuum at 40°C. Next, the samples were incubated in basal medium (5 mL per sample) without FBS at 37°C and under mild agitation. At each time point, samples were washed and dried under vacuum. The dried sample weights were measured and compared with the initial dry weights to determine fractional mass remaining.³⁸

Encapsulation and cultivation of hMSCs in SPELA hydrogels

Mechanical, biochemical, and cellular requirements for the superficial, middle, and calcified zones of articular cartilage are summarized in Table 1. hMSCs were cultivated at 5000 cells/cm² in a high glucose DMEM medium supplemented with 10% FBS, 100 units/mL penicillin and 100 µg/mL streptomycin (basal medium). All hMSCs were passaged <5 times prior to cell encapsulation according to supplier's instructions (Lanza, Allendale, NJ). For cell encapsulation, the sterile initiator and macromer solutions were mixed, the suspension of hMSCs in 100 µL DMEM was added to the precursor solution and mixed with a pre-sterilized glass rod as we described previously.⁴⁰ For the calcified zone gel, prior to cell addition, 3% (W/V) hydroxyapatite (HA) was suspended in the hydrogel precursor solution by sonication (0.5 h at 20 kHz) as we previously described.⁴⁸ The precursor cell suspension was injected between two sterile microscope glass slides and crosslinked by UV irradiation for 200 s. After crosslinking, the disk-shape samples were incubated in basal medium for 24 h with two medium changes. Next, the medium was replaced with chondrogenic medium consisting of DMEM (4.5 g/mL glucose, 50 µg/mL L-proline, 50 µg/mL ascorbic acid, 0.1 mM sodium pyruvate, 1% v/v insulin-transferrin-selenium (ITS) premix)²¹ and supplemented with the corresponding chondroinductive growth factors to each zone as specified in Table 1. Recent studies indicate that hypertrophy in chondrogenic differentiation of hMSCs is initiated in days 9-12^{8,9} which is marked by a significant increase in Col X expression. Therefore, the medium for hMSCs in the superficial zone gel was supplemented with the appropriate growth factors for 10 days to prevent hypertrophy whereas the medium for hMSCs in the middle and calcified zone gels were supplemented with the appropriate growth factors for 21 days. After day 10, the medium for hMSCs in the superficial zone gel was replaced with basal medium. Cell cultures were incubated at 37°C and 5% CO₂ for 3 weeks.

The experimental groups included hMSCs encapsulated in the gel simulating the superficial zone and cultured in chondrogenic medium supplemented with 3 ng/mL TGF-β1 and 100 ng/mL BMP-7 (hereafter referred to as "Superficial gel"), hMSCs encapsulated in the gel simulating the middle zone and cultured in chondrogenic medium supplemented with 30 ng/mL TGF-β1 and 100 ng/mL IGF-1 (hereafter referred to as "Middle gel"), hMSCs encapsulated in the gel simulating the calcified zone and cultured in chondrogenic medium supplemented with 30 ng/mL TGF-β1 and 10 mM β-glycerol phosphate (hereafter referred to as "Calcified gel"),^{49,50} and the control gel group. The control group was hMSCs encapsulated in the gel simulating the superficial zone (80 kPa modulus) but cultured in basal medium (no TGF-β1 or BMP-7). The cellular hydrogels were cultured for up to 21 days. At each time point, samples were assessed with respect to cell content, total collagen content, GAG, ALP activity, and markers specific to each cartilage zone biochemically, histologically and by mRNA analysis.

Viability of the encapsulated cells

For cell viability, disks were incubated with Acetomethoxy derivative of calcein (cAM) and ethidium homodimer (EthD) (1 µg/mL, Life Technologies, Grand Island, NY) to stain live and dead cells, respectively, as described.³⁸ Stained samples were imaged with an inverted fluorescent microscope (Nikon Eclipse Ti-ε, Nikon, Melville, NY). For imaging the

encapsulated cells, gels were rinsed twice with PBS and fixed with 4% paraformaldehyde for 3 h. After fixation, cells were permeabilized using PBS containing 0.1% Triton X-100 for 5 min, rinsed, and incubated with Alexa 488 phalloidin (1:200 dilution) and DAPI (1:5000 dilution) to stain actin filaments of the cell cytoskeleton and cell nuclei, respectively. Stained samples were imaged with a Nikon Eclipse Ti-E inverted fluorescent microscope as described.⁴⁵

Histology

The day 21 samples were fixed in formalin, embedded in paraffin, and sectioned using a microtome to a thickness of 10 μm as described.⁵¹ An improved cryosection method described by Ruan et al.⁵² based on a protein solution in place of sucrose solution in the dehydration step was used to produce less brittle blocks for sectioning. The sections were stained with Hematoxylin and Eosin (H&E) to ascertain morphology of the encapsulated cells and stained with Alcian blue to image GAG. Whole gels were also stained with Alcian blue and Alizarin red to image the extent of GAG accumulation and mineral deposits, respectively, in hMSC encapsulated gels.¹⁸ Histological sections were imaged with a Nikon Optiphot Epifluorescence System at the same instrument settings.

DNA content and ALP activity

At each time point, the gel samples were homogenized and sonicated to rupture membrane of the encapsulated cells. Double stranded DNA (dsDNA) content of the samples was measured using QuantiGreen assay (Molecular Probes) according to manufacturer's instructions. Samples were homogenized using the kit lysis buffer. 100 μL of the cell lysate was added to the kit's working solution and fluorescence was measured after 4 min incubation at ambient condition with a Synergy-HT plate reader at emission and excitation wavelengths of 485 and 528 nm, respectively.⁴⁴ Measured intensities were correlated to the equivalent amount of dsDNA using a calibration curve constructed with DNA standard solutions.³⁸ ALP activity was assessed using QuantiChrom ALP Assay Kit according to manufacturer's instructions. Briefly 190 μL of the kit's reagent solution containing 10 mM p-nitrophenyl phosphate and 5 mM magnesium acetate was loaded to each well of a 96 well plate, 10 μL aliquot of the homogenized sample was added, and absorbance was recorded at time zero and again after 4 minutes at 405 nm wavelength.⁴⁴ ALP activity was calculated using the equation $[(A_{t=4} - A_{t=0}) / (A_{\text{calibrator}} - A_{\text{ddH}_2\text{O}}) \times 808]$. The measured ALP activities were normalized to DNA content.³⁸

Total collagen and GAG content

Total collagen content was assayed based on selective binding of G-X-Y amino acid sequence of collagen to Sircol dye, according to manufacturer's instructions. Briefly 1 mL of Sircol dye was added to the sonicated cell lysate, incubated for 30 minutes and centrifuged at 10000 rpm for 5 min to separate the collagen-dye complex. After removing supernatant, the collagen-dye complex was mixed with 1 mL Sircol Alkali reagent and absorbance was measured on a plate reader at 555 nm wavelength. The assayed collagen was calibrated using reagent blanks and the kit's standard collagen solution, and normalized to DNA content. GAG production in the gel samples was determined using a dimethylmethylene blue (DMMB) assay. Briefly, the gel samples after washing twice with PBS were digested

with 1 mM EDTA, 2mM dithiothreitol and 300 µg papain in 1 mL of 20 mM sodium phosphate buffer (pH 6.8). After incubation at 60°C for 60 min, iodoacetic acid was added to 10 mM final concentration and the volume was adjusted to 5 mL by addition of 50 mM Tris buffer. The color reagent was prepared by dissolving 16 mg dimethylmethylene blue to 1 L water containing 3 g glycine, 2.37 g NaCl and 95 ml of 0.1 M HCl to the final pH of 3. Next, 2.5 mL of the color reagent was added to 100 µL of each sample and absorbance was recorded after 10 minutes at 525 nm wavelength. The assay was calibrated using reagent blanks and serial dilutions of the standard solution of whale chondroitin sulfate, and normalized to DNA content.¹⁸

mRNA analysis

Markers included SOX-9 as master regulator of chondrogenesis,⁵³ SZP as marker for the superficial zone,²² Col II and AGC as ECM components highly expressed in the middle zone,¹⁸ and Col X and ALP as markers for cartilage hypertrophy in the calcified zone.³ At each time point, total RNA of the homogenized samples was isolated using Trizol⁴⁴ and genomic DNA was removed using Deoxyribonuclease I (Invitrogen) according to manufacturer's instructions. After converting RNA (250 ng) using Promega reverse transcription system (Madison, WI), the converted cDNA was amplified by RT-qPCR with SYBR green RealMasterMix (Eppendorf, Hamburg, Germany) using Bio-Rad CFX96 PCR system (Bio-Rad, Hercules, CA) and the appropriate gene specific primers. Primers were designed and selected using Primer3 web-based software as described.⁵⁴ The list of primer sequences are provided in Table 2. Data were analyzed using $\Delta\Delta C_t$ Real time analysis method.³⁸ mRNA expressions were normalized against GAPDH as a house keeping reference gene and fold changes were compared to those in the same group at day zero.

Statistical analysis

All experiments were done in triplicate and quantitative data expressed as means \pm standard deviation. Significant differences between groups were evaluated using a two-way ANOVA with replication test, followed by a two-tailed Students t-test. A value of $p < 0.05$ was considered statistically significant.

Results

Characterization of the hydrogels

Figure 1A shows degradation rate of SPELAN hydrogels at 37°C. Mass loss of SPELA5, SPELA7.5 and SPELA10 hydrogels after 7 days of incubation was 9%, 13% and 18%, respectively. Further, mass loss of the hydrogels was between 17% and 32% after 14 days and between 29% and 47% after 21 days (Figure 1A). We previously showed that water content of SPELA gels did not change significantly with lactide segment length.⁴⁰ Therefore, the 2- and 1.5-fold higher mass loss of SPELA10 and SPELA7.5, respectively, compared to SPELA5 after 7 days was due to lactide segment length, not differences in water content. The effect of macromer concentration on compressive modulus of SPELAN hydrogels is shown in Figure 1B. The increase in modulus with concentration was attributed to higher crosslink density of the gels as modulus was directly related to density of active crosslinks in the gel network.⁵⁵ The gel with 15 wt% SPELA10 concentration simulated

compressive modulus of the superficial zone (80 kPa, open square). Likewise, the gel with 50 wt% SPELA7.5 concentration simulated modulus of the middle zone (2.1 MPa, open diamond). The gel with 35 wt% SPELA7.5 concentration reinforced with PLA nanofibers simulated modulus of the calcified zone (320 MPa, open circle).

Cell viability and histological characterization of the cellular gels

Images in Figure 2A-D compare live (stained green) and dead (stained red) stained hMSCs encapsulated in Superficial (B), Middle (C), and Calcified (D) gels after 21 days of incubation in chondrogenic medium with the control group cultivated in basal medium (A). The inset in Figure 2D shows cell clusters around nanofibers. Images in Figure 2E-H compare phalloidin (cytoskeleton, purple) and DAPI (nuclei, blue) stained hMSCs encapsulated in Superficial (F), Middle (G) and Calcified (h) gels after 21 days of incubation in chondrogenic medium with the control group cultivated in basal medium (E). Insets in Figure 2E-H show morphology of the encapsulated cells at higher magnification. hMSCs encapsulated in Superficial (Figure 2E) and Middle (Figure 2F) gels were rounded consistent with their morphology in the native tissue.^{4, 5} Figure 2H shows morphology of the cells in Calcified gel in merged bright and dark field image of the whole gel. The inset in Figure 2H shows extensive cell clustering adjacent to aligned nanofibers in Calcified gel.

Images in Figure 3A-D compare GAG stained histological sections (Alcian blue) of hMSCs encapsulated in Superficial (B), Middle (C) and Calcified (D) gels after 21 days incubation in chondrogenic medium with the control group cultivated in basal medium (A). Figures 3E-H compare H&E stained histological sections of hMSCs encapsulated in Superficial (F), Middle (G) and Calcified (H) gels after 21 days incubation in chondrogenic medium with the control group cultivated in basal medium (E). The rounded cells increased in size from Superficial (Figure 3F) to Middle (Figure 3G) and Calcified (Figure 3H) gels compared to that cultivated in basal medium (Figure 3E). The cells in Calcified gel were in isolated Lacunae consistent with their morphology in the native calcified layer of articular cartilage tissue. However, the observed cell clusters in images of the whole Calcified gel in Figure 2 (D and H) were not observed to the same extent in histological sections of Figure 3. We believe the fibers and their associated cell clusters, due to higher fiber stiffness, pulled out of the hydrogel matrix during sectioning, which led to unmatched microscopic images in Figures 2 and 3 for Calcified gel. Coburn et al.⁵⁶ previously reported changes in cell morphology at the fiber/gel interface and McCullen et al.²⁰ reported alignment of cell clusters along fiber direction. Images in Figure 4A-D compare GAG staining of the whole Superficial (B), Middle (C) and Calcified (D) gels after 21 days incubation in chondrogenic medium with the control cultivated in basal medium (A). GAG content of the whole samples progressively increased from Superficial (B with lighter blue) to Middle (C) and Calcified (D with darker blue) gels compared to those gels cultured in basal medium (A with the lightest blue). Images in Figure 4E-H compare Alizarin red staining for the whole Superficial (F), Middle (G), and Calcified (H) gels after 21 days incubation in chondrogenic medium with the control cultivated in basal medium (E). Figure 4H shows high mineral deposits in Calcified gel with minor amounts in Superficial and Middle gels (Figure 4F-G).

Analysis of mRNA expressions

mRNA expression of AGC, Col II, Col X, ALP, SZP, and Sox-9 for hMSCs encapsulated in the gels are shown in Figures 5A to 5F, respectively. mRNA expression of AGC and Col II as the main components of cartilage ECM (Figures 5A and 5B, respectively) showed an increasing trend with time for all groups. hMSCs encapsulated in Middle gel showed highest expression of AGC and Col II expression level with 112 and 45-fold increase in expression on day 21, respectively. Col X mRNA expression of hMSCs in Superficial, Middle and Calcified gels (Figure 5C) was 0.96 ± 0.04 , 4.3 ± 0.3 , and 16 ± 1 on day 14, respectively, and increased to 1.05 ± 0.03 , 7.3 ± 0.5 , and 26 ± 2 on day 21. On day 14 and 21, Calcified and Superficial gels had highest and lowest expression of Col X, respectively. Col X mRNA expressions of hMSCs were in agreement with ALP mRNA expressions as shown in Figure 5D. Calcified gel showed highest ALP expression level for all time points whereas Superficial and Middle gels had ALP expression levels similar to the control for all time points. Figure 5E shows mRNA expression of SZP, as a marker of the superficial zone, for the encapsulated hMSCs. Superficial gel showed highest SZP expression level relative to Middle and Calcified gels and the expression level had an increasing trend with incubation time during the exposure period of BMP-7 and TGF- β (first 10 days). There was no significant difference in SZP expression of hMSCs in Calcified gel relative to the control for all time points.

SOX-9 mRNA expression of hMSCs as a major regulator of chondrogenesis showed a biphasic trend with incubation time for all gels. SOX-9 expression increased significantly in the first 14 days of incubation and then decreased on day 21 (Figure 5F). hMSCs in Superficial gel showed highest SOX-9 expression level for all time points relative to Middle and Calcified gels.

Biochemical analysis of cellular gels

DNA (A), GAG (B), total collagen content (C) and ALP activity (D) of hMSCs encapsulated in Superficial (L1), Middle (L2), and Calcified (L3) gels with incubation time are shown in Figures 6A to 6D, respectively. DNA content of the hMSCs decreased slightly with incubation time for all gels (Figure 6A). GAG content of the hMSCs showed an increasing trend with incubation time for all gels with higher GAG content in Middle and Calcified gels relative to Superficial gel (Figure 6B). Total collagen content of the hMSCs showed an increasing trend with incubation time for all gels with higher collagen content in Middle gel relative to Superficial and Calcified gels (Figure 6C). ALP activity of hMSCs also showed an increasing trend with incubation time with Calcified gel having orders of magnitude higher ALP activity relative to Superficial and Middle gels (Figure 6D). The biochemical results for GAG/collagen (Figure 6), in particular higher GAG/collagen of the hMSCs in Middle gel, were consistent with mRNA expressions in Figure 5. hMSCs encapsulated in the gel simulating the calcified zone showed highest increase in ALP activity with time relative to Superficial and Middle gels reaching 1030 IU/mg DNA on day 21 (Figure 6D).

Discussion

The structure of articular cartilage tissue is anisotropic and spatially variant with complex interactions between resident cells, ECM, and soluble growth factors. This anisotropy is integral to the function of cartilage tissue including lubrication, compressive strength, and a rigid interface for load transmission to the underlying bone.^{1, 4, 20} Despite the importance of zonal organization of articular cartilage on tissue function, a viable approach to zonal regeneration of the tissue is not established. In this work, we propose a developmental approach to zonal regeneration of articular cartilage by recapitulating the contribution of biochemical, mechanical, and cellular regulatory factors during cartilage development. The organization of articular cartilage is initiated from the early stages of development by stimulation of undifferentiated MSCs in the lateral plate mesoderm followed by condensation, lineage commitment and eventually differentiation into mature chondrocytes (Figure 7A).^{2, 57} Wu et al.⁵⁷ isolated different layers of articular cartilage from femoral epiphysis of 5, 8, 12, and 17 weeks old human embryos by laser capture dissection method and measured changes in the expression pattern of surface markers. Results of his study show that de novo chondrocyte condensation of MSCs is observable up to 11-12 weeks of gestation and fetal articular cartilage is not a homogenous tissue from week 12.⁵⁷ Chondrocytes isolated from the superficial zone of the condensates were significantly different from the more differentiated and hypertrophic chondrocytes in the middle zone. Further, his results show that the vast majority of isolated chondrocytes from femoral epiphysis at week 17 of human development were zonally specified.⁵⁷ Subsequently, during fetal and postnatal development and in puberty,⁵⁸ the developing chondrocytes secrete specific marker proteins to remodel the zonal structure of articular cartilage (Figure 7B).^{2, 59} Results of Wu et al. also indicate that BMP-7 and IGF-1 play critical roles during early cartilage development.⁵⁷

In the superficial zone, BMP-7 in combination with TGF- β 1 and low matrix stiffness up-regulate the expression of SZP in resident chondroprogenitor cell population.^{22, 60} In the middle zone, IGF-1 as an anabolic growth factor in combination with TGF- β 1 and a moderately stiff matrix have a direct effect on up-regulation of Col II and GAG in pre-hypertrophic chondrocytes.³⁰ The calcified zone contains developmentally mature chondrocytes expressing proteins associated with hypertrophy such as Col X and ALP in a mineralized matrix.^{3, 4} The combination of TGF- β 1 signaling and high matrix stiffness is essential for the formation of calcified cartilage.^{5, 18} We simulated the three zones of articular cartilage with hydrogels mimicking mechanical and biochemical signaling factors involved in articular cartilage tissue development and homeostasis. Specifically, stiffness and degradation of the constructs as biophysical cues, concentrations of three growth factors (TGF- β 1, IGF-1 and BMP-7) as biochemical cues, and cell density as the cellular cue were used to control differentiation of the encapsulated human MSCs to the chondrocyte lineage corresponding to superficial, middle, and calcified zones. Results of previous clinical studies indicates that regeneration of superficial zone occurs 3-4 weeks post-surgery⁴¹ while regeneration of load-bearing full-thickness cartilage happens after 8-12 weeks.^{41, 42} Therefore, SPELA gels with resorption times of 4, 8 and 12 weeks were used to simulate the turnover rate of superficial, middle and calcified zones, respectively. Furthermore,

compressive modulus of the gels were adjusted to 0.08, 2.1 and 320 MPa based on biomechanics of the superficial, middle and calcified zones in the native articular cartilage tissue, respectively.⁵

DNA content of the hMSCs in Superficial, Middle, and Calcified gels decreased slightly with incubation time which was attributed to cell differentiation leading to telomere shortening and decrease in cell replication.⁶¹ The decrease in DNA content was consistent with the previously reported reduction in cell density with chondrogenic differentiation of MSCs in alginate gels with incubation time.⁶² mRNA results in Figure 5 showed an increase in the expression of chondrogenic markers AGC and Col II of the hMSCs with incubation time for all three gels which was attributed to the exposure of encapsulated hMSCs to TGF- β . TGF- β stimulates chondrogenic differentiation of MSCs in a 3D culture environment^{4, 63-65} in crosstalk with Wnt/ β -catenin pathway through Smad 2/3 phosphorylation. Next, phosphorylated Smad 2/3 forms a complex with Smad 4 and translocates to the nucleus to serve as a transcription factor for activation of Collagen II and AGC genes. In addition, Smad 3 and 4 interact with β -catenin⁶⁶ and form a complex for protection of β -catenin from ubiquitin-proteasome dependent degradation and mediation of β -catenin nuclear translocation. Smad3-mediated stability of β -catenin activates down-stream chondrogenic target genes including Col II and AGC.⁶⁶ TGF- β and Wnt/ β -catenin pathways synergistically regulate expression of downstream chondrogenic target genes through transcription factors TCF/ Lef1.⁶⁷

hMSCs in the gel simulating the middle zone (Middle gel) had highest AGC and Col II expression with 112- and 45-fold increase on day 21 (Figure 5A, 5B), respectively, which was consistent with the biochemical results in Figure 6 and zonal composition of AGC and Col II in the native cartilage tissue. The Middle gel had significantly higher Col II and AGC expression relative to Superficial and Calcified gels which could be attributed to the combined effect of TGF- β 1 and IGF-1 growth factors and moderately stiff gel (2.1 MPa). IGF-1 has been shown to stimulate synthesis of proteoglycans and collagen in pre-hypertrophic chondrocytes of the middle zone *in vivo* and *in vitro*.^{7, 30, 68} In an *in vivo* study to investigate the effect of IGF-1 on prenatal and postnatal cartilage tissue development, cartilage specific IGF-1 knockout mice (Cart/igf-1^{-/-}) died shortly after birth and showed disorganized chondrocyte columns, decreased cell proliferation, and increased apoptosis in their cartilage tissue.⁶⁹ Further, exposure of 3D MSC cultures or chondrocyte grafts to IGF-1 significantly increased GAG and collagen synthesis,⁷⁰ and IGF-1 stimulated synthesis of proteoglycans in pre-hypertrophic chondrocytes *in vitro*.^{30, 71}

Expression of SZP and Sox-9 were used as markers for differentiation of hMSCs to chondrocyte lineage of the superficial zone. The Superficial gel had significantly higher expression of SZP relative to Superficial and Middle gels which could be attributed to the combined effect of TGF- β 1 and BMP-7 growth factors and relatively low matrix compressive modulus (80 kPa).^{22, 72} To engineer superficial zone chondrocytes, Coates co-cultured alginate-encapsulated MSCs with explants from the superficial or middle/deep zone of bovine articular cartilage.⁷³ Co-cultures with the superficial zone explants without the addition of TGF- β 3 to the medium resulted in up-regulation of chondrogenic markers Sox-9 and Col II and modest increase in SZP in the encapsulated MSCs after 21 days compared to

standard chondrogenic medium. Remarkably, the middle/deep zone explants did not induce expression of chondrogenic markers in co-cultured MSCs.⁷³ The results of Coates et al. highlight the importance of growth factors secreted in the superficial zone on chondrogenesis whereas our results highlight the importance of BMP-7 in combination with TGF- β 1 for directing differentiation of hMSCs to the superficial zone phenotype. Another factor in our experiments that may have contributed to the high SZP expression (21 fold increase) in Superficial gel was the relatively high cell density of hMSCs (60×10^6 cells/mL) consistent with that in the native cartilage tissue.⁷⁴ GAG content of the superficial zone in native articular cartilage is lower than the remaining zones, consistent with our results (Figure 6B).⁷⁵ However contrary to our results (Figure 6C), collagen content of the superficial zone is higher in the native tissue compared to the remaining zones⁷⁵ and the orientation of collagen fibers is parallel to the cartilage surface.⁵⁸ Julkunen et al. reported significant changes in the orientation of collagen fibers in femoral medial condyles of rabbits with postnatal development.⁵⁸ Collagen content and orientation in the superficial zone is regulated by external load during postnatal skeletal maturation to limit deformation of the relatively compliant superficial zone in the direction parallel to the cartilage surface.⁷⁶ Since the cell-seeded gels in our experiments were not loaded externally, we simulated compressive stiffness perpendicular to the cartilage surface as opposed to tensile modulus parallel to the surface. As a result, collagen content of the superficial zone was lower than the remaining zones. Steward et al.⁷⁷ investigated the effect of matrix stiffness and hydrostatic pressure, with or without integrin blocker, on matrix production and gene expression of MSCs encapsulated in agarose gels. They reported that stiffer gels suppressed cartilage matrix production in chondrogenic medium but matrix synthesis returned to the levels in softer gels by blocking integrin receptor-ligand interaction with RGDS peptide.⁷⁷ It should be noted that the concentration of agarose in Steward et al. experiments ranged 1-4%⁷⁷ which corresponded to moduli of 10-70 kPa.⁷⁸ Conversely in our experiments, production of chondrogenic matrix (total collagen and GAG) by the hMSCs was suppressed in the compliant Superficial gel compared to stiff Middle and Calcified gels. The discrepancy between our results and those of Steward et al.⁷⁷ could be attributed to significantly higher modulus of the gels (80 kPa, 2.1 MPa, 320 MPa) in our experiments versus <70 kPa in Steward et al.⁷⁷ and absence of hydrostatic pressure in our experiments.

It is well established that TGF- β 1 signaling regulates Sox-9 expression in all chondroprogenitor cells and developing chondrocytes but not in mature hypertrophic chondrocytes.^{2, 79} Further, It has been shown that inactivation of BMP receptors in mice leads to loss of Sox-9 expression.⁸⁰ Since both TGF- β 1 and BMP-7 were used, the high expression of Sox-9 in Superficial gel could be contributed to TGF- β 1/BMP-7 pathway through downstream Smad and TAK1 (TGF- β 1 activated kinase) factors.⁸⁰

The Calcified gel had significantly higher Col X and ALP expression relative to Middle and Superficial gels which could be attributed to the combined effect of high matrix modulus (320 MPa), TGF- β 1, and HA. Khanarian et al. observed highest ALP activity in pre-hypertrophic chondrocytes derived from the deep zone of bovine articular cartilage and seeded in agarose gel with the addition of 3% HA.⁸¹ Based on Huang et al. results, HA concentrations higher than 75 μ g/mL induced expression of ALP in 3D cultures of rat MSCs at both mRNA and protein levels.⁸² In addition, mechanical strain along with HA addition

had a positive effect on mRNA expression of ALP and matrix mineralization.⁸² Several studies have shown the significance of matrix stiffness on ALP expression. Chatterjee et al. detected no mineralization in MC3T3-E1 osteoblasts encapsulated in polyethylene glycol (PEG) gels with <100 kPa compressive modulus.⁸³ The relatively low cell density of Calcified gel could have also contributed to up-regulation of Col X and ALP expression of the hMSCs in Calcified gel. Lower cell density can lead to lower oxygen consumption rate and hence higher oxygen partial pressure in the gel, thus reducing the risk of hypoxia. Leijten et al. has shown that hypoxia suppresses markers of hypertrophy and endochondral ossification in fetal mouse tibia explants.⁸⁴ Further, Eslaminejad and Nadri reported significantly higher mineralization by murine MSCs in 2D culture at lower cell seeding density.⁸⁵ In fabrication of Calcified gel, nanofiber microsheets were wrapped around a metallic rod sufficiently tight such that the suspended hMSCs came partially in contact with the surface of nanofibers. In other words, cells were encapsulated in the gel but could sense the fiber stiffness. Therefore, the fibrous nature of Calcified gel could have also contributed to hypertrophy in encapsulated hMSCs as it has been reported that a 3D apatite-fiber scaffold structure enhanced endochondral ossification by bone marrow-derived stem cells.⁸⁶

Previous studies revealed that substrate stiffness can direct lineage commitment of MSCs through a modulus-sensing mechanism mediated by integrin engagement and myosin-based contraction of actin cytoskeleton.⁸⁷ Shih et al. observed up-regulation of α_2 integrin and kinase activation of ROCK, FAK, and ERK1/2 of MSCs in stiff matrices.⁸⁸ Further, matrix rigidity affects phosphorylation of multiple proteins such as Akt, β -catenin, Jnk, and STAT5a.⁸⁹ Akt activation and increased β -catenin expression are implicated in the induction of skeletal phenotype,^{90, 91} whereas STAT5a has been shown to participate in up-regulation of 1,25-dihydroxyvitamin D3 which is critical to mineral deposition.^{92, 93}

Matrix stiffness plays an important role in regeneration of all cartilage zones. In a previous study, total collagen content of the chondrocytes encapsulated in PEG gels decreased with increasing compressive modulus from 30 to 950 kPa in the absence of chondrogenic growth factors whereas expression of Col II peaked in moderately crosslinked gels with compressive modulus of 360 kPa.⁹⁴ The lower optimum compressive modulus for Col II expression in the previous study⁹⁴ compared to that in the middle zone of the native articular cartilage could be attributed to suboptimal *in vitro* conditions with respect to biochemical signaling. Expression of TGF- β is regulated at multiple levels some of which intersect with mechanotransduction pathways.⁸³ Previous studies indicate that TGF- β 1 and optimum substrate stiffness have equivalent inductive effects on ATDC5 chondroprogenitor cells but the effect is synergistic when both are applied.⁸³ Previous studies also indicate that integrin receptors and cell-generated tension mediate the conversion of TGF- β receptors from a latent to an active form.⁹⁵ After receptor activation, TGF- β signaling propagates through multiple pathways including Rho/Rho-associated protein kinase (Rock) and mitogen-activated protein kinase (MAPK) pathways, and the canonical effectors of TGF- β signaling such as receptor-regulated Smad 2, 3 and 4, which leads to up-regulation of Sox-9 and Col II expression.^{96, 97}

Recently, Nguyen et al. used different combinations of PEG, chondroitin sulfate, and hyaluronic acid gels with compressive modulus of 100, 120 and 300 kPa, respectively, and

equal TGF- β concentration and MSC density in all three layers to regenerate superficial, middle and deep zones of articular cartilage.⁹⁸ Chondroitin sulfate and hyaluronic acid provided ECM signaling to the encapsulated MSCs. The calcified zone which has orders of magnitude higher modulus relative to the deep zone was not simulated in that study.⁹⁸ Considering that only gradients in matrix stiffness and ECM signaling were used to guide differentiation of MSCs in Nguyen experiments, he reported highest expression of Col II and GAG by MSCs encapsulated in the gels simulating superficial and deep zones and a significant increase in Col X expression in all three gels. Our results with gradients in cell density, matrix stiffness, and growth factors (TGF- β , BMP-7, and IGF-1) showed highest expression of Col II and AGC in Middle gel and highest Col X expression in Calcified gel, consistent with the observed zonal expression of Col II, AGC, and Col X in the native articular cartilage tissue.^{3, 5, 18}

Steele et al. generated bilayer scaffolds by electrostatic deposition of PCL microfibers onto porous scaffolds produced by porogen-leaching.²¹ After seeding the bilayer scaffolds with chondrocytes and culturing the constructs in chondrogenic medium supplemented with 10 ng/mL TGF- β 3, he observed higher sulfated glycosaminoglycan (sGAG) production in the zones with aligned fibers. However unlike our results, he observed uniform expression of SZP in all zones.²¹ McCullen et al. recently fabricated tri-laminar scaffolds composed of a layer of aligned fibers parallel to the scaffold surface laminated to two other randomly-oriented fiber layers to mimic microstructure and stiffness of the superficial, middle, and deep zones of articular cartilage, respectively.²⁰ Col II mRNA expression of the chondrocytes seeded on the tri-laminar scaffolds increased with incubation time but there was no significant difference in Col II expression and GAG accumulation between the scaffold layers.

In this work, we focused on the effect of combined gradients in cell density, matrix stiffness, and zone-specific growth factors on differentiation and marker expression of hMSCs. Due to the multifactorial nature of the study, it was not possible to decouple the effect of individual factors (matrix stiffness, cell density, growth factors) on zone-specific differentiation and marker expression of the hMSCs. Inclusion of control groups for the effect of individual factors would have resulted in a very large number of samples. Nonetheless, there are very few studies on combined effect of biochemical (growth factor), mechanical (matrix stiffness), and cellular (cell density) factors on differentiation of human MSCs to zone-specific chondrocytes. We recognize that zonal organization of articular cartilage is regulated by gradients of other equally important factors including tissue strain, oxygen tension, and glucose. Thorpe et al.⁹⁹ confined MSC-encapsulated constructs radially to half their thickness and applied dynamic compression to create gradients in oxygen tension and strain in the direction of construct thickness. The gradients (low oxygen tension in the bottom and high strain at the top) increased GAG in the bottom, increased collagen accumulation at the top, and suppressed hypertrophy and calcification throughout the construct, which led to the formation of an engineered tissue mimicking some aspects of the zonal organization of articular cartilage. Spitters et al. cultivated bovine chondrocytes in a gradient of glucose concentrations and found that GAG production rate was highest at 5 mM glucose level.¹⁰⁰ Recapitulating the gradients in all regulatory signals including matrix stiffness, cell density, growth factors, tissue strain, oxygen tension, and glucose

concentration in a tissue engineered cell-seeded matrix has the potential to regenerate zonal complexity of the articular cartilage. In summary, our results demonstrate that zonal structure of the native articular cartilage can be generated by adopting a developmental approach with gradients in cell density, matrix stiffness, and zone-specific growth factors.

Conclusions

We investigated the combined effect of cell density, matrix stiffness, and zone-specific growth factors within a 3D PEG-based hydrogel microenvironment on chondrogenic differentiation of encapsulated hMSCs into phenotypes corresponding to the superficial, middle, and calcified zones of articular cartilage. The superficial zone was simulated by encapsulating 60×10^6 cells/mL hMSCs in an 80 kPa SPELA gel loaded with 3 ng/mL TGF- β 1 and 100 ng/mL BMP-7. The middle zone was simulated by encapsulating 20×10^6 cells/mL hMSCs in a 2.1 MPa SPELA gel loaded with 30 ng/mL TGF- β 1 and 100 ng/mL IGF-1. The calcified zone was simulated by encapsulating 15×10^6 cells/mL hMSCs in a 320 MPa gel reinforced with nanofibers aligned perpendicular to the gel surface and loaded with 30 ng/mL TGF- β 1 and 3% HA. hMSCs in the simulated superficial zone gel showed up-regulation of Sox-9, the early-stage marker of chondrogenesis, and SZP whereas hMSCs in the simulated calcified zone gel showed up-regulation of Col X, the late-stage marker of hypertrophic chondrocytes, and ALP. Further, hMSCs in the simulated middle zone gel had highest expression of chondrogenic markers Col II, AGC, and GAG among all three gels. The results of this work can potentially lead to the design of more effective multilayer grafts with mechanical, biochemical, and cellular gradients for the treatment of articular cartilage defects.

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Insight, innovation, integration

Strategies that mimic zonal organization of the articular cartilage with complex interactions between resident cells, extracellular matrix and soluble growth factors are more likely to create an engineered tissue with more effective clinical outcome. A developmentally inspired approach is proposed to *in vitro* regeneration of articular cartilage tissue by recapitulating biochemical, mechanical, and cellular regulatory factors during cartilage development. Combination of TGF- β 1 and BMP-7 in a soft gel with a relatively high cell seeding density stimulated chondrogenic differentiation of human mesenchymal stem cells (hMSCs) to the superficial zone phenotype of articular cartilage. Conversely, combination of TGF- β 1 and hydroxyapatite in a stiff gel with a relatively low cell seeding density stimulated chondrogenic differentiation of hMSCs to the calcified zone phenotype. This work could potentially enable the design of effective multilayer grafts as a more sustainable solution to regeneration of articular cartilage defects.

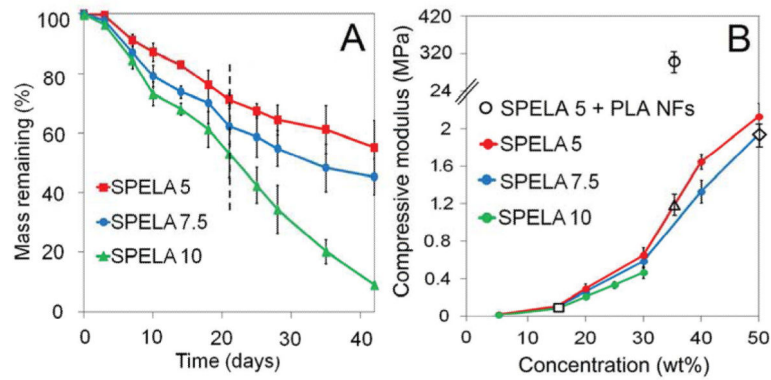


Figure 1.

A) Degradation of SPELA5 (red), SPELA7.5 (blue), and SPELA10 (green) gels with incubation time in basal medium at 37°C. The dashed line shows remaining mass of the gels after 21 days of incubation. B) Compressive modulus of SPELA5 (red), SPELA7.5 (blue), and SPELA10 (green) gels versus SPELA macromer concentration. The square shows modulus of 15% SPELA10 gel simulating the superficial zone; the triangle and circle show modulus of 35% SPELA5 gel before and after reinforcement with nanofibers aligned perpendicular to the gel surface, respectively, to simulate the calcified zone; and the diamond shows modulus of 50% SPELA7.5 gel simulating the middle zone. Error bars correspond to means \pm 1 SD for n=3.

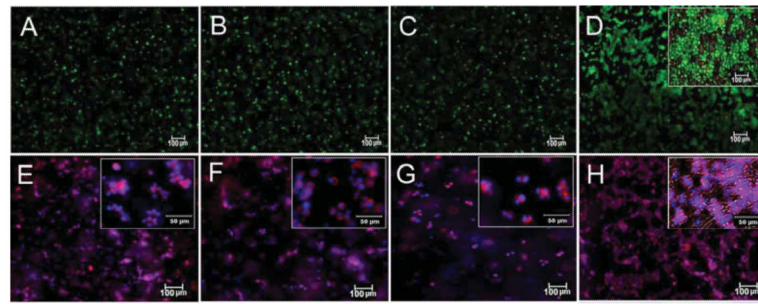


Figure 2.

Calcein AM (live, green) and EthD (dead, red) staining of the hMSCs encapsulated in Superficial (B), Middle (C), and Calcified (D) gels and incubated in chondrogenic medium supplemented with zone-specific growth factors. Phalloidin (purple for cytoskeleton) and DAPI (blue for cell nucleus) images of the hMSCs encapsulated in Superficial (F), Middle (G), and Calcified (H) gels and incubated in chondrogenic medium supplemented with zone-specific growth factors. Images A and E are for hMSCs encapsulated in the 80 kPa gel and incubated in basal medium as the control. The insets are higher magnification images.

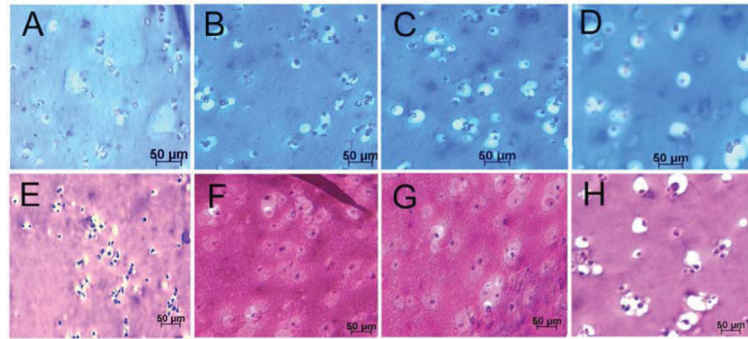


Figure 3.

Alcian blue (GAG) stained histological sections of the hMSCs encapsulated in Superficial (B), Middle (C), and Calcified (D) gels and incubated in chondrogenic medium supplemented with zone-specific growth factors. H&E stained sections for the hMSCs encapsulated in Superficial (F), Middle (G), and Calcified (H) gels and incubated in chondrogenic medium supplemented with zone-specific growth factors. Images A and E are for the hMSCs encapsulated in the 80 kPa gel and incubated in basal medium as the control.

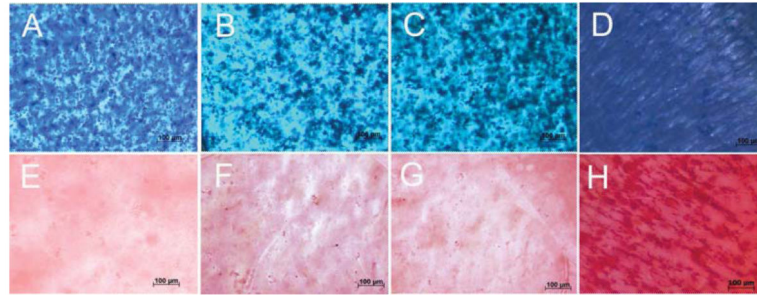


Figure 4.

Alcian blue (GAG) staining of whole gels to showing extent of GAG accumulation for the hMSCs encapsulated in Superficial (B), Middle (C), and Calcified (D) gels and incubated in chondrogenic medium supplemented with zone-specific growth factors. Alizarin red staining of whole gels showing mineral deposition of the hMSCs encapsulated in Superficial (F), Middle (G), and Calcified (H) gels and incubated in chondrogenic medium supplemented with zone-specific growth factors. Images A and E are for hMSCs encapsulated in the 80 kPa gel and incubated in basal medium as the control.

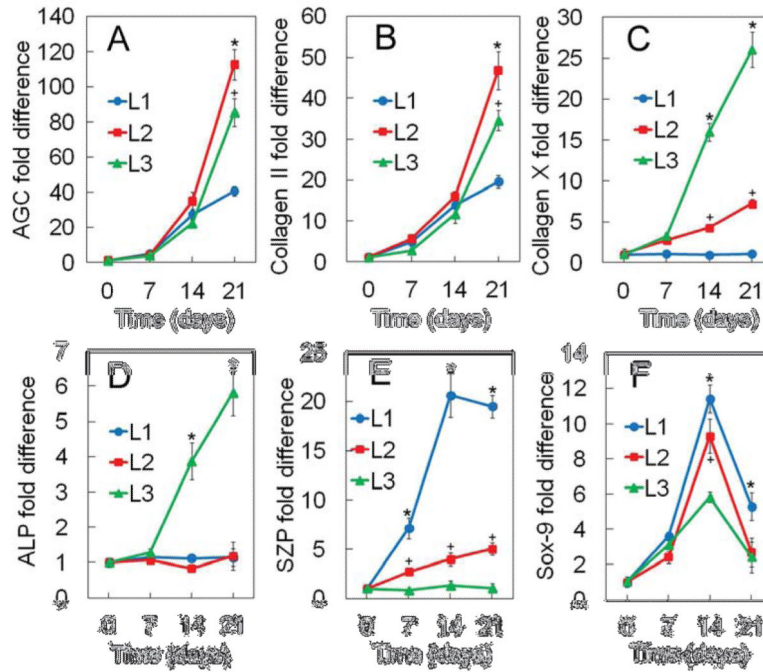


Figure 5. mRNA expression of AGC (A), Col II (B), Col X (C), ALP (D), SZP (E), and Sox-9 (F) for the hMSCs encapsulated in Superficial (L1, blue), Middle (L2, red), and Calcified (L3, green) gels with incubation time in chondrogenic medium supplemented with zone-specific growth factors. A star indicates a statically significant difference between the test and all other groups at each time point. Error bars correspond to means±1 SD for n=3.

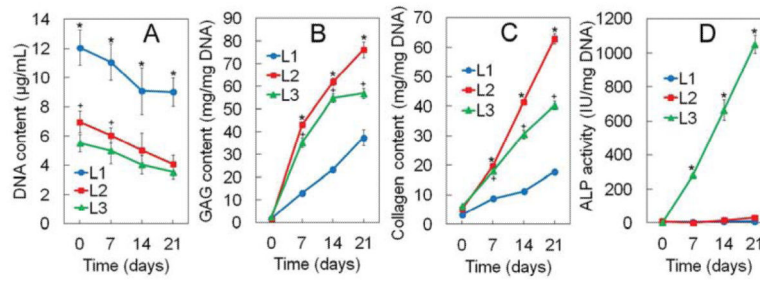


Figure 6.

DNA content (A), GAG content (B), total collagen content (C), and ALP activity (D) for the hMSCs encapsulated in Superficial (L1, blue), Middle (L2, red), and Calcified (L3, green) gels with incubation time in chondrogenic medium supplemented with zone-specific growth factors. A star indicates a statically significant difference between the test and all other groups at each time point. Error bars correspond to means±1 SD for n=3.

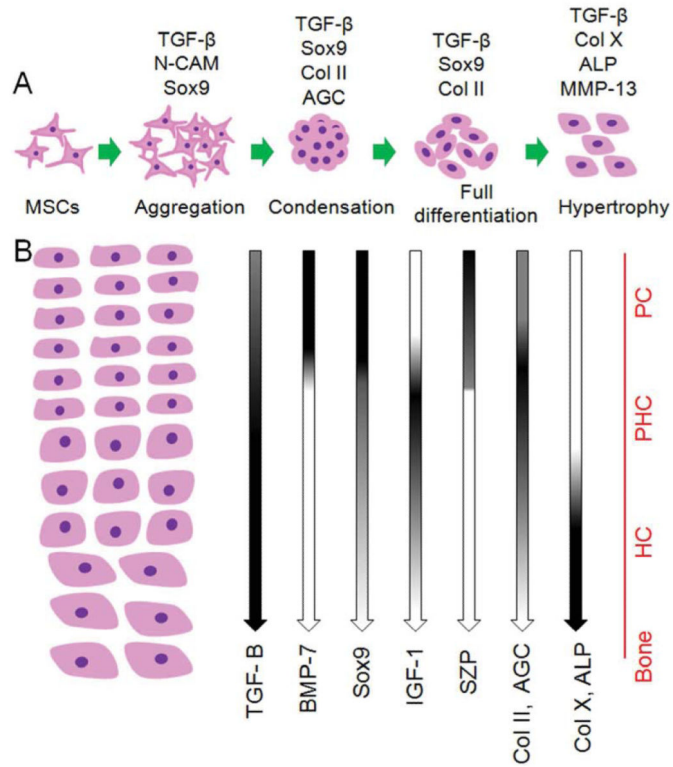


Figure 7. Schematic representation of the events during chondrogenesis. (A) Chondrogenesis consists of a dynamic series of events including stimulation of mesenchymal stem cells followed by mesenchymal aggregation and condensation and finally cartilage tissue differentiation and maturation. Expression pattern of specific cartilage genes at different stages of tissue development are shown. (B) Overview of the effect of specific growth factors on zonal organization of the articular cartilage tissue including superficial, middle/deep and calcified cartilage zones. The spatial distribution of specific growth factors in each zone is illustrated. The distribution of cartilage specific proteins is indicated by gradients of gene expression.^{2, 59} PC, PHC, and HC stand for pre-chondrocytes, pre-hypertrophic chondrocytes, and hypertrophic chondrocytes, respectively.

Table 1

Gel composition and degradation time, matrix stiffness, type, concentration and exposure time of growth factors, and hMSC cell density for simulating superficial, middle, and calcified zones of articular cartilage.

Simulated cartilage zone	Gel composition and degradation time	Matrix stiffness	Growth factors (GF)	GF Exposure time	Cell density (cells/mL)
Superficial	SPELA10 (15%) 4weeks	80kPa/	TGFβ1 (3 ng/mL) BMP-7 (100 ng/mL)	10 days	60×10 ⁶
Middle	SPELA7.5 (50%) 8 weeks	2.1 MPa/	TGFβ1 (30 ng/mL) IGF-1 (100 ng/mL)	21 days	20×10 ⁶
Calcified	PLA/SPELA5 (35%) fiber composite 3% HA 12 weeks	320 MPa/	TGFβ1 (30 ng/mL)	21 days	15×10 ⁶

Table 2

Reverse and forward sequences for PCR primers.

PCR Primer	Forward	Reverse
SZP	5'AACCACCACCAGACCTAACCAA3'	5'CAGCACCACCTGCATCTTCA3'
Sox-9	5'CCCAACAGATCGCCTACAGT3'	5'GAGTTCTGGTGGTCGGTGTAGTC3'
AGC	5'TCGAGGACAGCGAGGCC3'	5'TCGAGGGTGTAGCGTGTAGAGA3'
Col II	5'AGCCTGGTGCATGGGTTTC3'	5'GTCCCTTCTCACCAGCTTTGC3'
Col x	5'CTGTATAAGAATGGCACCCTGTA3'	5'GCACTCCCTGAAGCCTGATC3'
ALP	5'ATGGGATGGGTGTCTCCACA3'	5'CCACGAAGGGGAACTTGTC3'
GAPDH	5'CATGACAACCTTGGTATCGTGG3'	5'CCTGCTTACCACCTTCTTG3'

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