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The bio in the ink: cartilage regeneration with bioprintable hydrogels and articular cartilage-derived progenitor cells

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Abstract

Cell-laden hydrogels are the primary building blocks for bioprinting, and, also termed bioinks, are the foundations for creating structures that can potentially recapitulate the architecture of articular cartilage. To be functional, hydrogel constructs need to unlock the regenerative capacity of encapsulated cells. The recent identification of multipotent articular cartilage-resident chondroprogenitor cells (ACPCs), which share important traits with adult stem cells, represents a new opportunity for cartilage regeneration. However, little is known about the suitability of ACPCs for tissue engineering, especially in combination with biomaterials. This study aimed to investigate the potential of ACPCs in hydrogels for cartilage regeneration and biofabrication, and to evaluate their ability for zone-specific matrix production. Gelatin methacryloyl (gelMA)-based hydrogels were used to culture ACPCs, bone marrow mesenchymal stromal cells (MSCs) and chondrocytes, and as bioinks for printing. Our data shows ACPCs outperformed chondrocytes in terms of neo-cartilage production and unlike MSCs, ACPCs had the lowest gene expression levels of hypertrophy marker collagen type X, and the highest expression of PRG4, a key factor in joint lubrication. Co-cultures of the cell types in multi-compartment hydrogels allowed generating constructs with a layered distribution of collagens and glycosaminoglycans. By combining ACPC- and MSC-laden bioinks, a bioprinted model of articular cartilage was generated, consisting of defined superficial and deep regions, each with distinct cellular and extracellular matrix composition. Taken together, these results provide important information for the use of ACPC-laden hydrogels in regenerative medicine, and pave the way to the biofabrication of 3D constructs with multiple cell types for cartilage regeneration or *in vitro* tissue models.

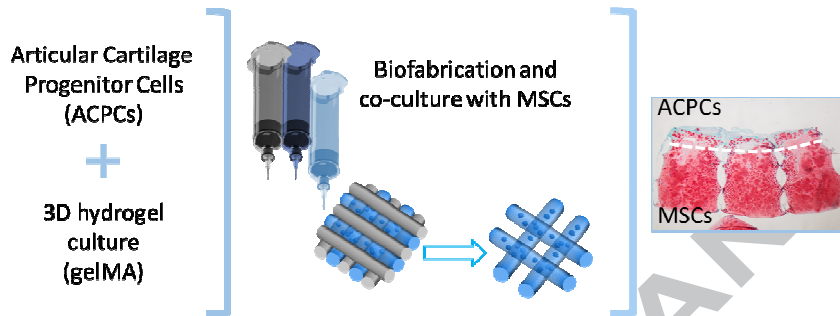
Keywords: Hydrogel, chondroprogenitor cells, biofabrication, cartilage regeneration, stem cells, co-culture

Statement of Significance

Despite its limited ability to repair, articular cartilage harbors an endogenous population of progenitor cells (ACPCs), that to date, received limited attention in biomaterials and tissue engineering applications. Harnessing the potential of these cells in 3D hydrogels can open new avenues for biomaterial-based regenerative therapies, especially with advanced biofabrication technologies (*e.g.* bioprinting). This study highlights the potential of ACPCs to generate neo-cartilage in a gelatin-based hydrogel and bioink. The

ACPC-laden hydrogel is a suitable substrate for chondrogenesis and data shows it has a bias in directing cells towards a superficial zone phenotype. For the first time, ACPC-hydrogels are evaluated both as alternative for and in combination with chondrocytes and MSCs, using co-cultures and bioprinting for cartilage regeneration *in vitro*. This study provides important cues on ACPCs, indicating they represent a promising cell source for the next generation of cartilage constructs with increased biomimicry.

Graphical Abstract



1. Introduction

Articular cartilage defects do not heal spontaneously and are prone to progress towards osteoarthritis, eventually resulting in impaired joint function, disability and a reduced quality of life [1]. Therapies involving the delivery of cells, including autologous chondrocyte implantation have substantially improved the outcome of treatments of such defects[2], but unfortunately often result in a low-performance repair tissue, which only delays the onset of degeneration and osteoarthritis[3].

In the quest for therapies that enhance cartilage healing, hydrogel-based constructs are particularly appealing for regenerative medicine, as they allow encapsulation of cells in a highly hydrated environment, analogous to that of native cartilage[4]. Moreover, cell-laden hydrogels can be used as bioinks, which are the building blocks of many biofabrication strategies[5]. Biofabrication allow the coordinated deposition of multiple cells and materials in a layer-by-layer fashion, enabling three-dimensional (3D) bioprinting of patient-specific and anatomically-shaped grafts[6]. Bioprinting also allows for the possibility of mimicking tissue-specific architecture such as the zonal and depth-dependent structure of articular cartilage[7] and also recreating the transition and interface between contiguous tissues[8].

While biofabrication holds the promise of introducing a new generation of regenerative therapies for cartilage[9], simply recapitulating the native structural and cellular composition during the printing process is only the first step of many. Regeneration is driven by the encapsulated cells (and, upon implantation, by the interplay with the host biomechanical and biochemical environment) and occurs over time, after the initial bioprinting step. Thus, in the design of hydrogels, and consequently bioinks, harnessing the potential of regeneration-competent cells is paramount. Nevertheless, the optimal cell source to be incorporated for cartilage-based cell therapies and tissue engineering is still subject of ongoing debate[10].

Chondrocytes are known to recover their ability to deposit cartilage-like matrix when cultured in a 3D hydrogel environment [11], but steadily lose their re-differentiation capacity after a few population doublings limiting their usefulness for cell therapy and tissue engineering of larger defects. To bypass this limitation, multipotent progenitor cells, such as mesenchymal stromal cells (MSC) derived from the synovium[12], umbilical cord[13], bone marrow[14] or adipose tissue[15] have also been applied to synthesize new cartilage in hydrogel matrices. However, while MSCs have been used with beneficial effects to the level of

clinical trial[16], their tendency to undergo hypertrophic differentiation and trigger endochondral ossification remains a major concern[17].

Recently, the identification and characterization of a population of a resident, cartilage-specific, multipotent progenitor cells has opened new avenues for cartilage repair[18,19]. Articular cartilage-derived progenitor cells (ACPCs) are found both in young and adult cartilage and make up about 0.1-1% of cartilage cell content. They are mainly located in the superficial zone of articular cartilage and express high levels of integrin alpha5, and thus can be enriched by differential adhesion to fibronectin, as no unique marker has been identified yet[18,20]. To date, ACPCs have been isolated from different species, including of human, equine and bovine.

ACPCs are sometimes referred to as cartilage progenitor/stem cells[21] or simply chondroprogenitors[22], although the latter term can create confusion, since it is also often used to describe any progenitor cell which has been driven towards chondrogenic differentiation, including MSCs, synovial fluid or synovial membrane-derived cells and ephyphial progenitors[21,22]. Much like MSCs, ACPCs are capable of *in vitro* self-renewal, and can be expanded to more than 60 population doublings while maintaining their potential to differentiate towards osteo-, chondro- and adipogenic lineages[18]. Interestingly, ACPCs play an important role in cartilage development[25], maturation[26], repair upon injury and response to osteoarthritic changes in the joint [21,27]. Moreover, unlike MSCs, ACPCs appear to show low or no expression of RUNX2, the master transcription factor for chondrocyte terminal differentiation and subsequent formation of calcified tissue[28]. This resistance to hypertrophy and the consistent production of hyaline-like cartilage from ACPC is maintained also during dynamic culture under multiaxial mechanical loading[29]. Additionally, ACPCs have been proposed as cell sources for autologous transplantation in cartilage in equine models[30], and even in a pilot clinical trial in humans with encouraging results[31]. Despite promising characteristics for cell and tissue therapies, the potential of ACPCs for tissue engineering is unexplored, in particular the behavior of ACPCs in 3D culture and biomaterial-driven tissue regeneration remains to be studied. Indeed, ACPCs have received limited attention in the biomaterials community as cell source for producing cartilage constructs, and only a few studies have highlighted their utilization in combination with scaffolds, even though adult, tissue-specific progenitor cells have been important resources for the regeneration of other tissues[32,33]. Harnessing the chondrogenic potential of ACPCs by combining them with a biomaterial matrix permissive

for cartilage production, and demonstrating their use as a bio-ink for advanced biofabrication strategies, can help advance the design of more functional implants for cartilage repair and regeneration.

Therefore, the aim of this study was to evaluate the potential of ACPC-laden hydrogel constructs for cartilage regeneration and to compare their extracellular matrix (ECM) synthesis capacity with native articular chondrocytes and bone marrow-derived MSCs. A photosensitive, gelatin methacryloyl (gelMA) hydrogel bioink was used as a platform for cell encapsulation and 3D culture, and the overall production of cartilage ECM by all three cell types was assessed, together with the mechanical properties of cultured constructs. Particular attention was paid to the expression of zonal markers, in order to evaluate if different progenitor cells display different preferential zonal affinities when cultured in a 3D gelMA-based milieu. Furthermore, layered co-cultures were performed as models for multi-layered cartilage constructs. Cross-talk between the different cell types and its effect on cartilage deposition were evaluated. Finally, proof-of-concept zonal cartilage constructs were biofabricated using a 3D bioprinting set-up.

2. Materials and Methods

2.1 Synthesis of gelMA

GelMA was synthesized from gelatin type A, obtained from porcine skin (Sigma-Aldrich, The Netherlands), as described previously[34]. Briefly, a 10% w/v solution of gelatin in PBS was reacted with 1:0.6 methacrylic anhydride (Sigma-Aldrich, The Netherlands) at 50°C for 1 hour, to achieve an 80% degree of modification of the lysine residues. Excess of methacrylic anhydride was removed by centrifugation. The resulting gelMA solution was neutralized with 1 M NaOH and dialyzed against distilled water. Eventually, the gelMA solution was sterile-filtered, freeze-dried and stored at -20°C until used.

2.2 Cell isolation

All animal tissue and cells used in this study were obtained from deceased equine donors, donated to science by their owner, and according to the guidelines of the Institutional Animal Ethical Committee. These were all skeletally mature horses, aged 3 to 7 years old, not suffering from any disorder of the joints from which cartilage was harvested or from any disorder possibly affecting MSCs in case of harvesting bone marrow.

Equine cells were chosen due to the morphological and biochemical similarity between human and equine cartilage tissue[35], and in the perspective of successive preclinical testing, as the equine model is important for testing cartilage repair strategies[36,37]. At the same time, horses are also patients when it comes to joint pathologies.

2.2.1 Chondrocytes and ACPCs

Macroscopically healthy cartilage from the metacarpophalangeal joint of two equine donors was harvested with a scalpel under sterile conditions, without damaging the tidemark. The tissue was minced and digested in 0.2% pronase for 2 hours, followed by incubation for 12 hours in a 0.075% collagenase type 2 solution. The tissue digest was sieved through a 70 μm cell strainer and the resulting single-cell suspension was centrifuged for 5 minutes at 300g. Pelleted chondrocytes (CH) were washed in phosphate buffered saline (PBS) and counted with a hemocytometer. An aliquot of the total cell harvest was used to isolate ACPCs, the remaining chondrocytes were stored in liquid nitrogen until further use. Prior to encapsulation in gelMA, chondrocytes were expanded to passage 1 in Dulbecco's modified Eagle medium (DMEM, 31966, Gibco, The Netherlands), supplemented with 10% v/v heat-inactivated fetal calf serum (FCS, Gibco, The Netherlands), 0.2 mM ascorbic acid-2-phosphate, 100 U/mL penicillin and 100 $\mu\text{g}/\text{mL}$ streptomycin.

ACPCs were isolated as previously described[18]. Briefly, an aliquot of the freshly isolated cartilage cells was pelleted by centrifugation, suspended in serum-free DMEM, and plated in fibronectin-coated tissue culture plates, at a density of 500 cells cm^{-2} . After 20 minutes, non-adherent cells were removed, and the attached cells were cultured in chondroprogenitor expansion medium (DMEM, supplemented with 10% v/v FCS, 0.2 mM ascorbic acid-2-phosphate, 100 U/mL penicillin, 100 $\mu\text{g}/\text{mL}$ streptomycin and 5 ng/mL basic fibroblast growth factor (bFGF, Peprotech, UK)). After 6 days, colonies with more than 32 cells were marked for cloning. Collected colonies were pooled and cells were expanded until passage 3 before being used for the study.

2.2.2 MSCs

Bone marrow aspirates were obtained from the sternum of two equine donors, as previously described [17]. Briefly, the mononuclear cell fraction was derived from the bone marrow aspirate in a Ficoll-paque density gradient (GE Healthcare, The Netherlands), after centrifugation for 30 minutes at 100g. The mononuclear

cell fraction was collected, washed with PBS and centrifuged again at 300g for 10 minutes. Finally, the cells were plated on tissue culture plastic and cultured in MSC expansion medium, consisting of α MEM (22561 Gibco, The Netherlands) supplemented with 0.2mM L-ascorbic acid 2-phosphate (Sigma), 10% FCS (Lonza, The Netherlands), 100 U/mL penicillin with 100 μ g/mL streptomycin (Life Technologies, The Netherlands) and 1 ng/mL bFGF. Cells grown to passage 3 were used for this study.

2.3 Multipotency and characterization of progenitor cells

For osteogenic and adipogenic differentiation, ACPCs and MSCs were expanded plated in 6 well plates at a density of 2×10^5 cells/well, cultured until subconfluent, and cultured for 21 days either in osteogenic medium (alpha-MEM supplemented with 10% FCS, 100 U/ml penicillin, 100 μ g/mL streptomycin, 0.2 mM L-ascorbic acid-2-phosphate, 20 mM β -glycerol phosphate, 100 nM dexamethasone, Sigma-Aldrich, The Netherlands), or adipogenic medium (α MEM supplemented with 10% v/v FCS, 100 U/mL penicillin, 100 μ g/mL streptomycin, 0.01 mM indomethacin, 83 mM isobutylmethylxanthine, and 1.72 μ M bovine pancreas-derived insulin, Sigma-Aldrich, The Netherlands). Medium was refreshed every 3 days. At the end of the culture, samples were stained for calcified matrix (alizarin red) and for intracellular lipid vesicles (oil red O). For chondrogenic differentiation, 2.5×10^5 cells were pelleted by centrifugation in 15 mL Falcon tubes and cultured in chondrogenic medium (DMEM supplemented with 1% v/v insulin-transferrin-selenous acid (ITS+ Premix, Corning, USA), 0.2 mM ascorbic acid-2-phosphate, 100 units/mL penicillin with 100 μ g/mL streptomycin (Life Technologies, The Netherlands), 100 nM dexamethasone and 10 ng/ml transforming growth factor- β 1 (TGF- β 1)). Medium was refreshed every 3 days. After 21 days, histological sections of the pellets were stained for sulphated glycosaminoglycans (GAG) with safranin-O.

Reverse transcription-polymerase chain reaction (RT-PCR), was used to characterize the gene expression of the cell surface markers CD13, CD29, CD31, CD44, CD45, CD49d, CD73, CD90, CD105, CD106, CD146 and CD166 in ACPCs and MSCs[38]. Hypoxanthine phosphoribosyltransferase-1 (HPRT1) was monitored as a housekeeping gene. The primer sequences are reported in the supplementary information (Table ST1). RNA isolation was performed on cells at passage 3 using the RNeasy mini kit (Qiagen, Germany), following the instructions of the manufacturer. Isolated RNA was quantified by UV-vis spectrophotometry

with a Nanodrop 2000 (Thermo Scientific, The Netherlands), and used as template for the PCR reaction. Amplification was carried out using a SuperScript® One-Step RT-PCR System with Platinum® Taq DNA Polymerase (Life Technologies, The Netherlands). The PCR products were run in agarose gel electrophoresis and stained with ethidium bromide.

2.4 Cell encapsulation in gelMA, and 3D monoculture and layered co-culture

10% w/v gelMA was dissolved in PBS, supplemented with 0.1% w/v 2-hydroxy-1-[4-(2-hydroxyethoxy)phenyl]-2-methyl-1-propanone (Irgacure 2959, BASF, Germany) as a photoinitiator. The temperature of this solution was stabilized at 37°C, prior to cell mixing. For the monoculture samples, $1.5 \cdot 10^7$ cells/mL (ACPC, MSC or CH) were homogeneously suspended in the macromer solution, and cast into a custom-made Teflon mold to produce cylindrical samples (height = 2 mm, diameter = 6 mm). The mixture was UV-irradiated for 5 minutes ($\lambda = 365$ nm, $E = 3$ mW cm⁻² at $h = 2$ cm; Vilber-Lourmat 144 portable UV-lamp) to trigger the free-radical polymerization and thus the chemical crosslinking of the hydrogel. Cell-free hydrogels, prepared in the same way, were used as control.

Layered co-cultures, consisting of two adjacent hydrogels cast on top of each other, were also prepared as models for zonal constructs. In each layer only one single cell type was encapsulated, and all the possible cell combinations (ACPC/MSC, ACPC/CH and MSC/CH, all ratios 1:1) were studied. To fabricate these constructs, a gelMA macromer solution laden with the first cell type (density = $1.5 \cdot 10^7$ cells/mL) was cast into custom-made cylindrical molds (height = 2 mm, diameter = 6 mm). The mixture was partially crosslinked via UV-irradiation for 2 minutes, to leave unreacted methacrylate groups available for chemical binding with the second gelMA layer. Subsequently, a second mold was aligned on top of the first one, and a gelMA solution laden with the second cell type was cast. The double-layered constructs were UV-irradiated for 5 additional minutes to complete the crosslinking. No delamination was observed when handling the layered gels. An optical microscopy picture and a schematic representation of the layered 3D co-culture system are reported in the supplementary information (Figure S1). All the samples (mono-, co-cultures and cell-free hydrogels) were cultured in chondrogenic differentiation medium as described in section 2.3.

2.5 Biochemical analysis

Samples (both monocultures and layered co-cultures) were harvested to measure DNA and sGAG content after 1, 28 and 56 days of culture (n = 4-6). The constructs were freeze-dried and digested overnight in 0.01 M cysteine, 250 µg/mL papain, 0.2 M NaH₂ PO₄ and 0.01 M EDTA at 60°C. Subsequently, the digest sample was reacted with dimethylmethylene blue (DMMB, Sigma-Aldrich, The Netherlands), and the sGAG content calculated by determining the ratio of the absorbance at 525 and 595 nm with a spectrophotometer (Bio-Rad, Hercules, CA). Known dilutions of chondroitin sulfate were used as standards for quantitative analysis. DNA content of the constructs was quantified on papain digests, using a Quant-iT PicoGreen dsDNA kit (Life Technologies, The Netherlands).

2.6 Mechanical testing

The mechanical properties of the monoculture constructs were studied in an unconfined uniaxial compression test, with a Dynamic Mechanical Analyzer (DMA Q800, TA Instruments, The Netherlands). Samples at days 1, 28 and 56 of culture were washed with PBS and compressed at a -20%/minute strain rate (n = 4-6). The compression modulus was calculated as the slope of the stress/strain curve in the 10% to 15% strain range.

2.7 Gene expression of cartilage and zonal markers

Gene expression analysis was performed by qPCR on monoculture samples taken at days 1, 28 and 56 of culture (n = 3). Constructs were harvested and mechanically ground in RLT buffer (Qiagen, Germany). The lysate was then processed with the RNeasy Mini kit to isolate mRNA. Amplification and cDNA synthesis were performed with a SuperScript III Platinum SYBR Green One-Step qRT-PCR Kit (LifeTechnologies, The Netherlands). The relative expression levels for aggrecan (ACAN), collagen type II (COL2A1), collagen type I (COL1A1), proteoglycan 4 (PRG4), collagen type X (COL10A1) and cartilage oligomeric matrix protein (COMP) were analyzed compared to the housekeeping gene HPRT1. The primers sequences are reported in the supplementary information (table ST2). Relative expression, Ct and efficiency values were calculated using the PCRminer algorithm[39].

2.8 Histological assessment

The deposition of main constituents of the cartilage ECM in the hydrogel was assessed on formalin-fixed, paraffin-embedded samples (n =3). Prior to the harvesting, cultured constructs were treated overnight with 0.1 μ M monensin to trap PRG4 intracellularly. Tissue sections (5 μ m) were sliced with a microtome and processed for the staining. sGAG content was visualized by Safranin O, collagen with fast green staining, and cell nuclei with haematoxylin. Immunohistochemistry was performed using the appropriate primary antibodies for collagen type I (sc-8784, Santa Cruz Biotechnology, USA), type II (DSHB, II-II6B3, USA) and PRG4 (ab28484, Abcam, The Netherlands). Samples were deparaffinized with xylene, hydrated in ethanol graded solutions, and treated with 0.3% v/v H₂O₂ to block endogenous peroxidases. For collagen type I and II, antigen retrieval was performed with pronase (1 mg/mL, Roche, USA) and hyaluronidase (10 mg/mL, H2126, Sigma Aldrich, The Netherlands), applied for 30 min at 37 °C. Tissue sections were blocked with bovine serum albumin (BSA, 5% w/v in PBS) for 1 hour at room temperature and the primary antibody was incubated overnight at 4°C. Depending on the primary antibody, appropriate IgGs were used as isotype controls. Sections were then incubated with a secondary antibody for 1 hour at room temperature, and the staining was developed with 3,3-diaminobenzidine-horseradish peroxidase (Sigma Aldrich, The Netherlands). Cell nuclei were counterstained with hematoxylin. Sections were mounted in DPX (Millipore, USA), and micrographs were taken with an optical microscope (Olympus BX51, Olympus, Germany).

2.9 Bioprinting of zonal-like constructs

Zonal-like constructs (12 x 12 x 2.16 mm) were fabricated using a 3DDiscovery bioprinter (regenHu, Switzerland). The printer path was drawn in vector graphic and translated into g-code with the BioCAD software (regenHu, Switzerland). Three materials were loaded for printing: i) a superficial zone-mimicking bioink, consisting of 10% w/v gelMA laden with $2 \cdot 10^7$ ACPC/mL, ii) a middle/deep zone-mimicking bioink, composed of 10% w/v gelMA laden with $2 \cdot 10^7$ MSC/mL, and iii) pluronic F-127 (40% w/v in PBS) as sacrificial ink to support gelMA during the biofabrication process. Each layer was obtained printing a frame of parallel pluronic filaments (room temperature, pressure = 0.180 MPa, 23G nozzle, translation speed = 20 mm/s) with a strand-to-strand distance of 1.2 mm. Subsequently, gelMA was dispensed in between the sacrificial strands using a microvalve (pressure = 0.04 MPa, 23G nozzle, dosing distance = 0.3 mm, opening time = 800 μ s, temperature = 37°C, translation speed = 30 mm/s). Printed layers were pre-crosslinked for 10

seconds via exposure to a built-in UV led ($\lambda = 365$ nm, $E = 240.2$ mW cm^{-2} at $h = 1$ cm). Layer height was set to 0.24 mm, and 3D constructs with a woodlog 0° - 90° structure were generated in a layer by-layer fashion. The first seven layers were printed with the MSC-laden bioink; the last two with the ACPC-laden bioink. After printing, crosslinking of gelMA was completed by exposure for 5 minutes to a Vilber-Lourmat 144 portable UV-lamp ($\lambda = 365$ nm, $E = 3$ mW cm^{-2} at $h = 2$ cm). Samples were soaked in cold PBS to remove the pluronic, cut into 3.5 x 3.5 mm squares and cultured for 56 days in chondrogenic differentiation media ($n = 3$). Cell viability was evaluated at day 1 and 14 after printing using calcein AM and ethidium homodimer-1 (Life Technologies) ($n = 3$), and compared to that of cast scaffolds. ECM deposition was qualitatively observed with histological analysis, as described in section 2.8. Morphology of the printed constructs was also observed from microscopy images and x-ray micro computerized tomography scans (μCT , Quantum FX, Perkin Elmer, USA, spatial resolution of 20 μm^3 of voxel size, scan time = 3 min, tube voltage = 90 kV and tube current = 180 μA). Additionally, a proof-of-concept fabrication of anatomically shaped constructs was performed. A computer-aided design (CAD) model of the caudal end of a human femur (maximum dimensions 26.5 x 26.3 x 23.1 mm) was processed with a CAD software (Rhinoceros, Robert McNeel and Associates, USA) to separate the distal end of the femoral condyle from the part modelling the proximal side and the underlying bone, thus generating two complementary models. The proximal part was converted to an STL file and the model was eventually printed with a digital light projection 3D printer (Ember, Autodesk, USA) using a proprietary PR48 resin (Autodesk). The height of each printed layer was set to 100 μm . Next, the distal end of the condyle was bioprinted on top of this model. A computer aided manufacturing software (CAM, BioCAM, regenHu) was used for slicing the condyle model and the g-code was generated with the BioCAD software (regenHu). The DLP-printed anatomical model was interfaced with the 3DDiscovery, and the caudal end of the condyle was printed using the approach and printing settings described above, via the co-printing of a sacrificial pluronic support and gelMA as a bioink.

2.10 Statistical analysis

Each experiment was performed in three to six replicates ($n = 3$ to 6). Quantitative results are expressed as mean \pm standard deviation (SD), and the statistical analyses were performed using the GraphPad Prism 6.0

software package (GraphPad Software, USA). Comparisons between the experimental groups at different time points were performed with a two-way ANOVA, with a Bonferroni post-hoc test. F-values and related degrees of freedom are listed in the supplementary information (Table ST3). A value of $p < 0.05$ was considered statistically significant.

3. Results

3.1 Characterization of ACPCs

Gene expression analysis showed high similarity between MSCs and ACPCs in their transcript profile of surface receptors important in defining the minimal criteria for mesenchyme stromal cell classification (Figure 1A). Notably, much like MSCs, ACPCs were positive for the stem cell markers CD73, CD90 and CD105, while being negative for the hematopoietic marker CD34 and for the leukocyte marker CD45 [40]. Additionally, both MSCs and ACPCs were CD29⁺, CD44⁺, CD49d⁺, CD106⁺, CD166⁺, and faintly positive for CD146, CD13⁻ and CD31⁻. ACPCs appeared to display higher expression of CD44, a cell membrane receptor for hyaluronic acid. Moreover, both bone-marrow derived MSCs and ACPCs were capable of differentiation towards osteogenic, chondrogenic and adipogenic lineage (Figure 1B).

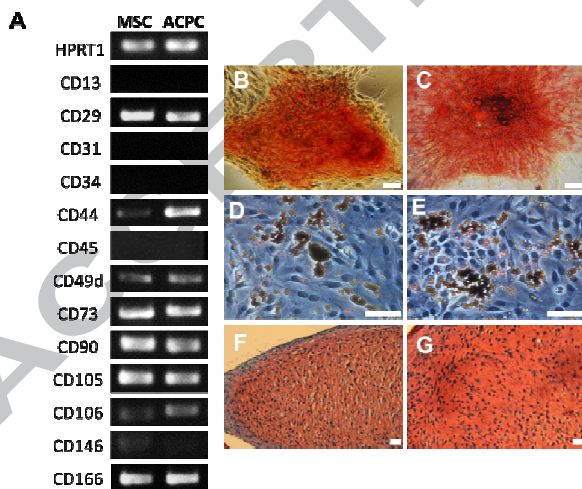


Figure 1: (A) Comparison of ACPC and MSC gene expression for several surface markers, as obtained from RT-PCR. Tri-lineage differentiation of (B,D,F) MSCs and (C,E,G) ACPCs, showing osteogenic (B,C, alizarin red staining), adipogenic (D,E, oil red O staining) and chondrogenic differentiation (F,G, Safranin O staining). Scale bars represent 50 μm.

3.2 3D culture and chondrogenic differentiation in mono-cultures

3.2.1 Biochemical evaluation of neo-cartilage

When cultured in 3D gelMA hydrogels, all three cell types, chondrocytes, ACPCs and MSCs were able to proliferate in the gel matrix. Significant differences in cell amount between experimental groups could be observed only at day 56, with MSC containing matrices containing less cells compared to ACPC-containing matrices, as estimated by the total DNA content of the hydrogel (Figure 2A). As for the deposition of neo-cartilage ECM, all three cell types were able to undergo chondrogenic differentiation, and synthesized sGAGs which were retained within the hydrogel matrix. ACPCs, however outperformed articular chondrocytes both in terms of total sGAG (Figure 2B) and sGAG normalized to DNA content (Figure 2C), displaying values more than 2.3-fold higher than chondrocytes. On the other hand, MSCs produced significantly more sGAG than cartilage-derived cells (2.6-fold vs ACPCs and 6.3-fold vs chondrocytes).

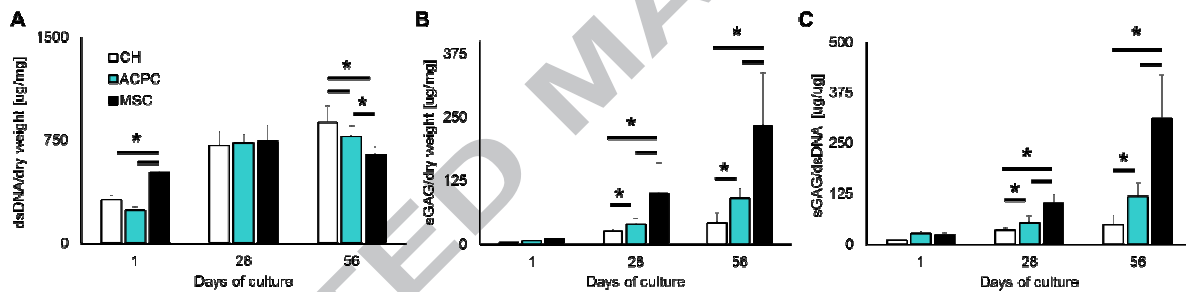


Figure 2: Quantification of (A) DNA and (B) sGAGs in the hydrogels for each cell type. Panel (C) shows sGAGs normalized per DNA content. ACPCs outperformed chondrocytes, while MSCs showed the highest sGAG/DNA ratio among the three cell types tested.

3.2.2 Neo-tissue deposition and mechanical properties

The compression modulus of the constructs was monitored over the culture time (Figure 3). Cell-free gelMA samples displayed no change in the compression modulus, which remained in the range of 18-23 kPa, suggesting no significant degradation of the hydrogel in culture medium over the two months of the assay. For all the cell-laden samples, cartilage matrix synthesis and accumulation in gelMA correlated with an increase of the mechanical properties of the hydrogel. After 4 weeks, chondrocytes, ACPC and MSC-laden constructs displayed a compressive modulus ranging between 42 to 47 kPa. At 8 weeks, a further increase in stiffness was observed and a significantly higher elastic modulus was found for MSC-laden construct (≈ 186.8 kPa vs 101.4 kPa and 125.5 kPa of ACPCs and chondrocytes).

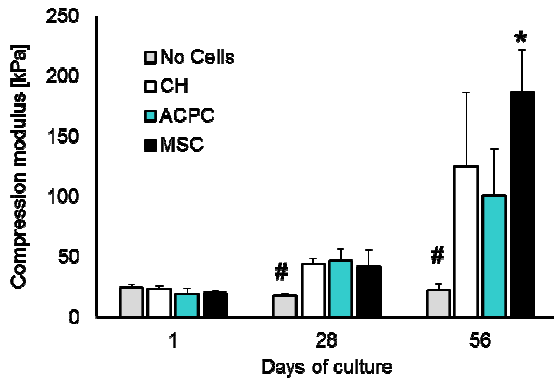


Figure 3: Compression modulus of cell-laden hydrogels along the culture period. * and # denote significant differences compared to all the other experimental groups ($p < 0.05$).

3.2.3 Differential mRNA expression of zonal markers in hydrogel culture

Gene expression of cartilage-specific transcripts increased over time, confirming chondrogenic differentiation of the cells in the hydrogels (Figure 4). In particular, aggrecan mRNA expression showed a correlation with sGAG synthesis, with higher values for MSCs (≈ 15 -fold at day 28 and day 56), followed by ACPCs (≈ 6.6 -fold at day 28 and 9.5-fold at day 56) and finally chondrocytes (4.1- and 2.1-fold at day 28 and 56 respectively; Figure 4A). A similar behavior was found for collagen type II, with MSCs displaying about a 145-fold (day 28) and 89.3-fold (day 56) values higher than ACPCs (19-fold and 30.7-fold) and chondrocytes (5- and 2.3-fold; Figure 4B). Collagen type I was also overexpressed in ACPCs and reached the highest values at day 28 (160.7- and 171.1-fold). In MSC-laden matrices, collagen type I showed a different trend: it was highly upregulated at the beginning of the culture (58.9-fold at day 1), increased to 81.2-fold at day 28 and decreased to a 48.1-fold overexpression at day 56 (Figure 4C), however this trend over the culture period was not significant. Moreover, COMP, the main non-collagenous protein in cartilage, displayed highly upregulated mRNA levels in all the experimental groups, compared to day 1 (Figure 4D). The analysis of the superficial zone marker PRG4 showed that the gene is strongly upregulated in ACPCs (21.2- and 99.5-fold at days 28 and 56), compared to chondrocytes (15.7- and 13.2-fold) and MSCs, which displayed the lowest levels (7.1- and 6.9-fold; Figure 4E). Conversely, collagen type X, which is a biomarker of endochondral ossification and a marker for calcified chondrocytes, was consistently downregulated in all experimental groups, but steadily increased over time in MSC-laden matrices (0.12- and 0.37-fold at day 1

and 56, respectively) and chondrocyte (0.28- to 0.48-fold), while decreasing in ACPC-laden matrices (0.35- to 0.18-fold; Figure 4F).

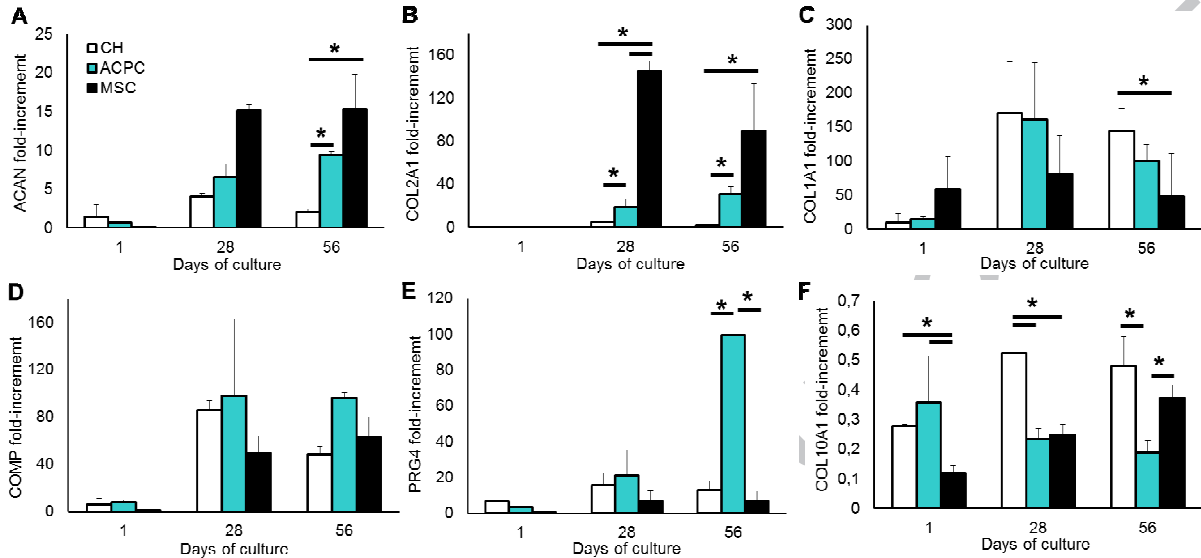


Figure 4: qPCR analysis of the cell-laden hydrogels, showing relative gene expression of (A) aggrecan, (B) collagen type II, (C) collagen type I, (D) COMP, (E) PRG4 and (F) collagen type X, normalized against the housekeeping gene HPRT1. Statistically significant differences are marked with an * ($p < 0.05$).

3.2.4 Histological evaluation of hydrogel mono-cultures

The presence and distribution of the main constituents of cartilage ECM in the hydrogel matrix were qualitatively observed on histological sections (Figure 5). A clear difference in the amount of sGAGs (Figure 5A) and collagen type II (Figure 5B) was detected between the three cell types, correlating with the quantitative analysis for sGAGs and collagen type II gene expression. Nevertheless, all experimental groups showed the formation of interconnected, homogeneously distributed staining for sGAGs at the last time point of the culture, while only ACPCs and MSCs displayed ubiquitous staining for collagen type II. Collagen type I was also in line with what was observed in the gene expression panel, with CHs having more intense and diffused staining at earlier time points than ACPCs, and MSCs in the last place (Figure 5C). In all experimental groups, most cells were positive for PRG4, with the least intense staining present for MSCs at 56 days of culture (Figure 5D). Interestingly, for all the samples, a layer of spread, elongated cells, with intense and continuous PRG4 positive staining was found at the outer rim of the hydrogel.

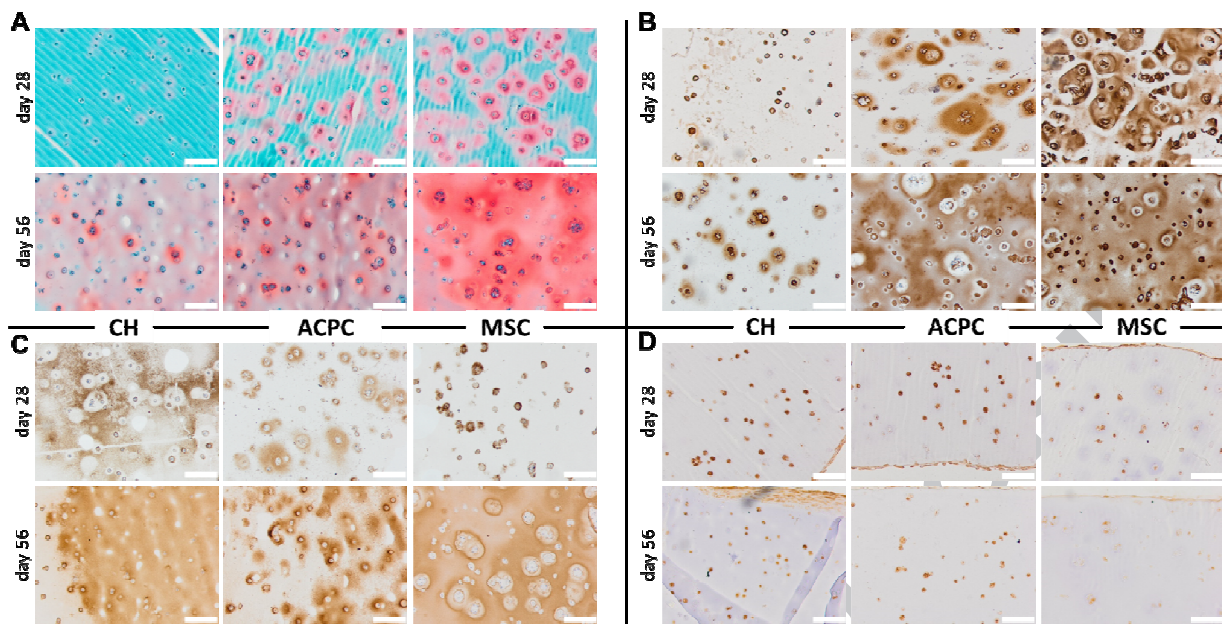


Figure 5: Histological analysis of the cultures at days 28 and 56. (A) Safranin O staining, and immunohistochemistry for (B) collagen type II, (C) collagen type I, and (D) PRG4. Scale bar is 200 μm .

3.3 3D co-culture in layered hydrogels

3.3.1 Biochemical analysis of layered co-cultures

In the co-culture system, all the possible cell combinations displayed higher amounts of DNA compared to their respective monoculture controls (Figure 6A). This effect appears to be more pronounced when MSCs are present in the construct. At days 28 and 56 MSC/chondrocyte matrices showed a 1.8- and 1.5-fold higher DNA production against monocultures, in comparison to a 1.4- and 1.5-fold increase for MSC/ACPC and a 1.3- and 1.2-fold for ACPC/chondrocyte samples, respectively. MSC/ACPC co-cultured matrices displayed the highest overall sGAG concentration ($\approx 53 \mu\text{g}/\text{mg}$ at day 28 and $136 \mu\text{g}/\text{mg}$ at day 56), followed by MSC/chondrocyte matrices (≈ 29 and $52 \mu\text{g}/\text{mg}$) and ACPC/chondrocyte matrices (≈ 25 and $55 \mu\text{g}/\text{mg}$; Figure 6B). It is worth highlighting that overall sGAG synthesis all co-culture groups, apart from MSC/chondrocyte samples, showed no significant difference with their respective mono-culture controls. As a consequence of having more cells but the same sGAG content, co-cultures tended to show a lower sGAG/DNA ratio

compared to their monoculture controls (Figure 6C). MSC/ACPC was the best performing co-culture group (≈ 51 and $150 \mu\text{g}_{\text{GAG}}/\mu\text{g}_{\text{DNA}}$ at days 28 and 56), performing better than MSC/chondrocyte (≈ 23 and $45 \mu\text{g}_{\text{sGAG}}/\mu\text{g}_{\text{DNA}}$ at days 28 and 56) and ACPC/chondrocyte (≈ 28 and $60 \mu\text{g}_{\text{sGAG}}/\mu\text{g}_{\text{DNA}}$ at days 28 and 56).

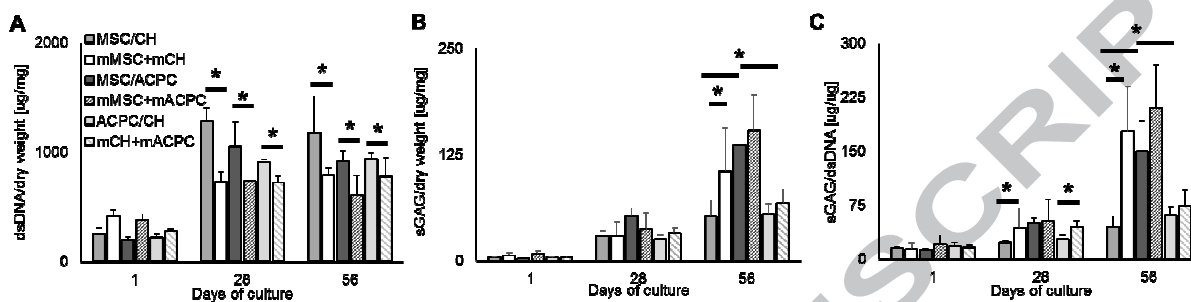


Figure 6: Biochemical analysis of the layered co-cultures, showing quantification of (A) DNA, (B) sGAGs, and (C) sGAGs normalized to DNA content. Statistically significant difference among the samples are indicated by an * ($p < 0.05$)

3.3.2 Histological evaluation of layered hydrogels

Qualitative evaluation of sGAGs (Figure 7A) and collagen type II (Figure 7B) confirmed the trends observed in monocultures, even when cells are co-cultured in adjacent hydrogel layers, with MSCs produced more matrix, followed by ACPCs and finally chondrocytes. These differences resulted in the formation of constructs with a distinct composition in each layer. Collagen type I was homogeneously distributed throughout the hydrogel matrix (Figure 7C), though antibody labelling was less intense compared to collagen type II labelling. PRG4 immunohistochemistry was positive intracellularly for all cells, and dense clusters of positively stained cells were found at the border of the construct, regardless of the zone of the hydrogel (Figure 7D).

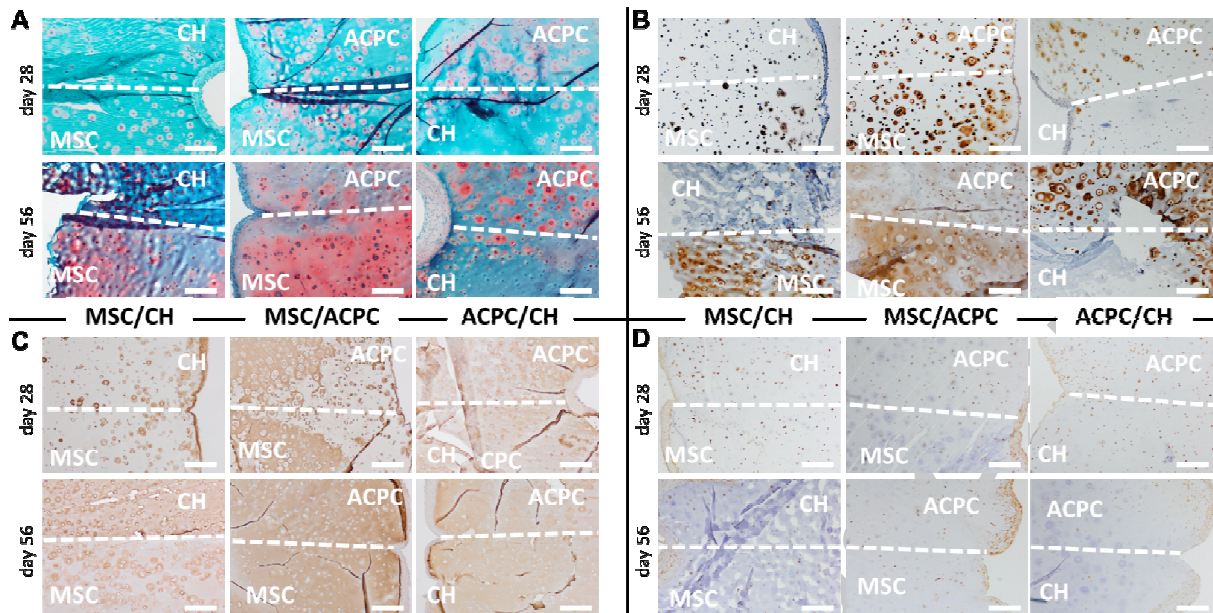


Figure 7: Histological sections of the layered co-cultures showing (A) safranin O staining for sGAGs, (B) collagen type II, (C) collagen type I, and (D) PRG4. The dotted line marks the interface between the two different cell-laden layers of the hydrogels, and the cell type residing in each zone is specified in overlay. Scale bar is 500 μm .

3.3.3 Bioprinted zonal-like constructs

Shape stable constructs, with zonal-like cell distribution were fabricated via bioprinting of cell-laden gelMA (Figure 8). After photocrosslinking and removal of the sacrificial pluronic supporting frame, the materials retained their shape, no delamination was observed, and interconnected pores were present after removal of the pluronic strands (Figure S2, Supplementary Video SV1). Cell viability was comparable between printed and cast constructs, ranging from about 75% to 90% at days 1 and 14 after fabrication (Supplementary Figure S3). Upon long-term culture, both ACPCs and MSCs in the printed layered constructs differentiated and produced cartilage matrix, with sGAG/DNA values comparable to those found in the mono- and co-cultures ($168.31 \pm 51.28 \mu\text{g}_{\text{sGAG}}/\mu\text{g}_{\text{DNA}}$, after 56 days of culture). Additionally, histological analysis confirmed a zonal difference in the distribution of sGAGs from the ACPC-laden to the MSC-laden zone. Collagen type II staining appeared more intense and homogeneous throughout the gel, compared to collagen type I, which was more intense at the border of the construct. Furthermore, anatomically-shaped models of the caudal end of a human femur condyle were successfully obtained with the proposed bioprinting approach (Figure 9).

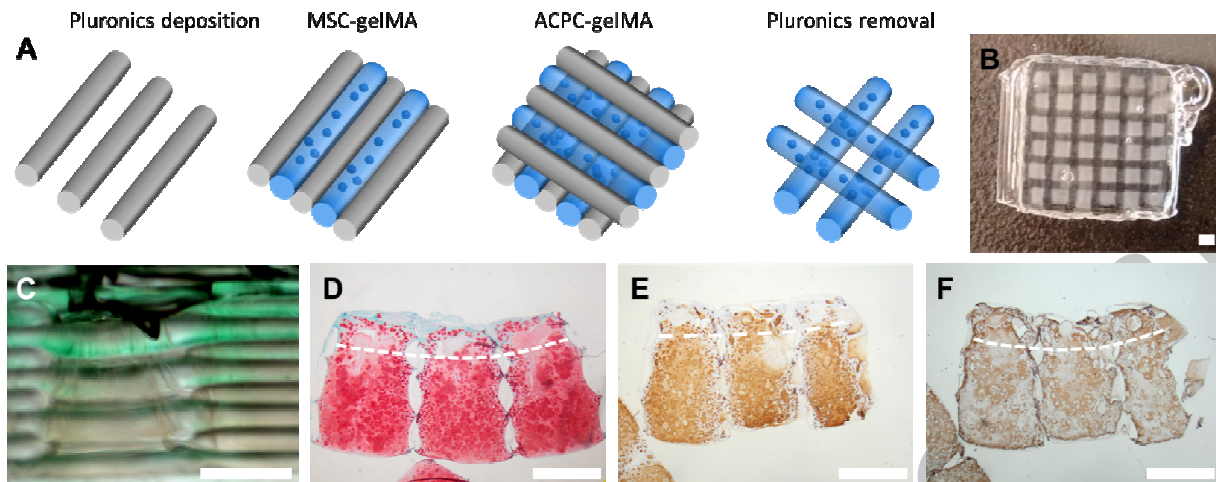


Figure 8: Bioprinted cartilage constructs with MSCs in the middle/deep layer and ACPCs in the superficial layer. (A) Scheme of the printing process, (B) view of the construct after printing with the pluronic frame, (C) lateral view of the construct, showing in green the superficial layer bioink. Histological staining after 56 days of culture for (D) sGAGs, (E) collagen type II, (F) collagen type I. The dotted line indicates the border between the ACPC-laden (top) and MSC-laden (bottom) zone. Scale bar represents 1 mm.

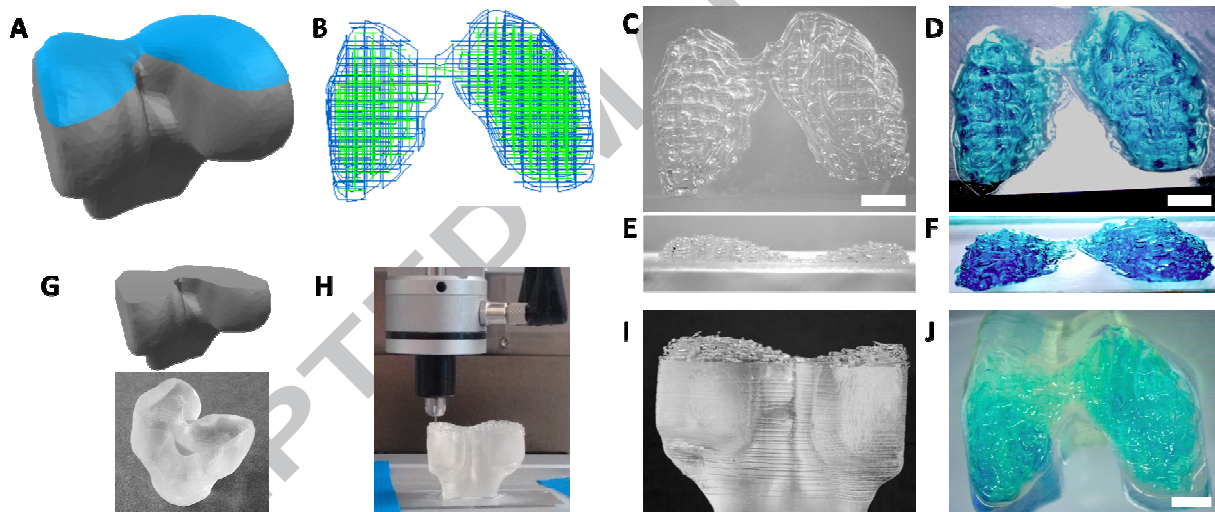


Figure 9: Proof-of-concept of bioprinting anatomical structures. (A) A CAD model of a femur condyle is obtained (highlighted in blue) and (B) used to generate the G-code and the path of the dual printing system (showing in blue the path for the supporting material and in green that of the bioink). (C, E) The top part of the femur condyle is printed together with the supporting hydrogel, (D, F) that can be then removed leaving only the bioink (stained in blue). (G) A model of the lower part of the joint and the underlying bone was produced using a DLP 3D printer and (H) the condyle structure was printed directly on top of it, as a proof-of-concept test to replace the missing anatomical part, via co-extrusion of the supporting sacrificial poloxamer and gelMA bioink. (I) This allows accurate printing of the shape both in presence of the supporting material and (J) after its removal. Scale bar is 5 mm.

4. Discussion

The results of this study demonstrate the ability of adult ACPCs to synthesize neo-cartilage in a hydrogel system. ACPCs appear to have potential to replace or complement chondrocytes and MSCs that are widely

established cell sources for cartilage cell-based therapies. In this study, layered cast and bioprinted constructs composed of two regions laden with distinct cell types were also produced, generating a zonal-like distribution of the main cartilage components.

GelMA was used as a hydrogel for cell encapsulation and printing. Overall, gelMA is becoming a widespread platform for 3D culture, organ models and tissue engineering[41], and it is also one of the most versatile bioinks for biofabrication[42–46]. Previous research already demonstrated that gelMA provides a permissive environment for neo-cartilage formation, using encapsulated chondrocytes[47,48], and MSCs [49]. In this study, all three tested cell types underwent chondrogenic differentiation and synthesized a cartilage-like matrix, which over time induced a considerable increase in the compressive properties of the hydrogel. A direct comparison can be drawn between chondrocytes and ACPCs, since these cells are obtained not only from the same donors, but also from the same tissue. More specifically, ACPCs are a subpopulation of the chondrocytes from full-thickness cartilage. Previous studies have shown that in contrast to full-depth dedifferentiated populations of articular chondrocytes, ACPCs from the same tissue retain SOX9 gene and protein expression, and therefore differentiation capacity even after extensive culture expansion [22]. In gelMA matrices, ACPCs outperformed chondrocytes in terms of cartilage matrix production (amount and distribution of sGAGs and collagen type II, less presence of collagen type I), a difference also reflected at the level of gene expression. These results, combined with the ability of ACPCs to undergo multiple passages without losing their chondrogenic potential[18], suggest that ACPCs could be a viable alternative to chondrocytes for improving cell-based cartilage therapies (such as matrix-induced autologous chondrocyte implantation), and further investigation in this direction is warranted.

ACPCs behave similarly to MSCs in terms of *in vitro* multipotency and self-renewal. In this study, we confirmed that ACPCs display gene expression of cell surface markers that is comparable with bone marrow-derived MSCs, which is in line with previously reported data[38]. Particularly, ACPCs, being CD34⁻, CD45⁻, CD73⁺, CD90⁺, C105⁺, feature the same minimal marker profile for the definition of MSCs in human tissue[40]. Despite this phenotypic similarity, marked differences were observed when chondrogenic differentiation was induced in 3D cultures. Overall, MSCs produced more sGAGs in gelMA, and displayed the highest collagen type II/type I ratio at mRNA level. Two inferences can be drawn from these data, first that the chondrogenic medium optimized for bone derived MSC differentiation may not be optimal for

ACPCs, secondly that there is a lack of tissue and cell signaling found within the native stem cell niche that may be required to enable ACPCs to differentiate into more mature mid and deep zone chondrocytes[50]. Clear differences between ACPCs and MSCs were observed in terms of zonal markers. Chondrocyte hypertrophy and associated calcified cartilage production is a common concern associated with MSC use for articular cartilage repair. Though collagen type X mRNA was only weakly expressed by MSCs, possibly thanks to the continuous supply of TGF- β 1 from the media, it steadily increased over time. While MSCs remain a versatile cell source for cartilage repair, ACPCs may be a better alternative to prevent inappropriate differentiation. This would be especially valuable when the chosen clinical strategy to treat cartilage defects has high risk of triggering MSC hypertrophy, for instance when osteochondral grafts which provide strong osteogenic signals in the bone component are implanted[51]. Over the culture period, ACPCs showed the highest transcript expression of PRG4, a key factor in joint lubrication and a well-known superficial zone marker, and the lowest expression of collagen type X, the hallmark of calcified cartilage. This suggests that ACPCs may be primed towards a superficial zone phenotype, perhaps as a result of pre-conditioning due to their original niche in native cartilage. Immunohistochemistry for PRG4 showed that all three cell types were positive for this proteoglycan, especially those at the external boundary of the hydrogels. However, since the antibody stains only intracellular targets, no information was obtained on the PRG4 secreted and incorporated in the hydrogel matrix. Several researchers have attempted to enhance PRG4 expression by cultured MSCs and chondrocytes in hydrogels via mechanical stimulation [52], through the biofunctionalization of the hydrogel matrix[53], use of co-cultures[54], varying cell density and growth factors[55,56], or by directly encapsulating zonally harvested chondrocytes [57]. Instead, the utilization of ACPCs may be a much simpler and still effective alternative for the engineering of salient features of the superficial zone.

Next, taking into consideration the differential expression of cartilage component by cells, we generated multiple combinations of bi-layered cartilage constructs. While no report is available on co-cultures involving ACPCs, MSC and chondrocytes, co-cultures have been widely described. Cell-cell contact has been identified as a key factor in enhancing GAG synthesis in MSC/chondrocytes systems[16]. At the polymer concentration used in this study, the density and stiffness of the gelMA network would limit cell migration, while mass transfer and diffusion of bioactive molecules are largely unhindered within the same

gel matrix[58]. Thus, in these layered co-culture models, the only effective communication between the cells in adjacent layers is through secreted factors. As a result, the co-cultures appeared to promote cell proliferation with a consequent reduction in overall sGAG/DNA ratio. Nevertheless, consistent chondrogenic differentiation was observed, especially in the MSC/ACPC constructs and the trends detected in each layer were similar to what was found for the monocultures, with MSCs producing more sGAGs and collagen type II, followed by ACPCs and chondrocytes. This suggests the feasibility of combining ACPCs and MSCs to create zonal constructs mimicking the zonal distribution of GAGs present in native cartilage, and in recreating the cell phenotypes present during rapid growth of cartilage in immature cartilage[59].

To verify this, bioprinted layered, zonal-like constructs were created as a proof-of-concept. The superficial zone of the construct was obtained from the ACPC-laden bioink, while the middle/deep region contained MSCs. Unmodified gelMA does not exhibit a marked yield stress, and is therefore difficult to print with high shape fidelity[9,60,61]. Thus, a sacrificial pluronic frame was used to preserve the architecture of the construct during printing. This supporting role of pluronic also allowed the generation of more complex, anatomical shapes, such as the caudal end of a femoral condyle. Nevertheless, reinforcing strategies, such as the co-printing with stiffer materials will be required to provide biomechanical stability, especially in the biofabrication of large joint components[62]. ACPCs were viable after the process of printing, pluronic removal and UV crosslinking, with cell viability values comparable to those observed for MSCs undergoing the same process, showing that the process is also non-harmful for ACPCs. The mechanism of crosslinking in methacryloyl-based hydrogels has been previously demonstrated to exert a protective effect on encapsulated cells, as it utilizes the free-radicals generated by the UV irradiation [63,64]. Nevertheless, to further reduce potential concerns of UV-A light, crosslinking chemistries that require reduced UV exposure, such as thiol-ene click reactions [65], or even alternatives based on visible light [46], could be applied in the future for encapsulating ACPCs, and in general for bioprinting.

During printing, cells are subjected to a range of shear stresses, depending on the gauge and shape of the extrusion nozzle, the extrusion mechanism, the printing pressure and the bioink rheological properties. Recent studies have highlighted that above a particular shear stress threshold chondrogenic differentiation is impeded, even in absence of harmful effects on viability, especially with viscous hydrogels using a

microvalve-based extrusion system[66]. The gelMA-based bio-ink, as used in this study, was also extruded through a microvalve, but we found that higher cell functionalities like differentiation potential and cartilage matrix deposition were preserved both for ACPCs and MSCs, further demonstrating the suitability of the gel and the chosen printing set-up for cartilage bioprinting. Overall, the zonal construct showed an inhomogeneous distribution of sGAGs and collagen type II, with the concentration of these macromolecules increasing from the ACPC- to the MSC-laden zone, a zonal difference approximating the one present in native cartilage.

In mono- and co-cultured bio-inks, all cell types displayed consistent production of collagen type I alongside collagen type II, as shown in the histological and mRNA analyses. In bioprinted constructs, the higher ratio of collagen type II to type I was more evident compared to that observed for the cast hydrogels, possibly suggesting a positive effect of stresses involved in the printing process on cell differentiation. Collagen type I is physiologically present in the superficial zone of native cartilage at the articulating boundary[67], and often in repair tissue[68], but is also usually associated with a fibrocartilage phenotype. Furthermore, the lack of mechanical loading may add to the potential of fibrous tissue formation, which has previously been observed in limb development studies utilizing the chick model[69]. Previous studies using gelMA as a bioink reported a more fibrocartilage-like matrix production from MSCs[70]. Modification of the bioink may be necessary to enhance the quality of the cartilage tissue in future experiments, as cell response can be tuned modifying the microenvironment of the hydrogel[71]. For instance, gelMA could be functionalized with hyaluronic acid, which has been previously demonstrated to inhibit the synthesis of collagen type I by encapsulated chondrocytes substantially[53], and has been successfully used to promote cartilage formation in other hydrogel systems [72]. Modification of gelatin bioinks with silk was also shown to modulate and improve chondrogenesis by MSC[73], and it may be interesting to evaluate ACPC. It is clear that the use of ACPCs in regenerative medicine is still in an early stage, and different biomaterials and culture conditions will have to be tested to optimize their chondrogenic potential. Mechanical conditioning has already been shown beneficial to guide ACPC differentiation[29]. Moreover, the differentiation media used in this study contain TGF- β 1 as main chondrogenic factor. This recipe has been optimized for MSCs, but a different formulation may be optimal for ACPCs[26], and it is expected that as more research will be performed involving these cells from both animal and human origin, the optimal culture conditions will be identified.

Another important aspect is the biomechanical profile of the constructs. While the *in vitro* maturation greatly increased the compressive properties of the gel, these are still not matching native cartilage. The compressive stiffness of mature equine cartilage ranges between 0.8 and 1.2 MPa[74], while our values fall in the range of immature cartilage[75], and may further be improved upon maturation through mechanical conditioning. Bioprinting of ACPCs may benefit with reinforcement strategies using highly organized microfibrinous meshes, *i.e.* produced by melt electrospinning writing[47] or through co-printing with reinforcing structures[6,76], which have already been previously described and demonstrated potential to approximate the behavior of native cartilage. Convergence of these biofabrication approaches may lead to a new generation of load-bearing, hydrogel-based cartilage constructs. These new strategies could be further characterized with an in-depth analysis of the biomechanical profile of cartilage constructs, *i.e.* via (nano)indentation, tensile testing, shear and friction at the surface, analysis of viscoelastic properties and response to cyclic stresses. While our present study focused on the biological performance of ACPCs, in relation to MSCs and chondrocytes, it did not provide such comprehensive mechanical characterization of the engineered constructs. Alongside accurate selection of cell sources for cartilage tissue engineering, future work should consider such mechanical testing, in order to evaluate the biomimicry of the anisotropic biomechanical profile of native articular cartilage, both under compression and tension, since achieving constructs mimicking the latter is still major challenge [77,78]. In this perspective, sets of mechanical tests, that include nanoindentation and single edge notch tests, have been previously proposed to compare native cartilage to engineered hydrogel constructs [79].

Finally, a new range of opportunities will be available for ACPCs for cartilage repair. Hydrogel encapsulation can be used with complementary strategies already proposed in the biomaterials community, including the use of hydrogels mimicking the developmental processes of cartilage, zonal gradients of growth factors and cell density, and mechanical stimulation and reinforcement strategies.

5. Conclusions

ACPCs are promising sources for cartilage regenerative medicine and biofabrication, and the encapsulation in gelMA hydrogels allowed the formation of 3D cartilage constructs *in vitro*. The interplay of ACPCs with

chondrocytes and MSCs supported neo-cartilage synthesis in layered co-cultures, indicating the possibility to use ACPCs also as a complementary cell source in cartilage constructs to produce functionally relevant differentiated tissue and to also act as a pool of stem cells for further growth and remodeling. The amount and quality of neo-cartilage matrix produced by ACPCs was superior to that generated by encapsulated expanded chondrocytes. Even though ACPC-laden hydrogels showed a lower production of ECM components compared to MSC-laden ones, ACPCs displayed distinctive phenotypic features, particularly a low expression of collagen type X and a high expression of PRG4, suggesting a priming toward a phenotype similar to superficial zone chondrocytes. When used as bioink, the ACPC-gelMA combination could be safely printed and combined with MSCs in a zonal-like architecture leading to a biomimetic GAG distribution. As the compressive mechanical properties of our hydrogel-only constructs did not reach those of adult articular cartilage, combination with reinforcement strategies or bioreactor culture will be recommended to fully address the complex mechanical behavior of cartilage under compression, but also in response to tensile and shear stresses. Consequently, future studies focusing on in-depth biomechanical characterization will play an important role in the field of cartilage tissue engineering. Importantly, further research on ACPC biology, 3D culture and bioprinting will be required to fully recapitulate the zonal organization of native cartilage. Overall, the results of this study provide important insights for the design of the next generation of cell- and biomaterial-based articular cartilage therapies.

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Conflict of interest

The authors declare no conflicts of interest.

Author Contributions

R.L., W.R.W., R.vW., I.M.K., J.M. designed the experiments; R.L., W.R.W., I.A.O. performed the experiments; A.M., M.vR., provided technical support and performed gene expression and histological analysis; R.L. W.R.W., R.vW., Y.Z., I.M.K., J.M. analyzed data; R.L. wrote the paper with inputs from all authors. All authors have approved the final article.

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Appendix A.

Supplementary Information

Table ST1: RT-PCR primers for the characterization of the progenitor cells from cartilage and bone marrow.

Target gene	Primer sequence (5' to 3')	Amplicon size (bp)
HPRT1	f: CAAGCTTGCTGGTGAAAAG r: GGCATATCCTACGACAACT	95
CD13	f: CTGAGTGGAGAGACAGAGTA r: CTGGAAATACTCGAAGAGGG	147
CD29	f: CTGGAGATGGGAAACTTGG r: GTTCCTACTGCTGACTTAGG	229
CD31	f: CAGAATCCTTCTCTATGCCC r: CATGGCCATCACTGAGTAG	194
CD34	f: GACTCAAGGTATCTGCCTG r: CCTGTTCTTTCTCACAGAGG	104
CD44	f: CTGGGGACTCTGCCTC r: TAGCGGCCATTTTTCTCC	99
CD45	f: TTGAACGGCCTTGAACC r: CTTGGCACCTTCAGTACC	153
CD49d	f: CTACAACCTGGACACCGAG r: GTCCGGTCTGGATTCTTTC	201
CD73	f: TCCGGACTTTATTTGCCG r: CAGAGGTGACTATGAATGGG	346
CD90	f: CTCTACACATGCGAACTCC r: CTCGCACTTGACCAGTTT	90
CD105	f: CAGTAATGAGGTGGTCGTC r: CTGAGGTAGAGGCCAG	108
CD106	f: ACTCTTACTTGTGCACGG r: CCACTGAAACTGATCTCTGG	103
CD146	f: TCCGTGTGTACAAAGCTC r: GTACCAGATGACCTGAGGA	137
CD166	f: GTCTTCTGCCTCTTGATCG r: CTGTCTTTGACTCTGGGAC	223

Table ST2: qPCR primers for neo-cartilage synthesis.

Target gene	Primer sequence (5' to 3')	Amplicon size (bp)
HPRT1	hypoxanthine phosphoribosyltransferase 1 f: AAGCTTGCTGGTGAAAAG r: GCATATCCTACGACAACT	95
ACAN	aggrecan f: AAGACAGGGTCTCGCTGCCAA r: ATGCCGTGCATCACCTCGCA	115
COL2A1	α 1 chain, collagen type II f: GGCAATAGCAGGTTACGTACA r: CGATAACAGTCTTGCCCCACTT	79

COL1A1	$\alpha 1$ chain, collagen type I	f: CGTGACCTCAAGATGTGC r: AGAAGACCTTGATGGCGT	94
PRG4	Proteoglycan 4	f: CTTCCCATTTACTTGTGCTG r: TAGAATACCCTTCCCCACAT	103
COL10A1	$\alpha 1$ chain, collagen type X	f: GGGAAACGGGATATGGTGCT r: GTCCCCTTTCTCCCGGAATG	168
COMP	cartilage oligomeric protein	f: CCACGTGAATACGGTCACAG r: ACGTCTGCTCCATCTGCTTC	104

Table ST3: F-values and degrees of freedom relative to the one way ANOVA tests performed to identify differences between the experimental groups at any given time point.

Monoculture - biochemical analysis and stiffness

	Day 1	Day 28	Day 56
DNA/dwt	F (2, 15) = 18.78	F (2, 15) = 0.2680	F (2, 14) = 16.34
GAG/dwt	F (2, 9) = 1.717	F (2, 12) = 34.45	F (2, 14) = 16.34
GAG/DNA	F (2, 10) = 2.544	F (2, 15) = 28.23	F (2, 14) = 15.66
DMA	F (3, 20) = 3.045	F (3, 20) = 12.00	F (3, 20) = 29.72

Monoculture - gene expression

	Day 1	Day 28	Day 56
ACAN	F (2, 6) = 3.031	F (2, 6) = 159.6	F (2, 6) = 38.70
COL2A1	F (2, 6) = 0.6004	F (2, 6) = 646.9	F (2, 6) = 16.08
COL1A1	F (2, 6) = 3.488	F (2, 6) = 2.661	F (2, 6) = 9.524
COMP	F (2, 6) = 4.989	F (2, 6) = 0.7556	F (2, 6) = 4.524
PRG4	F (2, 6) = 4.034	F (2, 6) = 2.221	F (2, 6) = 123.6
COL10A1	F (2, 6) = 12.02	F (2, 6) = 70.93	F (2, 6) = 28.80

Co-culture - biochemical analysis

	Day 1	Day 28	Day 56
DNA/dwt	F (5, 18) = 0.8620	F (5, 30) = 21.43	F (5, 28) = 9.219
GAG/dwt	F (5, 17) = 0.6024	F (5, 29) = 4.564	F (5, 30) = 18.21
GAG/DNA	F (5, 17) = 0.8402	F (5, 30) = 8.552	F (5, 29) = 15.39

Viability after printing

	Day 1	Day 14
LIVE/DEAD	F (4, 10) = 0.5730	F (4, 10) = 0.4508

Video SV1: Series of cross sections of the printed hydrogel constructs after removal of the sacrificial pluronic, showing the inner porosity of the construct.

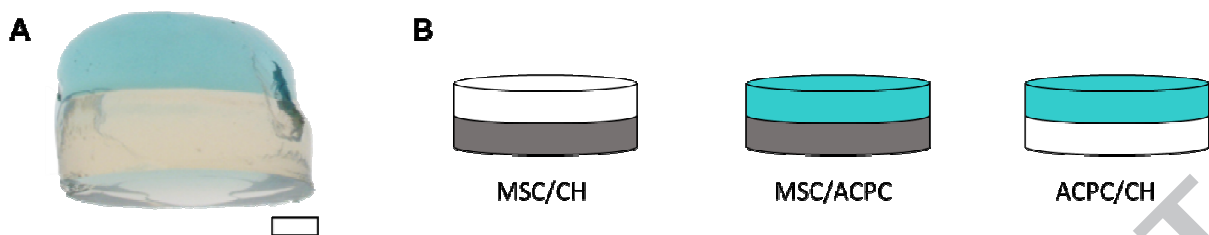


Figure S1: Representation of the layered 3D co-culture system. (A) Stereomicroscopy image showing the macroscopic appearance of the double layer constructs (side view). The upper layer is molded to display curved corners, to facilitate identification during histological analysis. Moreover, the upper layer is stained with a dye (fast green) for illustrative purposes. Scale bar is 1 mm. (B) Schematic summarizing the co-culture groups analyzed in this study.

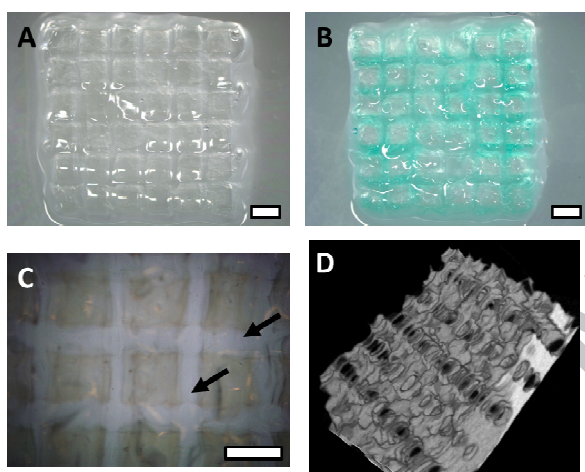


Figure S2: Example of a printed gelMA construct. (A) Once the pluronic filaments are removed, pores in the shape of open channels are left. (B) Injection of a blue coloured dye into this network shows the interconnection between these pores. (C) Front view of the printed constructs, with the black arrows pointing at the channels left by the pluronic removal. (D) μ CT 3D reconstruction showing the interconnected porous network inside the printed constructs. Scale bar is 1 mm.

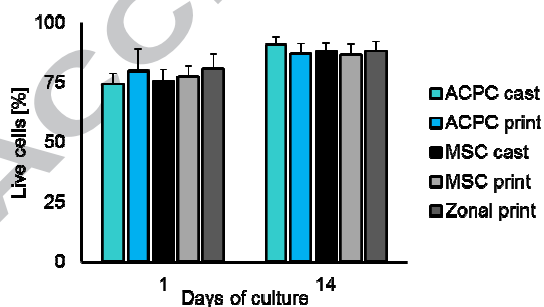


Figure S3: Cell viability in bioprinted MSC, ACPC and zonal constructs, compared to cast hydrogels. No significant difference was detected between the samples. Viability values ranged from about 75% (day 1) to 90% (day 14).