



PROGENITOR CELLS IN AURICULAR CARTILAGE DEMONSTRATE CARTILAGE-FORMING CAPACITY IN 3D HYDROGEL CULTURE

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Abstract

Paramount for the generation of auricular structures of clinically-relevant size is the acquisition of a large number of cells maintaining an elastic cartilage phenotype, which is the key in producing a tissue capable of withstanding forces subjected to the auricle. Current regenerative medicine strategies utilize chondrocytes from various locations or mesenchymal stromal cells (MSCs). However, the quality of neo-tissues resulting from these cell types is inadequate due to inefficient chondrogenic differentiation and endochondral ossification, respectively. Recently, a subpopulation of stem/progenitor cells has been identified within the auricular cartilage tissue, with similarities to MSCs in terms of proliferative capacity and cell surface biomarkers, but their potential for tissue engineering has not yet been explored. This study compared the *in vitro* cartilage-forming ability of equine auricular cartilage progenitor cells (AuCPCs), bone marrow-derived MSCs and auricular chondrocytes in gelatin methacryloyl (gelMA)-based hydrogels over a period of 56 d, by assessing their ability to undergo chondrogenic differentiation. Neocartilage formation was assessed through gene expression profiling, compression testing, biochemical composition and histology. Similar to MSCs and chondrocytes, AuCPCs displayed a marked ability to generate cartilaginous matrix, although, under the applied culture conditions, MSCs outperformed both cartilage-derived cell types in terms of matrix production and mechanical properties. AuCPCs demonstrated upregulated mRNA expression of elastin, low expression of collagen type X and similar levels of proteoglycan production and mechanical properties as compared to chondrocytes. These results underscored the AuCPCs' tissue-specific differentiation potential, making them an interesting cell source for the next generation of elastic cartilage tissue-engineered constructs.

Keywords: Auricular cartilage, tissue engineering, cartilage progenitor cells, hydrogel, 3D culture.

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Introduction

A significant challenge in cartilage tissue engineering is the recruitment of a sufficient number of cells for the generation of large tissue constructs (Bichara *et al.*, 2012). The necessity for larger constructs is exemplified by the case of auricular reconstruction required for the congenital disorder microtia and for defects of the auricle caused by injury and disease. For the human auricle, estimates of the number of cells required for tissue regeneration range between 100 and 150 million (Bichara *et al.*, 2012), with the obvious challenge of obtaining this number of cells from an autologous source.

Although considerable progress has been made using a variety of cells for auricular cartilage tissue

engineering, there is no definitive conclusion on which cell type is capable of providing the most favorable clinical outcome for tissue-engineered cartilage constructs. Originating from the native tissue, chondrocytes are a logical cell source for the generation of neocartilage. Auricular (Bichara *et al.*, 2014; Cao *et al.*, 1997; García-López *et al.*, 2015; Isogai *et al.*, 2006; Kusuhara *et al.*, 2008; Nakao *et al.*, 2017; Papadopoulos *et al.*, 2011; Park *et al.*, 2004; Pomerantseva *et al.*, 2016; Reiffel *et al.*, 2012; Sanz *et al.*, 2007; Tseng *et al.*, 2014; Xu *et al.*, 2005), microtia (Ishak *et al.*, 2015; Kamil *et al.*, 2004b; Nakao *et al.*, 2017), nasoseptal (Ávila *et al.*, 2015; Bichara *et al.*, 2010; Haisch *et al.*, 2002; Isogai *et al.*, 2006; Kusuhara *et al.*, 2008), costal (Isogai *et al.*, 2006; Kusuhara *et al.*, 2008) and articular chondrocytes (Isogai *et al.*, 2006;

Kamil *et al.*, 2004a; Kusuha *et al.*, 2008; Mizuno *et al.*, 2014) are all used for tissue-engineering auricular cartilage. Specific characteristics and ability for chondrogenesis can differ depending on the origin of the chondrocytes. Where the articular and costal cartilage are of mesenchymal origin, the developmental origin of the auricular cartilage of the *pinna* is still controversial. The auricle originates from two pharyngeal arches that have contributions from all three embryonic layers (Wright, 1997; Graham, 2001; Luquetti *et al.*, 2011; Tracy *et al.*, 2013). Nevertheless, fully differentiated cells from various cartilage origins generally demonstrate the capacity to produce cartilage-like extracellular matrix *in vitro* and *in vivo* (Ahmed and Hincke, 2014; Bichara *et al.*, 2010; Isogai *et al.*, 2006; Kusuha *et al.*, 2008; Pleumeekers *et al.*, 2013). However, only a small amount of tissue can be harvested from the patient, requiring extensive *in vitro* expansion of the isolated cells to obtain a sufficient cell's number for the production of clinically-relevant tissue-engineered cartilage constructs. The proliferative potential of chondrocytes is naturally low (Phull *et al.*, 2016) and their cartilage-forming ability is known to decline with extended cultivation (Homicz *et al.*, 2002; Schnabel *et al.*, 2002). In fact, repeated passaging of chondrocytes induces the loss of their chondrogenic phenotype (Saadeh *et al.*, 1999) and, ultimately, leads to progressive dedifferentiation. To some extent, re-differentiation can be attained under specific culture conditions, *e.g.* in three-dimensional (3D) environments (Benya and Shaffer, 1982) and in the presence of appropriate growth factors (Jakob *et al.*, 2001; van Osch *et al.*, 2001), yet the revenue is limited (van Osch *et al.*, 2001). The resulting neo-tissue is often of a fibrocartilagenous quality and exhibits inferior biochemical and mechanical properties as compared to native cartilage tissue (Chung *et al.*, 2006; Demoor *et al.*, 2014; Nabzdyk *et al.*, 2009; Williams *et al.*, 2010). In the past decade, stem cells have gained increasing interest for tissue engineering applications because of their capacity for self-renewal and multilineage differentiation. For example, bone-marrow-derived mesenchymal stromal cells (MSCs) can be extensively expanded (Gardner *et al.*, 2013) and demonstrate the ability to differentiate into various cell types, including chondrocytes, osteoblasts and adipocytes (Pittenger *et al.*, 1999). Numerous studies have successfully used MSCs for the generation of cartilage (Cheng *et al.*, 2014; Seda Tigli *et al.*, 2009; Vinardell *et al.*, 2012), sometimes in co-culture with auricular chondrocytes (Liu *et al.*, 2010; Pleumeekers *et al.*, 2015; Zhao *et al.*, 2016). Nevertheless, the risk of hypertrophic growth, terminal differentiation and subsequent tissue calcification remains an important limitation when using MSCs for cartilage tissue engineering (Williams *et al.*, 2010; Gawlitta *et al.*, 2010). In addition, bone marrow-derived MSCs are not involved in the development of native auricular

cartilage and are, as articular, costal and nasoseptal chondrocytes, not predisposed to produce the elastic fibers required for this elastic-type cartilage (Kusuha *et al.*, 2008).

Tissue-derived stem/progenitor cells exhibit stem-cell-like qualities, such as self-renewal and multipotency, yet are embedded within the target tissue in niches and are primed to differentiate to that tissue (Jayasuriya and Chen, 2015). The identification of resident progenitor cell populations in articular cartilage (Dowthwaite *et al.*, 2004; Williams *et al.*, 2010) as well as in auricular and tracheal perichondrium (Derks *et al.*, 2013; Kobayashi *et al.*, 2011b; Togo *et al.*, 2006) has opened up new pathways for cartilage tissue engineering. Like MSCs, which can undergo up to 70 population doublings (Christodoulou *et al.*, 2013), cartilage progenitor/stem cells retain proliferative ability for up to 60 population doublings (Williams *et al.*, 2010), demonstrating potential for accumulating large cell numbers starting from a single cell. In addition, the cells maintain multipotent differentiation ability while expressing chondrocyte-specific characteristics (McCarthy *et al.*, 2012). It is hypothesized that these tissue-specific progenitor cells are highly primed to differentiating into the chondrogenic lineage (Jayasuriya and Chen, 2015; McCarthy *et al.*, 2012).

Auricular cartilage tissue engineering may benefit greatly from a source of tissue-specific cells that can be expanded up to large numbers without losing their differentiation potential. Nevertheless, the auricular cartilage itself was, until recently, unknown to harbor a progenitor cell population. The presence of colony-forming, multipotent progenitor cells in auricular cartilage has recently been confirmed (Xue *et al.*, 2016), though these cells have not yet been explored for a tissue-engineering purpose. The objective of the current study was to evaluate the cartilage-forming ability of such auricular cartilage progenitor cells in a 3D hydrogel culture. It was hypothesized that these cells, despite *in vitro* expansion, could outperform auricular chondrocytes and bone-marrow-derived MSCs with regards to production of auricular cartilage.

Mimicking the hydrated environment of native cartilage, hydrogels are particularly attractive for cartilage regenerative strategies. Gelatin methacryloyl (gelMA)-based hydrogels are a versatile group of biomaterials shown to facilitate cartilage-like matrix production for chondrocytes, articular cartilage progenitor cells as well as MSCs, supporting both cell viability and mechanical properties (Levato *et al.*, 2017; Daly *et al.*, 2016; Gao *et al.*, 2015; Schuurman *et al.*, 2013). In this study, cell-laden gelMA hydrogels were cultured for up to 8 weeks in chondrogenic differentiation media and harvested at day 1, 28 and 56 for mechanical testing, biochemistry, gene expression, histology and immunohistochemistry for the assessment of cartilage-specific properties.

Material and Methods

Isolation of cells

Primary auricular chondrocytes (AuCHs) and auricular cartilage progenitor cells (AuCPCs) were obtained from deceased equine donors, which were kindly provided by a local slaughterhouse. Bone-marrow-derived MSCs were obtained from healthy equine donors. All tissues and cells were obtained according to the guidelines of the Institutional Animal Ethical Committee (the Netherlands).

AuCHs and AuCPCs were harvested from the auricles of fresh equine cadavers (3- to 10-year-old; $n = 3$). The ears were cut off at the base, shaved, thoroughly washed with soap and soaked for circa 15 min in Betadine® (Meda Pharma, Amstelveen, the Netherlands). Under sterile conditions, an incision through the skin was made on the dorsal side along the longitudinal axis and the skin and subcutaneous tissue were dissected. Ensuring the harvest of auricular cartilage exclusively, the perichondrium was fully removed by carefully scraping the tissue off with surgical tools. Cartilage chips were sectioned off the *scapha* of the ear, washed in sterile phosphate-buffered saline (PBS) and, subsequently, minced into 1 mm² pieces. The tissue was digested in 0.2 % pronase (Roche) for 2 h followed by a 16 h 0.075 % collagenase type II (Worthington Chemical Corporation, Lakewood, NJ, USA) digestion at 37 °C. Next, the solution was filtered through a 70 µm cell strainer and centrifuged for 5 min at 300 ×g to obtain a cell pellet. AuCHs were washed in sterile PBS, counted with a hemocytometer and stored at passage 0 in liquid nitrogen until further use.

For each donor, an aliquot of the freshly-isolated cells was saved for the isolation of AuCPCs, for which the cells were subjected to a fibronectin adhesion assay as previously described (Dowthwaite *et al.*, 2004; Williams *et al.*, 2010). Briefly, cells suspended in serum-free Dulbecco's modified Eagle medium (DMEM; 31966, Gibco) were plated at a density of 500 cells/cm² on fibronectin-coated tissue culture plates. After 20 min of incubation at 37 °C, the non-adherent cells were carefully removed. Attached progenitor cells were cultured in chondroprogenitor expansion media, consisting of DMEM supplemented with 10 % v/v fetal bovine serum (FBS; Lonza), 0.2 mM L-ascorbic acid 2-phosphate (Sigma-Aldrich), 100 U/mL penicillin (Life Technologies), 100 µg/mL streptomycin (Life Technologies) and 5 ng/mL basic fibroblast growth factor (bFGF; Peprotech, London, UK). After 6 d of culture, colonies consisting of > 32 cells were harvested. Monoclonal colonies were pooled and expanded until passage 3, when they were stored in liquid nitrogen until further use.

MSCs were obtained from bone marrow aspirates from the sternum of healthy equine donors (3- to 10-year-old; $n = 3$) and the mononuclear fraction was isolated following a previously described protocol using a Ficoll®-Paque density gradient (GE

Healthcare) (Visser *et al.*, 2014). After isolation, MSCs were cultured in MSC expansion medium, consisting of alpha modification minimum essential medium (αMEM) (22561, Gibco) supplemented with 0.2 mM L-ascorbic acid 2-phosphate (Sigma-Aldrich), 10 % FBS (Lonza), 100 U/mL penicillin (Life Technologies), 100 µg/mL streptomycin (Life Technologies) and 1 ng/mL bFGF (Peprotech) until passage 3 and subsequently stored in liquid nitrogen until further use.

Characterization of equine auricular chondroprogenitor cells

AuCHs (at passage 1), AuCPCs and MSCs (both at passage 3) were characterized by comparison of gene expression of cell surface markers and assessment of multilineage differentiation potential in two dimensional (2D) culture.

The expression of cell membrane markers was evaluated in duplicate by a reverse transcriptase-polymerase chain reaction (RT-PCR), comparing the transcriptome of AuCHs, AuCPCs and MSCs. Analyzed target genes included *CD13*, *CD29*, *CD31*, *CD34*, *CD44*, *CD45*, *CD49d*, *CD73*, *CD90*, *CD105*, *CD106*, *CD146* and *CD166* (primer sequences and expected amplicon sizes are reported in Table 1), which were compared to expression of the housekeeping gene hypoxanthine phosphoribosyltransferase (*HPRT1*). RNA was isolated using the RNeasy mini kit (Qiagen), according to the manufacturer's instructions. Isolated mRNA was quantified by UV-vis spectrophotometry with a Nanodrop 2000 (Thermo Scientific) to serve as template for polymerase chain reaction (PCR). The amplification of RNA was carried out using a SuperScript® One-Step RT-PCR System with Platinum® Taq DNA Polymerase (Life Technologies) and the PCR products were run on agarose gel stained with ethidium bromide. Subsequently, the amplicons were imaged using a UV transilluminator (ProXima 10 Phi; Isogen Life Sciences, De Meern, the Netherlands).

Multipotency was evaluated in duplicate through an *in vitro* trilineage differentiation assay in which cells were directed towards bone, fat or cartilage by culturing them in osteogenic, adipogenic or chondrogenic differentiation media, respectively. For osteogenic and adipogenic differentiation, cells were plated in 6-well culture plates at a density of 2×10^5 cells/well and cultured in chondroprogenitor expansion medium until sub-confluency, before initiating differentiation. Osteogenic differentiation medium consisted of αMEM (Gibco) supplemented with 10 % v/v FBS (Lonza), 100 U/mL penicillin (Life Technologies), 100 µg/mL streptomycin (Life Technologies), 0.2 mM L-ascorbic acid 2-phosphate (Sigma-Aldrich), 20 mM β-glycerol phosphate (Sigma-Aldrich) and 100 nM dexamethasone (Sigma-Aldrich). Adipogenic medium consisted of αMEM (Gibco) supplemented with 10 % v/v FBS (Lonza), 100 U/mL penicillin (Life Technologies),

Table 1. RT-PCR forward (Fw) and reverse (Rv) primers sequences used for characterization of AuCH, AuCPC and MSC.

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

100 µg/mL streptomycin (Life Technologies), 0.01 mM Indomethacin (Sigma-Aldrich), 83 mM 3-Isobutyl-1-methylxanthine (Sigma-Aldrich) and 1.72 µM bovine pancreas-derived insulin (Sigma-Aldrich). Culture medium was refreshed every 3 d. For chondrogenic differentiation, 2.5×10^5 cells were pelleted by centrifugation at $300 \times g$ in 15 mL Falcon® tubes. Subsequently, the pellets were cultured in chondrogenic differentiation medium, consisting of DMEM supplemented with 1 % v/v insulin-transferrin-selenous acid (ITS+ Premix; Corning), 0.2 mM L-ascorbic acid 2-phosphate (Sigma-Aldrich), 100 U/mL penicillin (Life Technologies), 100 µg/mL streptomycin (Life Technologies), 100 nM dexamethasone (Sigma-Aldrich) and 10 ng/mL transforming growth factor β1 (TGF-β1; Peprotech). After 14 d of culture, osteogenic commitment was evaluated by alizarin red S staining to observe calcified matrix deposition, whereas adipogenic differentiation was assessed with oil red O staining to visualize the formation of intracellular lipid vesicles. Cell pellets were embedded in paraffin and 5 µm-thick sections were stained with safranin O to visualize glycosaminoglycans (GAGs), indicative of chondrogenic differentiation.

Fabrication of cell-laden hydrogel constructs for 3D culture

Following a previously published protocol (Melchels *et al.*, 2014), the hydrogel gelatin methacryloyl (gelMA) was synthesized by functionalizing gelatin type A (obtained from porcine skin; Sigma-Aldrich) in PBS with methacrylic anhydride groups, to obtain a hydrogel with a 80 % degree of functionalization. Subsequently, a 10 % w/v solution of gelMA was supplemented with 0.1% w/v 2-hydroxy-1-[4-(2-hydroxyethoxy)phenyl]-2-methyl-1-propanone (Irgacure 2959; BASF, Ludwigshafen, Germany) as a photoinitiator. AuCHs, AuCPCs or MSCs, expanded beforehand until passage 1, 4 and 4, respectively, were homogeneously suspended in the hydrogel at 37 °C at a concentration of 1.5×10^7 cells/mL. The cell-laden gel was immediately casted into a custom-made Teflon™ mold to produce cylindrical samples (diameter = 6 mm, height = 2 mm) and subsequently chemically crosslinked by UV irradiation for 5 min (wavelength λ = 365 nm, intensity E = 3 mW/cm², at a height of 2 cm; 144 portable UV lamp, Vilber Lourmat, Eberhardzell, Germany), to trigger free-radical polymerization. Cell-free hydrogel samples (3 replicates per time point) were used as controls

Table 2. qPCR forward (Fw) and reverse (Rv) primers sequences used for evaluation of cartilage-like tissue synthesis.

Target gene	Full name	Primer sequence (5' to 3')	Amplicon size (bp)
HPRT1	Hypoxanthine phosphoribotransferase 1	Fw: AAGCTTGCTGGTGAAG	95
		Rv: GCATATCCTACGACAACT	
ACAN	Aggrecan	Fw: AAGACAGGGTCTCGCTGCCAA	115
		Rv: ATGCCGTGCATCACCTCGCA	
COL1A1	Collagen type I, α 1 chain	Fw: CGTGACCTCAAGATGTGC	94
		Rv: AGAAGACCTTGATGGCGT	
COL2A1	Collagen type II, α 1 chain	Fw: GGCAATAGCAGGTTACGTACA	79
		Rv: CGATAACAGTCTTGCCCCACTT	
COL10A1	Collagen type X, α 1 chain	Fw: GGGAAACGGGATATGGTGCT	168
		Rv: GTCCCTTTCTCCCGGAATG	
COMP	Cartilage oligomeric protein	Fw: CCACGTGAATACGGTCACAG	104
		Rv: ACGTCTGCTCCATCTGCTTC	
RUNX2	Runt-related transcription factor 2	Fw: GCAAGGTTCAACGATCTGA	248
		Rv: GGGACACCTACTCTCATACT	
ELASTIN	Elastin	Fw: TGGAGTCCCAGGTGTTGTTG	137
		Rv: CATAGCCAGGAACCCCGAA	

and were prepared following the same steps. All hydrogel samples were cultured in chondrogenic differentiation medium for 1, 28 and 56 d at 37 °C and 5 % CO₂, refreshing media 3 times per week.

Gene expression of cartilage markers after chondrogenic differentiation

The relative gene expression of cartilage markers in cell-laden hydrogels ($n = 3$) was evaluated by qPCR at 1 and 56 d of culture. Analyzed markers included aggrecan (*ACAN*), cartilage oligomeric matrix protein (*COMP*), collagen type II (*COL2A2*), collagen type I (*COL1A1*), collagen type X (*COLXA1*), runt-related transcription factor 2 (*RUNX2*) and elastin. The expression levels were normalized against the housekeeping gene *HPRT1*. Primer sequences for each transcript are reported in Table 2.

Gel samples were mechanically ground in RLT buffer (Qiagen). From the lysate, mRNA was isolated using the RNeasy Mini Kit (Qiagen) and subsequently quantified with a Nanodrop 2000 (Thermo Scientific). A SuperScript[®] III Platinum SYBR Green One-Step qRT-PCR Kit (Life Technologies) was used for amplification of the mRNA and cDNA synthesis, which was performed with a LightCycler[®] 96 (Roche). The PCRminer algorithm (Zhao and Fernald, 2005) was used to calculate relative gene expression, Ct and efficiency values.

Biochemical analysis of cell-laden hydrogels

After 1, 28 and 56 d in culture, 4-6 replicates of each group of cell-laden hydrogels were taken for quantification of DNA and GAG content. Cell-laden hydrogel samples were frozen at -20 °C and subsequently lyophilized. The wet and dry weights were recorded during this process. Digestion of samples occurred overnight at 60 °C in 200 μ L papain digestion buffer (P3125; Sigma Aldrich),

consisting of 0.2 M NaH₂PO₄ (Merck) and 0.01 M ethylenediaminetetraacetic acid (EDTA; VWR) in milliQ water (pH = 6.0) supplemented with 250 μ L/mL papain solution (16-40 units/mg of protein) and 0.01 M cysteine (C9768; Sigma Aldrich).

Total DNA content was quantified using a Quant-iT PicoGreen dsDNA assay (Life Technologies) and compared to a standard of known concentrations of DNA. The fluorescence was measured at 485 nm excitation and 520 nm emission by a spectrofluorometer (Bio-Rad).

Total GAG content, as a measure of cartilage-specific matrix production, was quantified using a dimethylmethyleneblue (DMMB; pH = 3.0) assay. The 525/595 nm absorbance ratio was measured with a VersaMax plate reader (Molecular Devices, Warriner, UK). The sulfated GAG (sGAG) content was calculated using a standard of known concentrations of chondroitin sulfate C and corrected for the dilution factor.

Both dsDNA and sGAG content were normalized against dry weight. The ratio of GAGs per DNA was calculated to display the activity of single cells in producing cartilage-specific matrix.

Compressive mechanical testing of hydrogel constructs

An unconfined uniaxial compression test was performed to evaluate the mechanical properties of the cell-laden hydrogel samples after 1, 28 and 56 d in culture (4-6 replicates per time point). Using a dynamic mechanical analyzer (DMA Q800; TA Instruments, Assen, Belgium), samples were compressed at a -20 %/min strain rate to -30 %. Young's modulus was calculated as the slope of the initial linear segment (10-15 % strain) of the stress/strain curve.

Histology and immunohistochemistry

Deposition of the main components of cartilage extracellular matrix in cell-laden hydrogels were visualized by histology and immunohistochemistry on formalin-fixed, paraffin-embedded samples.

After 1, 28 and 56 d in culture, samples from each group were fixated in 4 % neutral-buffered formalin. Then, they were dehydrated through a graded ethanol series (70 %, 96 % and 100 % ethanol), cleared in xylene and embedded in paraffin. The samples were sectioned into 5 μ m-thick slices and deparaffinized prior to staining. For the identification of cartilage glycosaminoglycan deposition, a triple stain, consisting of hematoxylin (cell nuclei), fast green (collagens) and safranin O (proteoglycans), was applied. Deposition of collagens was evaluated by immunohistochemistry, with appropriate primary antibodies for collagen type II (II-II6B3; DSHB, Iowa City, IA, USA), collagen type I (sc-8784; Santa Cruz Biotechnology) and collagen type VI (5C6; DSHB). In addition, appropriate IgG were used as isotype controls. After deparaffinization, samples were treated with 0.3 % v/v H₂O₂ to block endogenous peroxidases. Antigen retrieval was performed with 1 mg/mL pronase (Roche) and 10 mg/mL hyaluronidase (H2126; Sigma-Aldrich), both applied for 30 min at 37 °C. Subsequently, tissue sections were blocked with bovine serum albumin (BSA, 5 % w/v in PBS) for 1 h at room temperature. The primary antibodies were incubated overnight at 4 °C, followed by a HRP-tagged secondary antibody for 1 h at room temperature, after which the staining was developed with 3,3-diaminobenzidine-horseradish peroxidase (Sigma-Aldrich). Cell nuclei were counterstained with hematoxylin. A von Kossa staining was applied to detect calcium precipitates indicating tissue mineralization. All stained sections were mounted in DPX mounting media (Millipore) and examined using a light microscope (Olympus BX51; Olympus).

Statistical analysis

Quantitative results are expressed as mean \pm standard error of the mean (SEM). The means of the experimental groups were compared at different time points by performing a two-way ANOVA with a Bonferroni *post-hoc* test. The statistical analyses

were performed using Graphpad Prism 7 (Graphpad software). A value of $p < 0.05$ was considered statistically significant.

Results

Equine auricular cartilage progenitor cells displayed stem cell qualities

Histological evaluation confirmed that the perichondrial layer attached to auricular cartilage was successfully removed and, therefore, it was assumed that a population of exclusively cartilage-derived cells was obtained after tissue digestion, prior to isolation of cartilage progenitor cells using differential adhesion to fibronectin (Fig. 1).

Gene expression of several surface markers was analyzed for isolated AuCHs, AuCPCs and MSCs by RT-PCR (Fig. 2a), demonstrating that the transcript profile of AuCPCs shared similarities with both AuCHs and MSCs. Like MSCs, AuCPCs were positive for the stem cell markers *CD73*, *CD90* and *CD105* and negative for the hematopoietic marker *CD34* and for the leukocyte marker *CD45*. Thus, at the transcript level, AuCPCs satisfied the minimal requirements to be classified as human MSCs (Dominici *et al.*, 2006). Furthermore, all three cell types were *CD29*⁺, *CD31*⁺, *CD106*⁺ and *CD166*⁺. Differences among cell types arose in the expression of *CD13*, *CD49d* and *CD146*. *CD13* was highly positive in AuCPCs, to a lesser extent in MSCs and negative in AuCHs. *CD49d* appeared positive in MSCs and AuCPCs and negative in AuCHs. Lastly, *CD146* was faintly positive in both MSCs and AuCPCs and negative in AuCHs.

Multipotency of AuCPCs was assessed through a trilineage differentiation assay. Like MSCs (Fig. 2c,e,f), AuCPCs demonstrated to be capable of differentiating towards bone (Fig. 2b), adipose tissue (Fig. 2d) and cartilage (Fig. 2f), confirming their multipotent potential.

Differential mRNA expression of cartilage markers in hydrogel culture

As measured by qPCR, AuCHs, AuCPCs and MSCs embedded in 3D hydrogels exhibited increased relative-fold expression levels of cartilage-specific

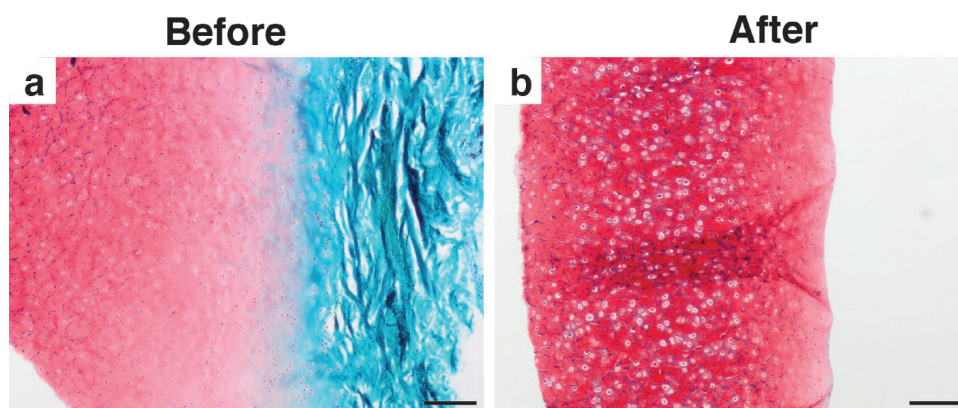


Fig. 1. Safranin O staining (a) before and (b) after utilizing the scraping method for the removal of perichondrium, confirming its complete removal from the auricular cartilage. Scale bars: 200 μ m.

gene transcripts over time, confirming their differentiation in the hydrogel (Fig. 3).

After 56 d of culture, aggrecan expression (Fig. 3a) was highest in AuCHs (48.8-fold upregulation), followed by AuCPCs (37.7-fold) and MSCs (15.7-fold). The same correlation was already apparent at day 1: mRNA expression of aggrecan was significantly more upregulated in AuCHs and AuCPCs (10.8- and 8.6-fold, respectively) than MSCs, expressing a 0.1-fold reduction. The expression of aggrecan increased over time in all three cell types, yet this increase was significant only in AuCHs.

A similar pattern was observed for the mRNA expression of *COMP* (Fig. 3b), which is a prominent non-collagenous matrix protein present in cartilage (Hedbom *et al.*, 1992; DiCesare *et al.*, 1995; Newton *et al.*, 1994). After 56 d of culture, AuCPCs exhibited a significantly higher expression level of *COMP* than MSCs (186.2-fold *versus* 64.2-fold, respectively), with AuCHs displaying a 146.7-fold increment. Likewise, AuCPCs already demonstrated the highest upregulation of *COMP* (39.1-fold) at day 1, followed by AuCHs (20.5-fold) and MSCs (0.9-fold). Both AuCHs and AuCPCs displayed a significant increase in expression levels over time, whereas MSCs did not.

After 56 d of culture, AuCHs and AuCPCs expressed *COL2A1* (Fig. 3c) at comparable levels, with a 51- and 60-fold increment, respectively. MSCs

exhibited the highest levels of *COL2A1* mRNA with an 89.1-fold increment at day 56, yet this difference was not significant. Expression of *COL2A1* was considerably lower (< 3.5-fold) in all three cell types at the beginning of the culture, indicating a significant increase over time in the case of AuCPCs and MSCs. Conversely, *COL1A1* (Fig. 3d) was already highly upregulated at the beginning of the culture in MSCs (59.9-fold increment), compared to a significantly lower 19.3-fold in AuCHs and 22.8-fold in AuCPCs. This trend reversed during chondrogenic differentiation, as in AuCHs, the expression of *COL1A1* significantly increased over time. Differences among cell types were not significant at day 56, yet the highest expression at day 56 was measured in AuCHs (77.2-fold increment), followed by equivalent levels in AuCPCs and MSCs (54.2- and 48.1-fold, respectively). AuCHs exhibited significantly higher levels of *COL10A1* at the end of culture (3-fold) in comparison to both AuCPCs and MSCs, which exhibited a 0.4-fold lesser expression (Fig. 3e). Both AuCHs and MSCs showed increasing yet non-significant levels of *COL10A1*, whereas, in AuCPCs, *COL10A1* expression levels decreased non-significantly over time.

At day 56, the expression of *RUNX2* (Fig. 3f), the master transcription factor during osteogenesis and endochondral ossification, was significantly higher in

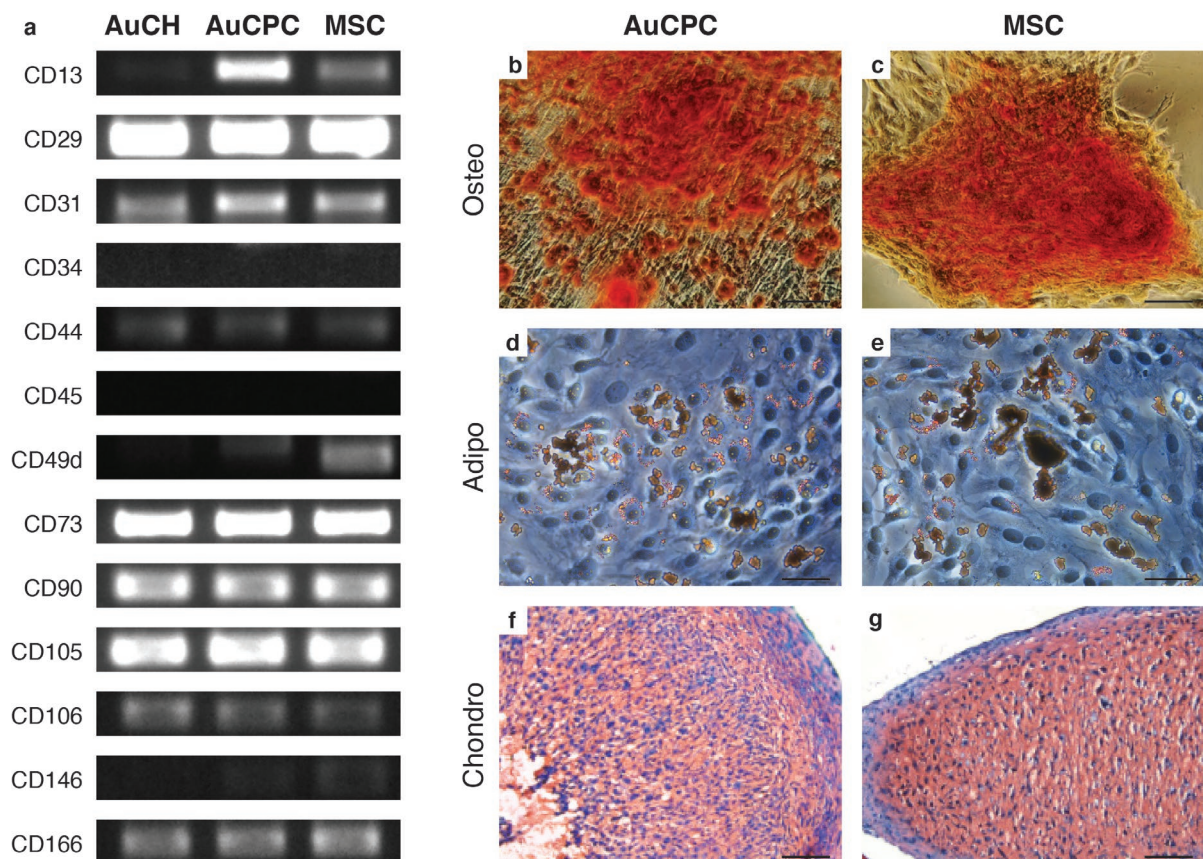


Fig. 2. Characterization of AuCPCs in comparison to AuCHs and MSCs. (a) The expression of several surface markers, analyzed by RT-PCR, was compared amongst the three cell types. Trilineage differentiation of AuCPCs and MSCs: (b,c) alizarin red staining demonstrating osteogenic differentiation, (d,e) oil red O staining demonstrating adipogenic differentiation and (f,g) safranin O staining demonstrating chondrogenic differentiation. Scale bars: 100 µm.

MSCs (2.5-fold) as compared to AuCHs (0.3-fold) and AuCPCs (0.0-fold). At the start of the culture, *RUNX2* was upregulated by 1.1-fold in AuCHs, by 1.9-fold in AuCPCs and by 2.1-fold in MSCs. AuCPCs showed a significant decrease in *RUNX2* expression over time to virtually zero at day 56.

The critical structural component of elastic cartilage is elastin. AuCHs exhibited higher mRNA expression levels of elastin (1.4-fold at day 1 and 2.6-fold at day 56) than MSCs (0.1-fold at day 1 and 0.4-fold at day 56). Although not significant, AuCPCs showed the highest upregulation of elastin, with

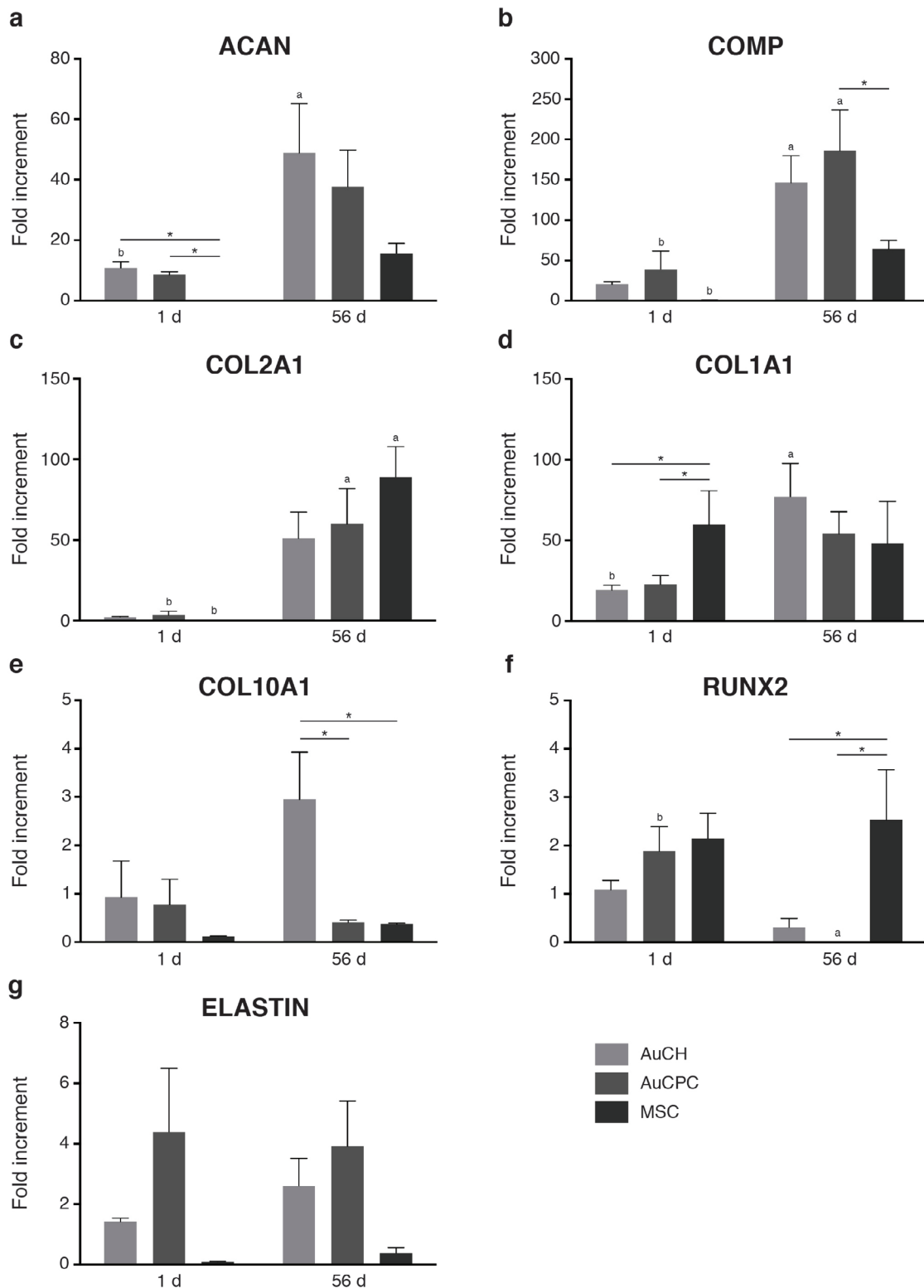


Fig. 3. Relative gene expression of (a) collagen type II, (b) collagen type I, (c) collagen type X, (d) aggrecan, (e) cartilage oligomeric matrix protein, (f) *RUNX2* and (g) elastin, as obtained from qPCR analysis of cell-laden hydrogels. Statistically significant differences of $p < 0.05$ for comparisons between cell types are marked with *. For each cell type, statistically significant differences of $p < 0.05$ are marked with ^a for a significant difference to day 1 and ^b for a significant difference to day 56.

4.4-fold and 3.9-fold increments at day 1 and 56, respectively (Fig. 3g).

Chondrogenic differentiation in hydrogels resulted in cartilage-specific matrix deposition

Cell proliferation and extracellular matrix deposition in cell-laden hydrogels was assessed by quantification of dsDNA and sGAG content, representative of cell number and proteoglycan quantity, respectively. As indicated by the increase in dsDNA content in the first 28 d of culture, AuCPCs and MSCs showed an ability to proliferate in the 3D gelMA hydrogel environment. The dsDNA content of AuCHs was significantly higher than AuCPCs and MSCs in the beginning of the culture period and stayed stable during the first 28 d; then, it significantly decreased to levels comparable to the other two cell types. These differences in cell numbers among groups at the beginning of culture could possibly be attributed to cell-loading inconsistencies. Nevertheless, dsDNA content among groups appeared to equalize over time (Fig. 4a).

The synthesis of neocartilage matrix was evaluated by the amount of sulfated GAGs present in the hydrogel samples. Over the course of chondrogenic culture, AuCHs, AuCPCs and MSCs produced significantly increasing amounts of sGAG, which were retained in the hydrogel matrix. AuCHs and

AuCPCs showed similar trends in total sGAG content (Fig. 4b) as well as in sGAG normalized to dsDNA content (Fig. 4c). After 56 d of culture, total sGAG content in AuCHs was $409.9 \pm 35.7 \mu\text{g}/\text{mg}$ and in AuCPCs $458.4 \pm 30.9 \mu\text{g}/\text{mg}$; sGAG/dsDNA was $509.9 \pm 56.8 \mu\text{g}/\text{mg}$ and $565.2 \pm 40.8 \mu\text{g}/\text{mg}$, respectively. MSCs initially appeared to lag behind in total sGAG content ($214.5 \pm 25.1 \mu\text{g}/\text{mg}$ at day 28, compared to $292.7 \pm 29.5 \mu\text{g}/\text{mg}$ for AuCHs and $308.3 \pm 15.8 \mu\text{g}/\text{mg}$ for AuCPCs), yet encompassed lower cell numbers in the samples. Accordingly, when normalized to dsDNA content, AuCHs, AuCPCs and MSCs performed similarly at the 28-d time point (304.9 ± 42.6 , 329.7 ± 19.9 and $296.9 \pm 38.1 \mu\text{g}/\text{mg}$, respectively). However, after 56 d of culture, when dsDNA levels were equivalent among groups, MSCs ($789.5 \pm 95.4 \mu\text{g}/\text{mg}$) significantly outperformed both cartilage-derived cell types, displaying sGAG/dsDNA values 1.6-fold higher than AuCHs ($508.9 \pm 56.8 \mu\text{g}/\text{mg}$) and 1.4-fold higher than AuCPCs ($565.2 \pm 40.8 \mu\text{g}/\text{mg}$).

Mechanical properties of cell-laden hydrogels increased over time

Compressive mechanical testing was performed to evaluate the progressive changes in the stiffness of the constructs over time (Fig. 5). After 28 d of chondrogenic culture, the compressive Young's

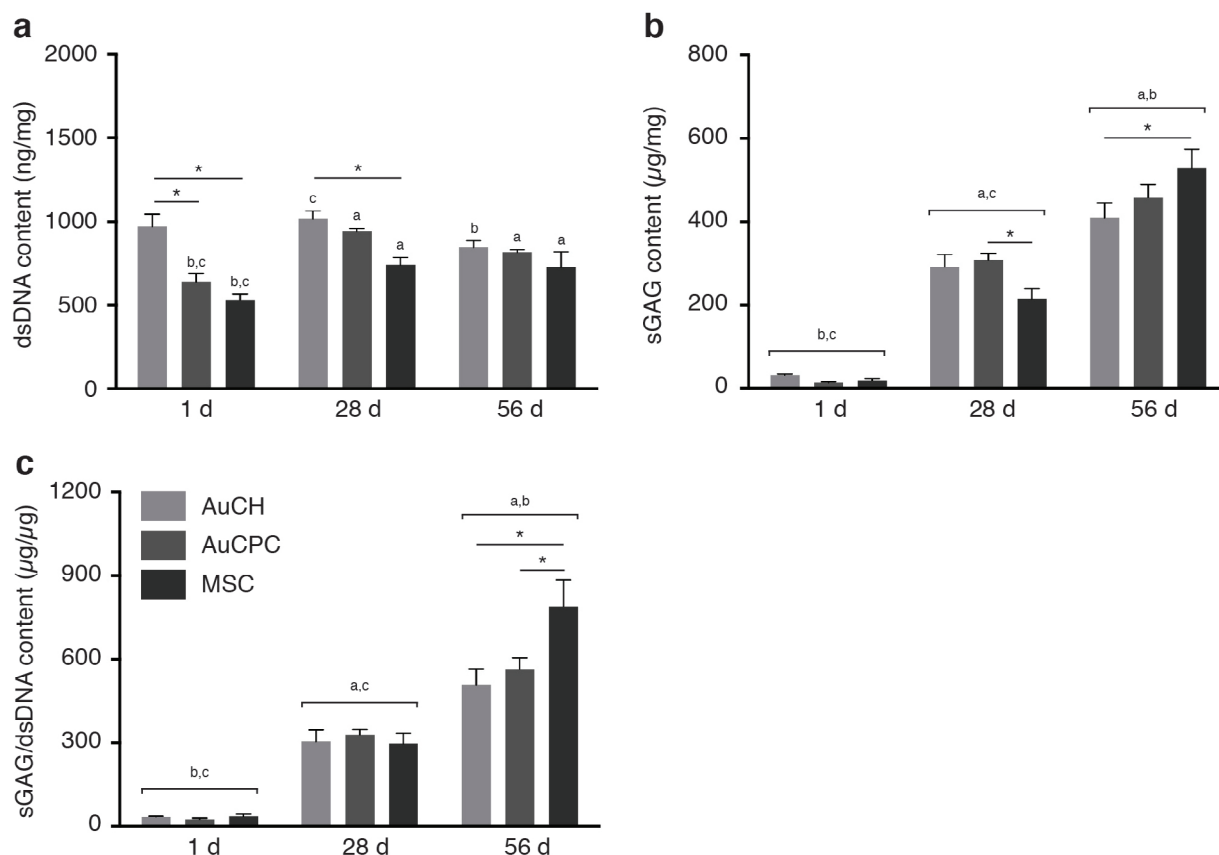


Fig. 4. Quantification of (a) sulfated GAG content and (b) dsDNA content in cell-laden hydrogels after 28 and 56 d of chondrogenic culture, both normalized against dry weight. The sGAG per dsDNA ratio is shown in c. Statistically significant differences of $p < 0.05$ for comparisons between cell types are marked with *. For each cell type, statistically significant differences of $p < 0.05$ are marked with ^a for a significant difference to day 1 and ^b for a significant difference to day 56.

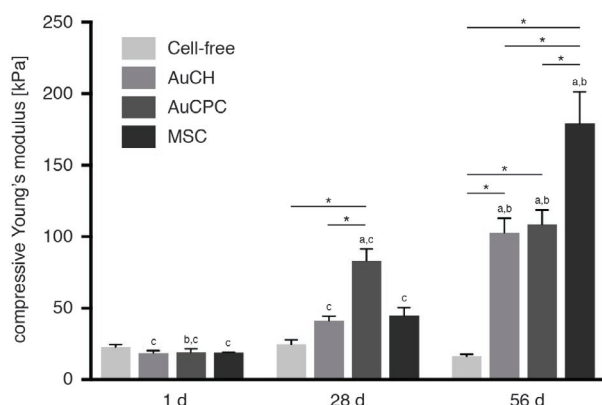


Fig. 5. Compressive Young's modulus of cell-laden hydrogels after 28 and 56 d of culture as compared to cell-free samples. Statistically significant differences of $p < 0.05$ for comparisons between cell types are marked with *. For each cell type, statistically significant differences of $p < 0.05$ are marked with ^a for a significant difference to day 1 and ^b for a significant difference to day 56.

modulus of AuCHs (41.3 ± 3.0 kPa) and MSCs (44.9 ± 5.6 kPa) increased non-significantly of 1.6-fold and 1.8-fold, respectively, in comparison to cell-free samples. Both cell types were outperformed by AuCPCs (83.2 ± 8.1 kPa), which exhibited a significant 3.3-fold increase. When comparing these results to the total sGAG contents per sample, a 1.4-fold difference between AuCPCs and MSCs was noted; however, the observed difference in mechanical properties between AuCPCs and AuCHs was not reflected in total sGAG content.

At the end of the culture period, MSCs (179.2 ± 22.2 kPa) significantly outperformed both AuCHs (102.8 ± 10.2 kPa) and AuCPCs (108.6 ± 10.1 kPa) in terms of compressive Young's modulus (1.7-fold and 1.6-fold difference, respectively), correlating with the observed trends in total sGAG and sGAG/dsDNA content. In comparison with cell-free hydrogels, cell-laden constructs demonstrated a significant increase in the compressive modulus of 10.5-fold for MSCs, 6.4-fold for AuCPCs and 6.1-fold for AuCHs.

The compressive Young's modulus of cell-free constructs remained in the range of 16–25 kPa at all time points, indicating no notable degradation of the bulk properties of the hydrogel during the period of *in vitro* culture. In general, cartilage matrix synthesis in cell-laden hydrogels, as indicated by the total sGAG content, correlated with an increase in mechanical properties of the samples.

Histology and immunohistochemistry confirm matrix deposition

Histological sections displayed the presence and distribution of several main components of cartilage extracellular matrix in the hydrogel, including proteoglycans as well as collagen type II, I and VI (Fig. 6).

AuCHs, AuCPCs and MSCs showed inhomogeneous distribution of synthesized

proteoglycans throughout the hydrogel constructs (Fig. 6a–c), with a gradation of decreasing labeling from the pericellular territorial to inter-territorial matrices. In all cases, there was an evident increase in the intensity of safranin O staining for proteoglycans over time. In line with the trends in total sGAG content, MSCs qualitatively displayed less proteoglycans at 28 d of culture, yet exhibited the most intense staining at the end of the culture period in comparison to AuCPCs and to a greater extent to AuCHs. In all experimental groups, the deposition of collagen type II and type I seemed to occur predominantly in the outer rim of the cylindrical hydrogel constructs, with a lighter staining in the center. The diameter of this outer rim appeared to widen over time, indicating increased matrix deposition towards the center of the construct. MSCs and AuCPCs exhibited a more homogeneous distribution of collagen type II, whereas AuCHs displayed clusters of intense pericellular labeling (Fig. 6d–f). A similar trend was noted for collagen type I (Fig. 6g–i), where rims with the highest intensity staining were found in AuCH samples, followed by MSCs and AuCPCs. These observations correlated with *COL1A1* mRNA expression profiles at day 56. Collagen type VI-labelled clusters (Fig. 6j–l) were observed pericellularly after 56 d of culture in AuCPCs and to a lesser extent in MSCs, with only sporadic staining in AuCHs. Finally, von Kossa staining for mineralization did not show any black staining indicative of calcium in either group (Fig. 6m–o).

Discussion

Cell selection for the generation of clinically relevant-size cartilage tissue constructs remains a notable challenge in tissue engineering strategies. Cartilage-derived progenitor cells represent a promising cell source for use in tissue engineering-based clinical therapies, since they can generate large numbers of cells while maintaining chondrogenic differentiation potential (Williams *et al.*, 2010). In this study, progenitor cells originating from auricular cartilage demonstrated cartilage formation capacity in a 3D hydrogel system by generating a cartilage-like matrix *in vitro*.

Traditionally, chondrocytes from various cartilage tissues (auricular, articular, costal and nasoseptal) are predominantly used for the engineering of auricular structures. Autologous cells have the greatest clinical potential in view of infectious considerations and adverse immunological response (Saadeh *et al.*, 1999). Nevertheless, the acquisition of a sufficient number of cells remains an important limitation in current tissue engineering approaches, since extensive expansion is required to obtain sufficient numbers of autologous cells for the creation of a large tissue constructs (Bichara *et al.*, 2012). After only few population doublings *in vitro*, auricular, articular, septal and costal chondrocytes undergo

dedifferentiation and lose their capacity to produce cartilage-specific glycosaminoglycans and collagens (Chung *et al.*, 2006; Schulze-Tanzil *et al.*, 2004; Schnabel *et al.*, 2002; Homicz *et al.*, 2002; Lee *et al.*, 2007; Nabzdyk *et al.*, 2009; Williams *et al.*, 2010). Although some studies demonstrate cartilage-like tissue formation using extensively expanded chondrocytes, these dedifferentiated cells required the addition of fresh chondrocytes in order to salvage some of the chondrogenic phenotype characteristics (Pomerantseva *et al.*, 2016; Tseng *et al.*, 2014). Hence, primary chondrocytes seem less suitable as a single donor source for cartilage tissue engineering of large constructs, yet they may be beneficial additions to co-cultures (Kang *et al.*, 2012; Lv *et al.*, 2012; Pleumeekers *et al.*, 2015).

MSCs are also successfully applied for the generation of cartilage-like tissue constructs (Cheng *et al.*, 2014; Pleumeekers *et al.*, 2013; Seda Tigli *et al.*, 2009; Vinardell *et al.*, 2012). MSCs are attractive for cartilage tissue engineering as they are harvested from the bone marrow with minimally invasive procedures, can be expanded to yield high cell numbers without losing their tissue-specific phenotype (Gardner *et al.*, 2013) and have multipotent differentiation ability (Pittenger *et al.*, 1999). In 3D culture, MSCs can be directed towards the chondrogenic lineage and are consequently applied in numerous studies

for the generation of cartilage (Cheng *et al.*, 2014; Pleumeekers *et al.*, 2013; Seda Tigli *et al.*, 2009; Vinardell *et al.*, 2012). In fact, their usage is already being explored in clinical trials such as NCT02037204 (IMPACT), NCT00885729 and NCT01227694 (as registered on clinicaltrials.gov). Nevertheless, the usage of MSCs for cartilage engineering harbors the risk of terminal differentiation of cells and subsequent calcification and ossification of tissues (Williams *et al.*, 2010). This results in calcification of the extracellular matrix (Gawlitta *et al.*, 2010) – a phenomenon that is also observed in the costal cartilage framework implanted during auricular reconstruction surgery – causing an increasingly rigid construct (Jessop *et al.*, 2016). This is an unfavorable outcome for engineered auricular tissue structures as elasticity is one of the key features of the external ear (Nimeskern *et al.*, 2015b; Pappa *et al.*, 2013; Xu *et al.*, 2005).

Tissue-specific progenitor cells maintain stem-cell-like proliferative potential, yet also display tissue-specific phenotypes since they are harvested directly from the target tissue. *In vivo*, environmental influences from their niche prime progenitors to regulate proliferation and differentiation towards the target tissue, providing a clear advantage over non-tissue-specific stem-cells-like MSCs. Their proliferative ability addresses the important limitation in cell number acquisition that continues

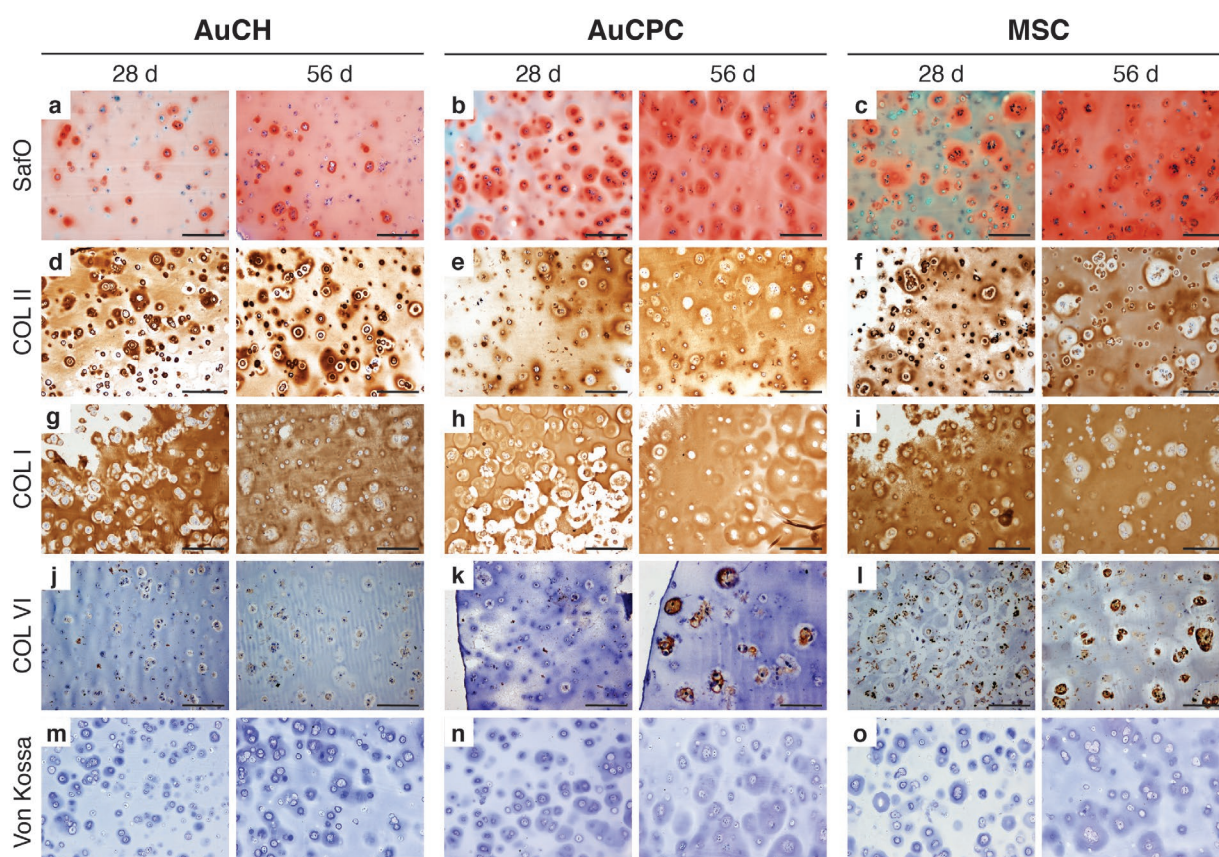


Fig. 6. Histological analysis of cell-laden hydrogels after 28 and 56 d in chondrogenic culture. Safranin O staining visualizing proteoglycan deposition in (a) AuCH, (b) AuCPC (c) and MSC samples, as well as immunohistochemistry for (d,e,f) collagen type II, (g,h,i) collagen type I and (j,k,l) collagen type VI. (m,n,o) Von Kossa staining demonstrating the absence of mineralization in all three cell types. Scale bars: 150 μ m.

to hamper the translation of large tissue-engineered constructs to clinical application. Cartilage stem/progenitor cells were first identified in the superficial zone of the articular cartilage of the knee (Dowthwaite *et al.*, 2004); yet, conclusive definitions on the identity of these cells remain elusive (Jiang and Tuan, 2015). Nevertheless, it is clear that cartilage tissue harbors a potent subpopulation of cells with distinct abilities from primary chondrocytes and a similar nature to MSCs. The current work presented the first identification and evaluation of this subpopulation of progenitor cells in equine auricular cartilage tissue and their potential for cartilage tissue engineering approaches.

Previous literature reporting progenitor cells from auricular tissue is principally focused on cells originating from the perichondrial layer (Kobayashi *et al.*, 2011a; Kobayashi *et al.*, 2011b; Togo *et al.*, 2006), which is a fibrous connective tissue consisting of fibroblasts and perichondrocytes. Recently, the first demonstration of the presence of an auricular cartilage progenitor population separated from the perichondrium has been presented by Xue *et al.* (2016) in a porcine species. The proliferation rate of these auricular cartilage stem/progenitor cells (CSPCs) is comparable to bone marrow-derived MSCs (Xue *et al.*, 2016). Although the authors report a higher proliferation rate for perichondrium stem/progenitor cells (PSPCs), chondrogenic differentiation potential is greater for auricular cartilage stem/progenitor cells (Xue *et al.*, 2016). PSPCs can differentiate into chondrocytes, yet maintaining a fibroblastic morphology (Kobayashi *et al.*, 2011a; Xue *et al.*, 2016). Cells derived from the cartilage tissue appear more apt to differentiate towards the chondrogenic and osteogenic lineages, whereas perichondrium-derived progenitor cells are inclined towards adipogenic differentiation (Xue *et al.*, 2016). The cartilage-forming potential of cartilage stem/progenitor cells has not yet been explored in 3D biomaterial culture for tissue engineering purposes. Hence, the present study focused on the use of cartilage progenitor cells, derived from the cartilage after complete removal of the perichondrium, for elastic cartilage tissue engineering. Besides fundamental characterization of these putative cartilage progenitor cells, their potential for cartilage regeneration for future therapeutic applications is of major interest.

The results from the characterization of AuCPCs demonstrated that they had a behavior similar to MSCs in terms of multipotency. Similar to MSCs, AuCPCs had the ability to differentiate towards multiple lineages, as confirmed by positive stainings for bone, adipose tissue and cartilage. Furthermore, AuCPCs, being $CD73^+$, $CD90^+$, $CD105^+$, $CD34^-$ and $CD45^-$, displayed a gene expression profile for surface markers that is consistent with the minimal criteria for human MSCs (Dominici *et al.*, 2006). Additional cell surface marker analysis showed similar gene expression profiles between AuCPCs and bone marrow-derived MSCs. A main difference among

AuCHs, AuCPCs and MSCs arose in the expression of $CD13$, which is a marker widely expressed by a variety of cell subpopulations, including stem cells (Calloni *et al.*, 2013; Dondossola *et al.*, 2013). These data, in addition to the ability to differentiate towards multiple lineages, demonstrated a behavior similar to bone marrow-derived MSCs, both genetically and functionally.

The limited knowledge on AuCPCs demands further investigation into specific cell characteristics, yet our study focused on advancing towards the utilization of these cells. Their stem cell characteristics, in addition to a differentiation capacity likely primed towards their source tissue, makes AuCPCs a highly interesting cell source for cartilage tissue engineering strategies. These cells provide an opportunity to overcome the drawbacks of the cell types currently used in cartilage tissue engineering and, thereby, can increase the likelihood of using tissue-engineered auricular cartilage structures for clinical application. Hence, the behavior of AuCPCs in 3D culture for tissue regeneration was explored in comparison to AuCHs and MSCs. Growth and differentiation in the 3D provides a more natural environment for cells and allows the preservation of tissue-specific characteristics (Pampaloni *et al.*, 2007). In this study, cells were encapsulated in a 3D hydrogel system (gelMA) proven to be a permissive environment for neocartilage production (Klotz *et al.*, 2016; Levato *et al.*, 2017; Levett *et al.*, 2013; Schuurman *et al.*, 2013). GelMA is a widespread platform for tissue engineering and bioprinting applications, owing to its natural bioactivity and tailorability (Klotz *et al.*, 2016). This versatile hydrogel can be functionalized for improved performance in supporting cells [for instance with hyaluronic acid (Levett *et al.*, 2013)], printing resolution [e.g. with gellan gum (Mouser *et al.*, 2016)] or mechanical stability [by incorporating reinforcing fibers (Visser *et al.*, 2015), scaffolding materials, such as stiff and elastic hydrogels (Melchels and Blokzijl *et al.*, 2016), or thermoplastic polymers (Mouser *et al.*, 2017)]. Its biocompatibility and clinical grade options make gelMA a beneficial biomaterial choice with respect to future clinical translation (Klotz *et al.*, 2016).

In 3D hydrogel culture, all three cell types demonstrated increasing extracellular matrix production over the culture period, as indicated by sGAG/dsDNA quantification and confirmed by histology. Whereas AuCHs and AuCPCs displayed a similar matrix-synthesizing behavior, MSCs outperformed both cell types in terms of sGAG deposition. Proteoglycan aggregates are the major structural matrix components contributing to the mechanical properties of the tissue and the trend in matrix synthesis clearly matched the compressive moduli among cell types at the end of the culture period. Nevertheless, the observed differences in total sGAG content at 28 d of culture did not reflect the concurrent mechanical properties. At this time point, AuCPCs were outperforming MSCs in terms of total

sGAG content and compressive Young's modulus, but the marked difference in Young's modulus between AuCPCs and AuCHs was not observed in the total sGAG content. Since the mechanical properties of cartilage tissue are impacted by the organization of the tissue (Wu and Herzog, 2002), a potential explanation is a more homogeneous distribution of proteoglycans and collagens in both AuCPCs and MSCs upon histological examination, whereas AuCHs displayed a more cluster-like organization of matrix components. In our study, the mechanical properties of tissue-engineered cartilage constructs ranged between 102.8 and 179.2 kPa, which was at least a factor 10 inferior to native auricular cartilage. Griffin *et al.* (2016) report the compressive moduli of native human auricular cartilage to range between 1.41 and 2.08 MPa, whereas Nimeskern *et al.* (2015a) determine an instantaneous modulus of 3.27–11.02 MPa, a maximum stress of 0.87–3.11 MPa and an equilibrium modulus of 2.22–7.23 MPa. Thus, tissue-engineered cartilage constructs require improved mechanical properties, which can be attained through supporting frameworks (Cervantes *et al.*, 2013), by fiber reinforcement (Visser *et al.*, 2015) or possibly through mechanical loading of the developing neo-tissue, which is shown to increase cellular production of matrix components in articular cartilage (Musumeci, 2016).

Differential mRNA expression of cartilage-specific markers confirmed cell differentiation towards the chondrogenic lineage in all three cell types. Although collagen type II expression was highly upregulated over the 56-d culture period, all three cell types demonstrated concurrent collagen type I expression, confirmed by immunohistochemistry. Improvement of neocartilage quality may be attained through functionalization of the hydrogel or by optimizing the recipe of the differential culture media. For example, the incorporation of hyaluronic acid in gelMA is shown to decrease collagen type I production (Levett *et al.*, 2013), whereas supplementation with insulin-like growth factor-1 (IGF-1) enhances the generation of elastic fibers in addition to improved overall tissue formation (Rosa *et al.*, 2014).

The auricle is a strong, yet flexible structure composed of cartilage of the elastic type. Its composition is similar to that of hyaline cartilage, consisting of negatively charged proteoglycan aggregates attracting water and a dense collagen type II network (Nabzyk *et al.*, 2009). Nevertheless, the auricular cartilage is unique in that it harbors an intricate network of elastic fibers. Elastin has a defining role in the mechanical properties of elastic cartilage, allowing flexibility and a swift return to its original shape after minor loads (Nimeskern *et al.*, 2015b). AuCPCs exhibited an upregulated expression of elastin in the cell-laden hydrogels. This feature offers the potential prime advantage of using AuCPCs over MSCs for tissue engineering of auricular cartilage, since the latter did not display elastin

upregulation. Quantification and/or visualization of the production of elastin in tissue-engineered constructs could confirm this potential and would be advisable for future studies.

One common problem with cartilage tissue engineering is calcification of the neo-tissue (Jessop *et al.*, 2016). Primary chondrocytes may terminally differentiate and become hypertrophic, which can lead to calcification and eventually ossification of the neo-tissue (Gerstenfeld and Shapiro, 1996; Phull *et al.*, 2016). Collagen type X is a typical marker of chondrocyte hypertrophy (Martin *et al.*, 2001), although its presence is demonstrated in native non-mineralized auricular cartilage (Dahl *et al.*, 2011; Hellingman *et al.*, 2011). Several studies report the expression of collagen type X in cartilage tissue engineered from auricular chondrocytes, yet this does not result in mineralization of neo-tissues in both *in vitro* and *in vivo* conditions (Dahl *et al.*, 2011; Hellingman *et al.*, 2011). Indeed, in this study, the mRNA expression of collagen type X was highly upregulated in AuCHs, indicating chondrocyte hypertrophy. Nonetheless, our findings confirmed that this did not result in mineralization of the neo-tissue up to 56 d of *in vitro* culture. The relatively low expression of *COL10A1* in both AuCPCs and MSCs might indicate preservation of the phenotype in these cell types under *in vitro* chondrogenic conditions. Nevertheless, the expression of *RUNX2* was significantly upregulated in MSCs as compared to AuCPCs and AuCHs at 56 d of culture. *RUNX2* drives osteogenic differentiation and inhibits differentiation of MSCs into chondrocytes (Komori, 2006). In contrast, AuCPCs demonstrated a significant reduction in *RUNX2* expression over time. MSCs from the bone marrow form bone *in vivo* through the endochondral ossification pathway (Gawlitza *et al.*, 2010), whereas AuCPCs originate from the cartilage itself and their niche may have primed them towards the target tissue, maintaining their specific phenotype. The lack of hypertrophy in AuCPC cultures, as well as the decreased *RUNX2* and increased elastin expression levels pointed to a stable phenotype that is amenable for tissue engineering.

In conclusion, the identification of a cartilage progenitor subpopulation in the auricular cartilage provided access to a promising cell source for tissue engineering strategies for auricular reconstruction. Although under the current culturing conditions, bone marrow-derived MSCs seemed to perform better in terms of matrix production, major advantages of AuCPCs include the ability to generate high cell numbers (Williams *et al.*, 2010; Xue *et al.*, 2016), upregulation of the elastin gene and a limited endochondral ossification potential. Taken together, these advantages make progenitor cells from the auricular cartilage a highly interesting candidate as a cell source for future tissue-engineering-based clinical therapies.

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Discussion with Reviewers

Oliver Gardner: Articular cartilage progenitor cells have a highly specific location within the tissue, which relates to their role in the growth and maintenance of the tissue. Do the authors believe that AuCPCs have a specific niche and role within auricular cartilage?

Authors: For auricular cartilage progenitor cells there is currently no proof of a specific localization, also due

to the limited amount of studies focusing on these cells. Since there is a lack of specific markers for these cells, such a localization cannot yet be easily defined. As it may be possible that AuCPCs have a specific niche within the auricular cartilage, future studies on the developmental origin, exact tissue localization and role of AuCPCs would be very interesting for the field of cartilage biology.

Karoliina Pelttari: Can a big enough biopsy be harvested from patients without donor site morbidity? What about patients with bilateral microtia?

Authors: Cartilage progenitor cells are thought to form up to $\pm 0.7\%$ of the cell population in the articular cartilage and are reported to be able to proliferate to over 60 population doublings (Williams *et al.*, 2010). Similar characteristics in progenitors from other cartilage sources are expected. AuCPCs demonstrated a potent ability to proliferate up to passage 4 – when, in the present study, the cells were utilized for tissue engineering purposes – meaning that large amounts of functional cells can be generated. Although exact calculations on the number of progenitor cells obtainable from a specific cartilage volume are outside the scope of this study, the proliferation potential of progenitor cells suggest that, with only a small biopsy from the auricular cartilage, a sufficient number of cells can be generated for the production of a full-sized tissue-engineered auricle. Cells can be obtained from healthy auricular cartilage or from the microtia remnant, although the presence of progenitor cells in the latter tissue has not yet been reported. Depending on the proliferative and cartilage-forming ability of microtia-derived cells, this source can potentially overcome donor site morbidity in both unilateral and bilateral microtia patients.

Karoliina Pelttari: Are cells from patients with congenital microtia different from healthy auricular chondrocyte? Gu *et al.* (2017) (additional reference) show an inferior cartilage-forming capacity of microtia-chondrocytes.

Authors: The microtic auricle is a congenital deformity. Its etiology is not fully understood, yet there is evidence for the involvement of environmental factors besides genetic factors (Luquetti *et al.*, 2012, additional reference). On a macroscopic level, the microtic cartilage is reported to possess similar cell morphology and tissue organization (Melgarejo-Ramirez *et al.*, 2016, additional reference). Various studies observed no marked differences between healthy and microtia chondrocytes in terms of growth rates, expression of cartilage genes and ability to generate cartilage-like tissue *in vivo* (Kamil *et al.*, 2004; Ishak *et al.*, 2011, additional reference; Ishak *et al.*, 2015; Nakao *et al.*, 2017). Indeed, Gu *et al.* (2017) report a biochemically and biomechanically inferior quality neocartilage produced by microtia chondrocytes. These contrasting results indicate the

need for a further exploration of differences between microtia and healthy chondrocytes and their ability for cartilage production under various conditions.

Karoliina Pelttari: Have such AuCPC with similar characteristics been identified in human?

Authors: Cartilage progenitor cells are a relatively established cell population in the articular cartilage of various animal species (Dowthwaite *et al.*, 2004; Williams *et al.*, 2010; McCarthy *et al.*, 2012; Tran *et al.*, 2014, additional reference), yet only little is known about the presence of these cells in the auricular cartilage. Cartilage-derived progenitor cells are identified in porcine and now equine auricular cartilage. Following this observation, the presence of such a subpopulation in the human auricular cartilage is likely.

Karoliina Pelttari: Is the hydrogel used in this study an already approved material for clinical use?

Authors: The hydrogel used in this study, gelatin methacryloyl (gelMA), is not yet approved for clinical use. Nevertheless, it demonstrates good biocompatibility when synthesized with the endotoxin-free gelatin type B (Sirova *et al.*, 2014, additional reference). Although the biocompatibility of gelMA and its degradation products requires more extensive analysis, the biomaterial has potential for clinical application. In this study, this hydrogel was chosen as it continuously proves to be a suitable 3D system for tissue engineering, as multiple research groups repeatedly demonstrate (amongst others: Yue *et al.*, 2015; Loessner *et al.*, 2016; Klotz *et al.*, 2016; Brown *et al.*, 2017, additional references).

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