

Isolation and characterization of human articular chondrocytes from surgical waste after total knee arthroplasty (TKA)

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Cartilage tissue engineering is a fast evolving field of biomedical engineering, in which the chondrocytes represent very commonly used cell type in related studies. Since research in tissue engineering always consumes a lot of cells, an efficient and cheap isolation method could form a powerful basis to boost such studies and enable their faster progress to the clinics. Isolated chondrocytes can be used for autologous chondrocyte implantation in cartilage repair, and are the base for valuable models to investigate cartilage phenotype preservation, as well as enable studies of molecular features, nature and scales of cellular responses to alterations in the cartilage tissue. In this study, we developed a protocol for isolation and consequent cultivation of primary human adult articular chondrocytes with the desired phenotype from the surgical waste obtained during total knee arthroplasty (TKA). TKA is a common and very frequently performed orthopaedic surgery during which often both femoral condyles are removed. The latter present the ideal source for an efficient and relatively cheap isolation of chondrocytes. Cartilage specific gene expression of collagen type 2 and aggrecan have shown that the proposed protocol enables the chondrocyte specific phenotype preservation up to the second passage. Collagen type 1 expression was performed to evaluate the dedifferentiation of the isolated cells into a fibroblast-like phenotype.

1 **Isolation and characterization of human articular chondrocytes from surgical waste after**
2 **total knee arthroplasty (TKA)**

3 Short title: Alternative human chondrocyte isolation

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33 **AUTHOR CONTRIBUTIONS**

34 J.N., L.G., M.V. and U.M. designed the study. J.N., L.G. and M.G. performed the experiments.
35 J.N., L.G. and U.M. wrote the manuscript and prepared the figures. All authors reviewed the
36 manuscript.

37 **ABSTRACT**

38 Cartilage tissue engineering is a fast evolving field of biomedical engineering, in which the
39 chondrocytes represent very commonly used cell type in related studies. Since research in tissue
40 engineering always consumes a lot of cells, an efficient and cheap isolation method could form a
41 powerful basis to boost such studies and enable their faster progress to the clinics. Isolated
42 chondrocytes can be used for autologous chondrocyte implantation in cartilage repair, and are the
43 base for valuable models to investigate cartilage phenotype preservation, as well as enable studies
44 of molecular features, nature and scales of cellular responses to alterations in the cartilage tissue.

45 In this study, we developed a protocol for isolation and consequent cultivation of primary human
46 adult articular chondrocytes with the desired phenotype from the surgical waste obtained during
47 total knee arthroplasty (TKA). TKA is a common and very frequently performed orthopaedic
48 surgery during which often both femoral condyles are removed. The latter present the ideal source
49 for an efficient and relatively cheap isolation of chondrocytes. Cartilage specific gene expression
50 of collagen type 2 and aggrecan have shown that the proposed protocol enables the chondrocyte
51 specific phenotype preservation up to the second passage. Collagen type 1 expression was
52 performed to evaluate the dedifferentiation of the isolated cells into a fibroblast-like phenotype.

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54 **KEYWORDS:** human articular chondrocytes, isolation protocol, total knee arthroplasty, TKA,
55 phenotype preservation, gene expression, collagen 2, aggrecan

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58 1 INTRODUCTION

59 Damage to articular cartilage has important clinical implications since the cartilage tissue
60 possesses a limited intrinsic healing potential and tends to an incomplete regeneration by local
61 chondrocytes, accompanied with an inferior fibrocartilage formation (Camp et al. 2014; McNickle
62 et al. 2008; Richter et al. 2016). Surgical intervention is often the only option, although the repair
63 of damaged cartilage is often less than satisfactory, and rarely restores full function or returns the
64 tissue to its native state (Kerker et al. 2008; Kock et al. 2012; Tuli et al. 2003). Over the past
65 decade a number of viable options of cartilage regeneration have been introduced into clinical
66 practice (Camarero-Espinosa et al. 2016; Hettrich et al. 2008; Schrobback et al. 2011). Among
67 these, autologous chondrocyte implantation (ACI) seems the most promising since it relies on the
68 use of biodegradable materials that serve as temporary cell-carriers, enabling *in vitro* cell growth
69 and subsequent implantation into the defective cartilage (Bomer et al. 2016; Niemeyer et al. 2016;
70 Robb et al. 2012).

71 Tissue engineering of articular cartilage remains challenging due to the specific structure of
72 cartilage tissue, i.e. its multiphasic cellular architecture together with a remarkable weight-bearing
73 characteristics (e.g. resistance to mechanical stress and wear) (Kim et al. 2012; Su et al. 2012).
74 Good understanding of the cartilage structure, physiology, and the molecular basis of
75 chondrogenesis is key to *in vitro* cartilage production, either for use in tissue engineering or clinics
76 (Bhat et al. 2011; Lee et al. 2013; Li et al. 2012). The state-of-the-art concept of *in vitro* cartilage
77 tissue development combines the use of biocompatible and biodegradable carrier materials, the
78 application of growth factors, the use of different cell types (stem or already differentiated) and
79 different approaches to simulate the native mechanical stimulation (Gardner et al. 2013; Hildner
80 et al. 2011; Khan et al. 2013; Naranda et al. 2016).

81 More specific challenges of articular cartilage tissue engineering remain the high consumption of
82 cells and related costs, as well as the preparation of an ideal host scaffold. Although solutions to
83 both mentioned challenges have been introduced in recent years (Bassleer et al. 1998; Stellavato
84 et al. 2016), is the cell part gaining far less research momentum. Therefore, it comes to no surprise
85 that novel approaches for chondrocyte isolation are highly desired, especially considering the high
86 prices of ordered cells. Optimisation of isolation yields, abundant cell sources and efficient
87 culturing procedures that lead to preparation of desired, reproducible and relatively affordable cell
88 cultures or/and material-cell constructs with good durability, are therefore highly rated novelties
89 in recent research (Dehne et al. 2009; Naranda et al. 2016; Otero et al. 2012).

90 The most demanding part in the process of *in vitro* culturing still presents the preservation of the
91 desired phenotype to a high enough passage to yield sufficient cells to perform planned
92 experiments (Pei & He 2012; Rosenzweig et al. 2012; Schnabel et al. 2002a; Wu et al. 2014). Since
93 the latter depends on numerous factors and can therefore be confirmed only by a combination of
94 (often) expensive techniques (different microscopies, molecular analysis, immunohistochemistry
95 etc.), it is important to prepare protocols for an easier and cheaper preliminary phenotype
96 confirmation by means of methods, available in most cell laboratories around the world. Since the
97 desired phenotype can be identified by chondrocyte specific production (Han et al. 2010), we
98 believe that the easiest and safest preliminary method to prove phenotype preservation could be

99 the analysis of gene expression. More specifically, this analysis should include the evaluation
100 expression of genes related to cartilage specific markers (e.g. collagen type 2 and aggrecan). To
101 follow-up possible dedifferentiation towards the fibroblastic phenotype (Goldring et al. 2006;
102 Makris et al. 2015; Otero et al. 2012), we propose simultaneous measurement of up-regulation of
103 collagen type 1.

104 Based on all mentioned it is clear that chondrocyte isolation from an abundant source with a high
105 yield, together with an effective and cheap preliminary phenotype confirmation method, would be
106 greatly beneficial to boost the cartilage tissue engineering (Cetinkaya et al. 2011; Goepfert et al.
107 2010; Schrobback et al. 2011). This study was therefore designed to provide a relatively simple,
108 yet effective procedure for isolation and culturing of human tissue derived primary chondrocytes
109 up to the second passage. As the preliminary method of phenotype confirmation we chose the
110 evaluation of chondrocyte specific gene expression, together with morphological evaluation of
111 cells. Such an approach provides a cheap and effective protocol to be considered an alternative to
112 other available methods (Hu et al. 2002; Li et al. 2015; Strzelczyk et al. 2001; Xu & Zhang 2014),
113 especially suitable for other laboratories to boost their respective entry level cartilage tissue
114 engineering studies.

115 **2 MATERIALS AND METHODS**

116 **2.1 Materials**

117 All used materials and chemicals were of laboratory grade and purchased from Sigma-Aldrich,
118 Germany, if not stated otherwise. For specific parts of the isolation process and cultivation, all
119 used labware and chemicals were additionally sterilized using the standard autoclavation
120 procedure (Avtoklav A-21, Kambič, Slovenia).

121 **2.2 Isolation of primary chondrocytes**

122 Full-thickness cartilage was surgically removed from the femoral condyle of arthritic knee of a 50
123 years old patient who underwent total knee arthroplasty (TKA) performed at the University
124 Medical Centre Maribor, Slovenia (application reference: 123/05/14). Prior to surgery, no systemic
125 disease or any treatment was reported for the donor patient. The study was conducted in accordance
126 with the *Declaration of Helsinki* and its subsequent amendments and was approved by the
127 Republic of Slovenia National Medical Ethics Committee (Ljubljana, Slovenia). The patients'
128 informed consent was obtained.

129 The cartilage tissue was surgically removed under sterile conditions, transferred into a previously
130 sterilized 250 ml glass bottle filled with phosphate buffered saline (PBS, Sigma-Aldrich,
131 Germany) and immediately transferred to the cell isolation laboratory. The cartilage tissue was
132 transferred to a petri dish filled with PBS to prevent drying of the tissue and subsequently cut into
133 pieces of approximately 2 x 2 mm. PBS was carefully removed by a pipette and the petri dish was
134 immediately filled with 10 mL solution of 0.25 wt.% Trypsin/EDTA (Sigma, France). The as-
135 prepared cartilage pieces were incubated for 3 hours at 37°C and 5 wt.% CO₂ (CO₂ Incubator
136 MCO-19AICUVH-PE, Panasonic, Japan), followed by addition of 20 mL of Advanced Dulbecco's
137 modified Eagle's medium (Advanced DMEM, Gibco, Grand Island, NY, USA) to the cell

138 suspension. The suspension was transferred to a 50 mL falcon tube and centrifuged at 300 x g for
 139 10 minutes (Centrifuge 5804 R, Eppendorf, USA). The supernatant was carefully discarded and
 140 the cell pellet was re-suspended in 20 mL of Advanced DMEM and centrifuged at 200 x g for 5
 141 minutes (Centrifuge 5804 R, Eppendorf, USA). The supernatant was again carefully discarded and
 142 the cell pellet re-suspended in 10 mL Advanced DMEM supplemented with 100 IU/ml Penicillin,
 143 1 mg/ml Streptomycin, 2mM L-glutamine and 5 wt.% foetal bovine serum (FBS, Gibco, Grand
 144 Island, NY, USA) and plated on 25 cm² flasks (in triplicates). The primary chondrocytes were then
 145 left until confluence was reached. Growing cells were regularly observed with an Axiovert 40
 146 inverted optical microscope (Zeiss, Germany) at several magnifications. The culturing medium
 147 was changed every three days. The general steps of the procedure are schematically depicted in
 148 **Fig. 1**.

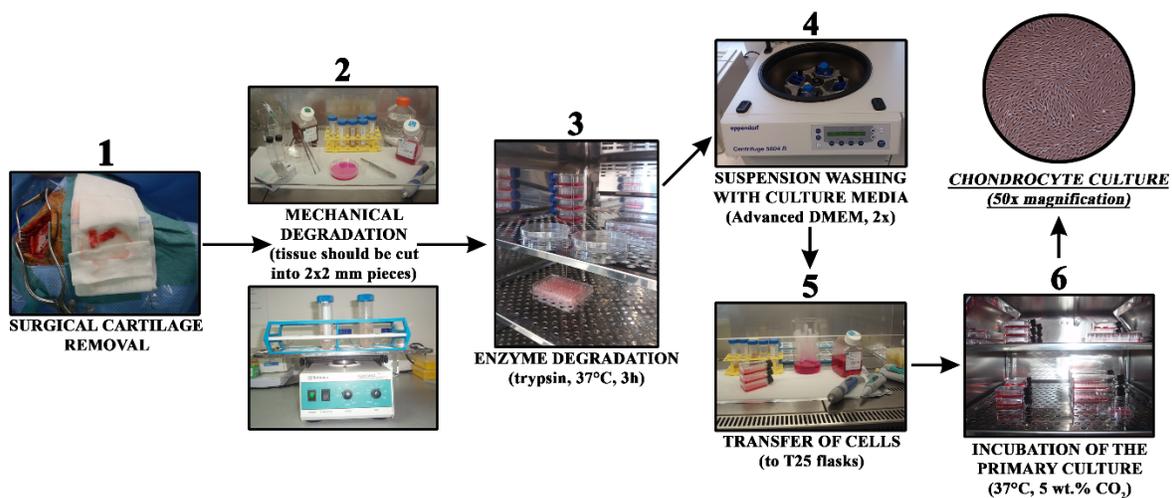


Fig. 1: Chondrocyte isolation from cartilage in a short overview of the most important preparation steps.

149 2.3 Isolated primary chondrocyte cell culture characterization

150 Gene expression analysis of cartilage specific markers collagen type 2 (COL2) and aggrecan
 151 (ACAN) was performed in order to determine the primary chondrogenic phenotype. Possible
 152 dedifferentiation to a more fibroblast like cell type was evaluated by monitoring the expression of
 153 collagen type 1 (COL1). After confluence was reached in all respective samples (triplicates) (see
 154 above section 2.2. for details), the cell suspension was transferred to micro-centrifuge tubes, and
 155 1.4 mL of TRI reagent (Sigma-Aldrich, Germany) was added. The tubes were vortexed for 30 min
 156 at room temperature. Afterwards, 280 μ L of chloroform (Sigma-Aldrich, Germany) was added and
 157 the tubes were further vortexed for 15 min and centrifuged at 12.000 rpm and 4°C. RNA extraction
 158 was carried out according to the manufacturer's instructions (Chomczynski 1993; Louveau et al.
 159 1991). Concentration and purity of the extracted cellular RNA was determined using NanoDrop
 160 2000c (Thermo Scientific, Delaware, USA) through optical density readings at 260 nm and a
 161 260/280 nm ratio. cDNA was obtained by using a cDNA reverse transcription kit (Applied
 162 Biosystems, California, USA). Primer sequences for cartilage target genes ACAN and COL2 were
 163 obtained from Caterson et al. (Caterson et al. 2001), while the corresponding mRNA sequences
 164 were retrieved from PubMed Nucleotide database (www.ncbi.nlm.nih.gov/nucleotide/) and the

165 AceView database (Thierry-Mieg & Thierry-Mieg 2006). Primers for the target gene COL1 were
 166 designed using IDT oligo analyser (eu.idtdna.com/calc/analyser). The primer sequences with the
 167 corresponding mRNA sequences and the corresponding NCBI accession numbers, are given in
 168 **Table 1**. 2 μ L of each cDNA sample with concentration of 15 ng/ μ L was used for quantitative real
 169 time PCR (qPCR) analysis performed using LightCycler 480 thermocycler (Roche, Switzerland)
 170 and with 2 \times Maxima SYBR Green qPCR master mix (Life Technologies, California, USA)
 171 according to the manufacturer's instructions. The quality and specificity of PCR amplicons were
 172 checked using melting curve analyses and agarose gel electrophoresis (see below **Fig. 4**). All
 173 shown results are presented as average values with the standard errors.

174 **Table 1:** Primer sequences with the corresponding mRNA sequence and the corresponding
 175 NCBI accession numbers.

Gene	Gene name	Accession number	Primer sequence 5'→3'
<i>ACAN</i>	<i>Aggrecan</i>	NM_013227.3 NM_001135.3	TGAGGAGGGCTGGAACAAGTACC GGAGGTGGTAATTGCAGGGAACA
<i>COL1</i>	<i>Collagen type 1, alpha 1</i>	NM_000088.3	CGGCTCCTGCTCCTCTTAG CACACGTCTCGGTCATGGTA
<i>COL2</i>	<i>Collagen type 2, alpha 1</i>	NM_001844.4 NM_033150.2	TTTCCCAGGTCAAGATGGTC CTGCAGCACCTGTCTCACCA
<i>GAPDH</i>	<i>Glyceraldehyde-3-phosphate dehydrogenase</i>	NM_001289745.1 NM_002046.5 NM_001289746.1 NM_001256799.2	GGGCTGCTTTTAACTCTGGT TGGCAGGTTTTTCTAGACGG

176 3 RESULTS

177 3.1 Isolation of primary chondrocytes

178 As mentioned in the Materials and methods section, the full-thickness cartilage was obtained from
 179 the femoral condyle of an arthritic knee during knee arthroplasty (TKA) performed at the
 180 University Medical Centre Maribor, Slovenia. TKA is a common procedure at the mentioned
 181 hospital, considering that approximately 700 such surgeries are performed each year (Univerzitetni
 182 klinični center 2014). Since the removed cartilage tissue is considered surgical waste, this presents
 183 a reliable and continuous source for isolation of primary chondrocytes.

184 The primary chondrocytes were isolated as described in the Materials and methods section. During
 185 their cultivation, their morphology and proliferation were regularly observed using inverted optical
 186 microscopy (**Fig. 2**). **Fig. 2A** shows the thin slice of cartilage that was used for their cultivation,
 187 while **Fig. 2B-D** present the primary human chondrocytes in a monolayer culture at different
 188 cultivation times. This initial examination was performed to follow possible morphological
 189 changes in the cell shapes, which would indicate possible dedifferentiation.

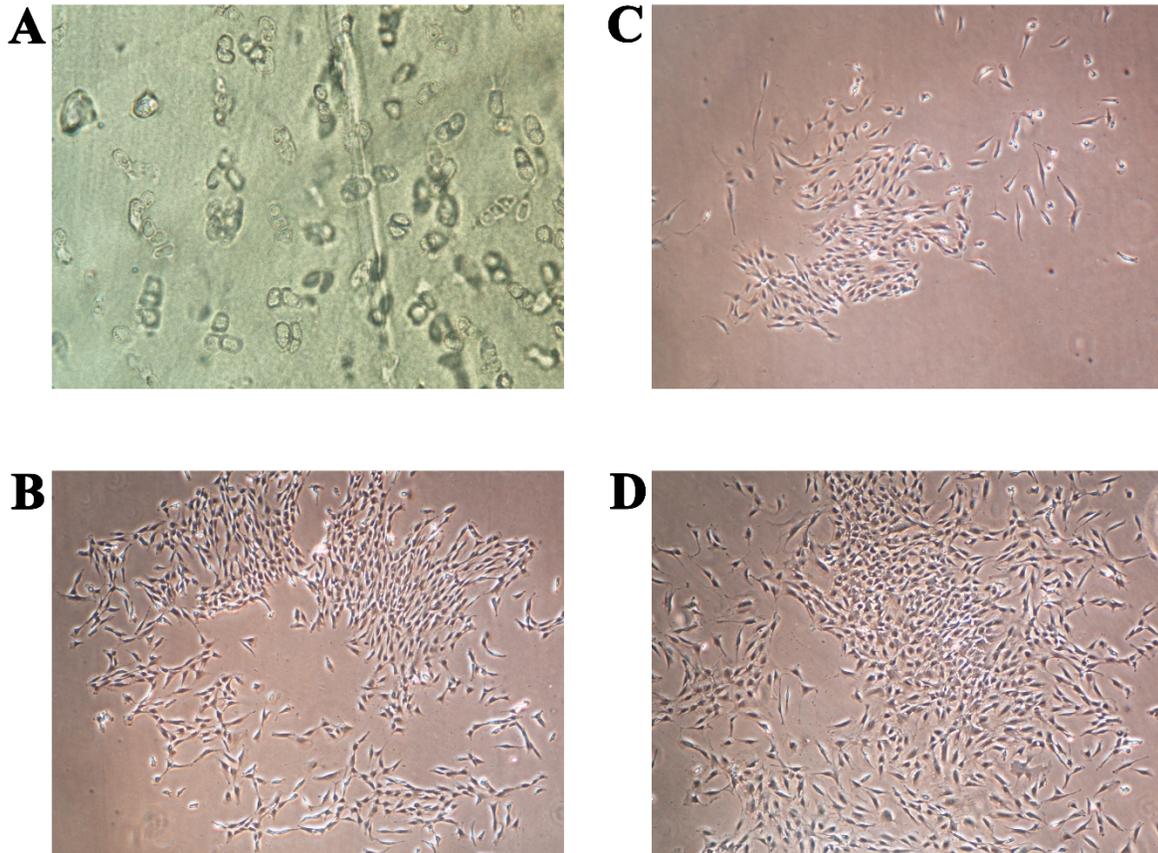


Fig. 2: A) Thin slice of cartilage for primary chondrocyte isolation; B-D) the primary human chondrocyte culture in a monolayer after 3, 6 and 9 days, respectively. The magnification of all shown images is 50x.

190 The full confluence of the isolated cells for the first and second passage was reached after two (14
191 days) and after one (7 days) week, respectively. Cell growth stopped presumably due to contact
192 inhibition (Lackie 2013). A comparison between the primary chondrocyte culture and the obtained
193 chondrocyte cultures after the first and second passages, is shown in **Fig. 3**. The cells formed
194 confluent monolayers (after the above mentioned cultivation times) and appeared polygonal in
195 shape (**Fig. 3A-C**). It can be observed that the chondrocyte morphology became more spindle-like
196 in the second passage (**Fig. 3C**), showing their tendency for dedifferentiation, most likely towards
197 fibroblast like cells (Hong & Reddi 2013). Observing the mentioned changes was an indication
198 that the third passage will not yield a high percentage of chondrocytes only, using the proposed
199 cultivation conditions.

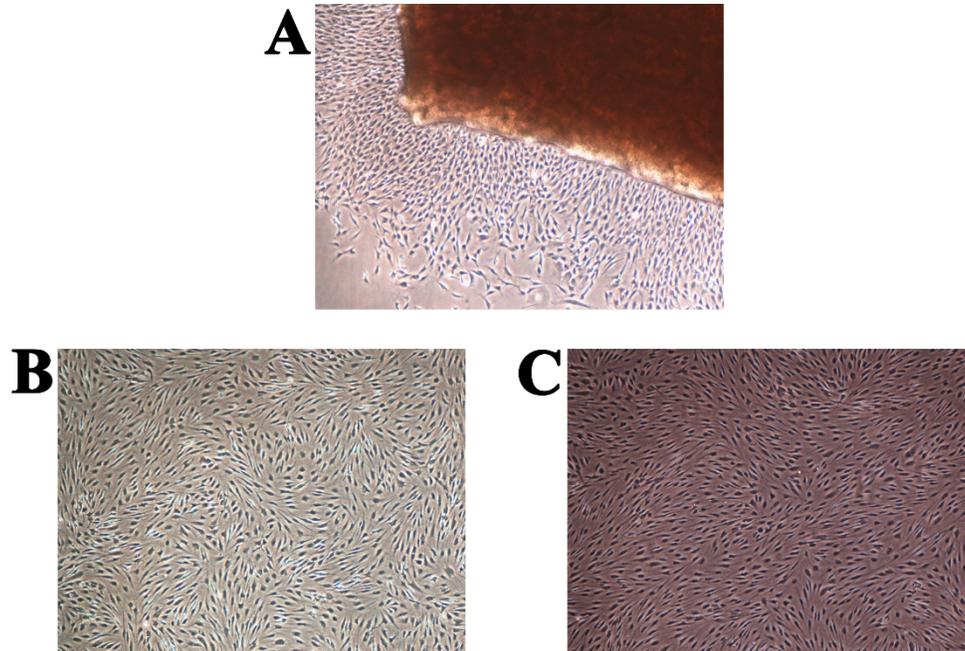


Fig. 3: Human chondrocyte culture: A) the explant culture of chondrocytes (“passage zero”), B) monolayer of chondrocytes after first passage, and C) monolayer of chondrocytes after the second passage. The magnification of all shown images is 50x (the inlay images were taken with a magnification of 100x).

200 3.2 Characterization of isolated chondrocytes

201 Now that we determined the suitable number of passages presumably yielding a high percentage
202 of chondrocyte cells, we performed additional characterization to confirm the chondrocytes’
203 desired phenotype. Analysis of gene expression was chosen due to its affordability and availability.
204 The isolated cells from the human articular cartilage were characterized in regard of the genes
205 related to specific chondrogenic production, namely collagen type 2 (COL2) and aggrecan
206 (ACAN). To detect possible dedifferentiation towards fibroblast like cells, expression of collagen
207 type 1 (COL1) was also determined. Expression of all three mentioned genes was performed after
208 the confluence was reached for the first (after 14 days – not shown) and second passage (after 7
209 days). As shown in **Fig. 4 and Fig. 5**, both cartilage specific genes (COL2 and ACAN) and also
210 the marker of fibrocartilage (COL1) were expressed in the isolated chondrocytes in both passages.
211 qPCR results are presented as absolute Ct values. Reference gene GAPDH was used as an internal
212 control (Chen et al. 2016).

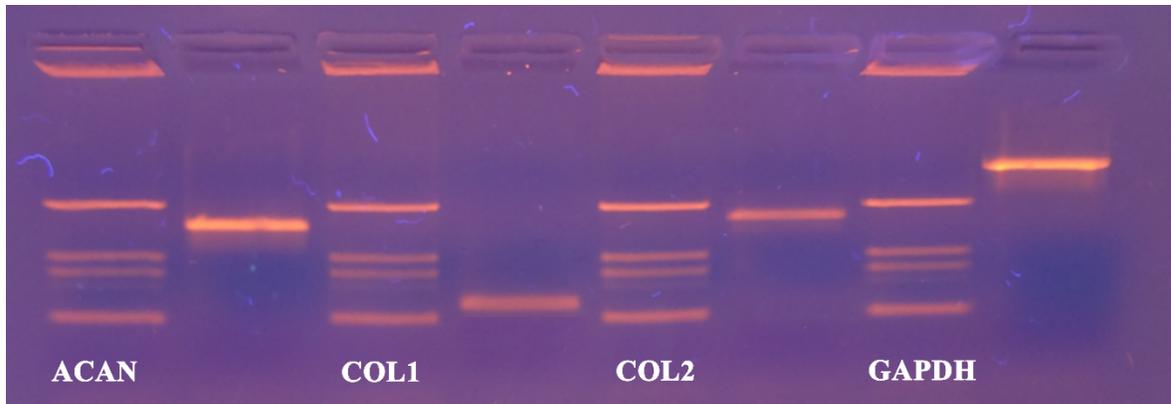


Fig. 4: cDNA products of analysed genes (GAPDH, collagen type 1, collagen type 2 and aggrecan) at the end-point of qPCR on agarose gel electrophoresis.

Analyzed genes: GAPDH (702 bp), COL2 (377 bp), COL1 (137 bp), ACAN (350 bp) and DNA markers (433 bp, 245 bp, 203 bp, 114 bp).

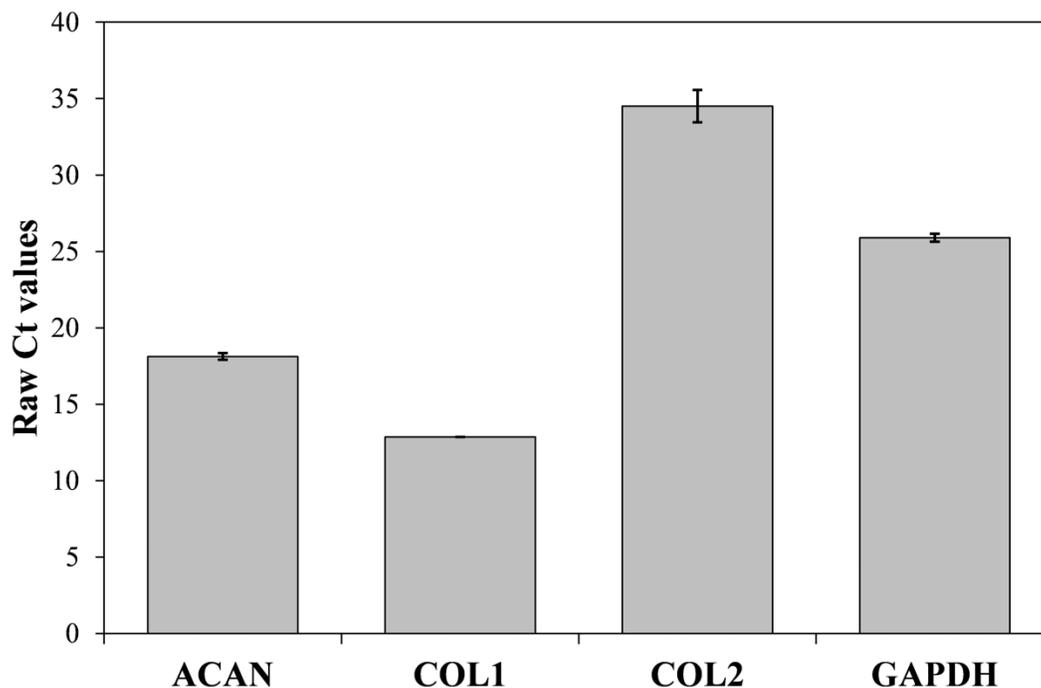


Fig. 5: Results of qPCR analysis presented as absolute Ct values of target genes expression (ACAN, COL1, COL2 and GADPH). The results are presented as average values with the standard errors of a triplicate.

213 4 DISCUSSION

214 The development of novel solutions related to any tissue engineering application consumes a huge
 215 amount of cells to prove safety and efficiency (Groeber et al. 2012; Maver et al. 2015; Mohd Hilmi
 216 & Halim 2015; Rodriguez-Vazquez et al. 2015). Cartilage tissue engineering is no exception, and
 217 hence large scale expansion of chondrocytes is required either for novel scaffold testing,
 218 determination of potential cytotoxic effects of medical devices and implants for orthopaedic use

219 (Bomer et al. 2016; Camarero-Espinosa et al. 2016; Makris et al. 2015). Cultivation of such high
220 cell counts is a demanding task, especially considering the low number of obtained cells in the
221 primary culture, and an often limited amount of available tissue. As a consequence, further
222 expansion and consecutive passages are needed, which on the other hand can lead to
223 dedifferentiation (Mirando et al. 2014; Shortkroff & Spector 1999; Thirion & Berenbaum 2004).
224 The latter is evident by morphological changes of the cells from polygonal to more elongated, as
225 well as through a reduction in the growth rate (Cetinkaya et al. 2011; Haudenschild et al. 2001;
226 Otero et al. 2012). For example, development of novel scaffolds for cartilage tissue engineering
227 often requires a million cells per experiment (the number depends on the size of the scaffold to be
228 tested), exposing the high demand for cells and at the same time one of the major bottle necks in
229 development of novel tissue engineering solutions. At later passages, the quality of chondrocytes
230 gradually decreases and is characterized with many of the phenotypic traits of fibroblast like cells
231 and an increased synthesis of collagen type 1, rather than type 2 (Bonaventure et al. 1994; Diekman
232 et al. 2009; Schnabel et al. 2002b).

233 A sufficient amount of cells can be ensured either through significant expenses (purchase of cells
234 from different cell banks) or isolation of desired cells from tissues. While the first scenario requires
235 sufficient funds, the latter requires appropriate tissue sources, an approval of respective
236 Committees of Medical Ethics, and a rigorous final analysis to confirm the isolation of the desired
237 cell type only. Since we are a small laboratory relying on limited available funds, yet work in the
238 close proximity and in tight collaboration with the local University Medical Centre, the second
239 scenario was more convenient. Our goal was to prepare an efficient and relatively less expensive
240 protocol, which would include the isolation of primary chondrocytes from full-thickness cartilage
241 that is surgically removed from the femoral condyle of an arthritic knee during total knee
242 arthroplasty (TKA). As a preliminary prove of the protocols' efficiency, we considered gene
243 expression analysis as the best option, since it is affordable and the required instrumentation (PCR,
244 inverted optical microscope) is most likely available in most cell biology laboratories.

245 In general, the chondrocyte isolation protocol can be divided into different stages: isolation,
246 seeding and chondrocytes grow in culture, although description and number of steps can vary
247 (Gosset et al. 2008; Thirion & Berenbaum 2004). After initial plating of the primary cultures, the
248 chondrocytes spread out after 2-3 days and after 4–7 days the sufficient amounts of total RNA may
249 be extracted. Primary cartilage phenotype (often confirmed by evaluating the presence of COL2
250 and ACAN mRNAs) may be initially preserved, but the expression of nonspecific collagens (e.g.
251 COL1) begins to appear already 7 days after isolation (Otero et al. 2005). Moreover, adult articular
252 chondrocytes are strongly contact-inhibited and undergo a rapid change in phenotype and gene
253 expression, termed “dedifferentiation”, when isolated from cartilage tissue and cultured on
254 culturing plastics (Haudenschild et al. 2001). Therefore, primary chondrocyte cultures should be
255 used for experimental analyses immediately before or just after confluence is reached to assure
256 optimal matrix synthesis and cellular responsiveness (Schneevoigt et al. 2016).

257 The protocol of chondrocyte isolation described in this article led to successful chondrocyte growth
258 and proliferation up to the second passage, yielding suitable number of cells for further studies.
259 The characterization of primary human chondrocytes by molecular analysis showed the expression

260 of cartilage specific genes (COL2 and ACAN), as well as a sign of dedifferentiation towards
261 fibrocartilage for the second passage (indicated by the expression of COL1).

262 The purpose of our study was not to revolutionize the chondrocyte isolation procedures, but rather
263 to push the evolution of cartilage tissue engineering. As such, our desire was to present an
264 alternative, affordable and relatively simple approach of chondrocyte isolation, especially suitable
265 for laboratories working closely together with orthopaedic clinics. Such laboratories have the
266 unique opportunity to use surgical waste materials, occurring during TKA. Since TKA is a very
267 common surgery (considering the present demographics, the incidence will only increase (Peterson
268 et al. 2015)), this approach could make cartilage related studies far more available also for
269 laboratories with limited resources, and hence push the overall development of this field towards
270 novel and cheaper therapeutic solutions. Based on our results, we can claim that the combination
271 of the use of surgical waste occurring during TKA, and analysis by inverted optical microscopy
272 and chondrocyte specific gene expression, indeed results in an alternative and affordable means to
273 boost cartilage related research in the future.

274 **5 CONCLUSION**

275 In this study we describe an efficient, simple and affordable procedure of isolation and cultivation
276 of human articular chondrocytes with a preserved chondrogenic phenotype to the second passage.
277 As the source material we propose the surgical waster occurring during total knee arthroplasty
278 (TKA). Chondrocyte cells are crucial not only for development of therapeutic approaches in
279 cartilage repair (e.g. autologous chondrocyte implantation – ACI), but are necessary in cartilage
280 tissue engineering to allow the development of functional cell models and novel scaffolds. For this
281 purpose, chondrocytes have to be isolated in sufficient quantities and their phenotype should be
282 preserved. Since all mentioned is related to very high costs, we propose alternative isolation and
283 testing protocols that are cheaper and could especially boost the preliminary studies related to
284 cartilage research.

285

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