1 Characterisation of the cells in repair tissue following autologous 2 chondrocyte implantation in mankind: a novel report of two cases.

3

4 Abstract

Aims: Autologous chondrocyte implantation (ACI) is used worldwide for the treatment of cartilage defects. This study has aimed to assess for the first time those cells that are contained within human ACI repair tissues several years post-treatment. We have compared the phenotypic properties of cells from within the ACI repair to adjacent chondrocytes and subchondral bone derivedmesenchymal stem cells (MSC).

11 **Materials and methods:** Two patients undergoing arthroplasty of their ACI-12 treated joint were investigated. Tissue and cells were isolated from the repair 13 site, adjacent macroscopically normal cartilage and MSC from the subchondral 14 bone and characterised for their growth kinetics, morphology, immunoprofile and 15 differentiation capacity.

Results: ACI repair tissue appeared fibrocartilaginous and ACI repair cells were heterogeneous in morphology and size when freshly isolated, becoming more homogeneous, resembling chondrocytes from adjacent cartilage, after culture expansion. The same weight of ACI repair tissue resulted in less cells than macroscopically normal cartilage. During expansion ACI repair cells proliferated faster than MSC but slower than chondrocytes. ACI repair cell immunoprofiles resembled chondrocytes, but their differentiation capacity matched MSC.

23	Conclusion: This novel report demonstrates that human ACI repair cell				
24	phenotypes resemble both chondrocytes and MSC but at different stages of their				
25	isolation and expansion in vitro.				
26	Keywords: Autologous chondrocyte implantation; histochemical analysis; repair				
27	cell characterisation, phenotype, morphology, growth kinetics, immunoprofile,				

28 differentiation potential.

30 Introduction

31 There is a huge interest worldwide in the development of tissue engineering and 32 cell based therapies for the treatment of cartilage defects. Autologous chondrocyte implantation (ACI) is a procedure that has been used for more than 33 34 20 years for the treatment of cartilage injury and osteoarthritis[1-3]. Our centre 35 has provided cells for over 400 ACI procedures since its inception, of which 81% 36 were a success, as indicated by a postoperative increase in Lysholm score[4]. 37 What happens to the culture expanded chondrocytes after implantation and the 38 contribution that they make to the repair tissue compared to cells from 39 surrounding tissues is still largely unknown[5-7]. Few preclinical studies have 40 labelled and tracked transplanted chondrocytes in ACI models. Those that have, 41 show that varying proportions of the cells injected form the cellular component of 42 the tissue at the site of ACI. In these studies transplanted cells have been shown to contribute in part to the formation and integration of repair tissues. However, 43 numerous unlabelled cells also form a major constituent, which suggests that 44 45 cells of unknown origin migrate to ACI-treated lesions and combine with 46 transplanted cells as part of the healing process[6-7].

The purpose of this study is to describe for the first time the phenotype of those cells that are contained within the tissue at the site of ACI in humans several years after treatment. Characterisation of the cells that are present at the site of ACI and hence, that are likely to produce and remodel the repair tissue, is critical to our understanding of the biological process in ACI. In previous studies we have only been able to assess the quality of ACI repair tissues in the clinic via magnetic resonance imaging (MRI) and histological analyses of small regions

(<1mm diameter cores)[8-11]. We have obtained two rare samples which have 54 55 provided us with the opportunity to isolate and examine the behaviour and 56 phenotypic properties of ACI repair cells in culture, in comparison to both 57 chondrocytes in the adjacent cartilage and MSC from the subchondral bone. 58 Observing ACI repair cells in culture will help to provide novel information on the cellular component of ACI repair tissues which we can then compare to 59 histological analyses and clinical outcome. In addition, by analysing the 60 61 properties of ACI repair cells in contrast to the phenotypes of cells isolated from 62 neighbouring tissues (e.g. cartilage and bone) we may begin to elucidate ACI 63 repair cell origin. Herein we describe the analysis of repair tissues and cells from two former ACI patients that have returned to our clinic for arthroplasty of their 64 ACI-treated joints several years post-ACI. 65

67 Materials and methods

68 **Patient information**

Following Local Research Ethical Committee approval and with informed consent, tissues obtained from two patients undergoing joint replacement surgery were included in this study (one knee – Patient 1 and one hip – Patient 2). Both were males aged 49 and 44 years at the time of arthroplasty, which was 11 and 5 years, respectively, after previous ACI treatment (Table 1).

74 Histological analysis

For histological examination decalcified wax-embedded tissue sections from the 75 76 region bridging macroscopically normal cartilage (MNC) and ACI repair tissues 77 (with subchondral bone) from Patients 1 and 2 were examined using haematoxylin and eosin (H&E) and toluidine blue stains as described 78 previously¹². For H&E histological examination, sections were flooded with 79 80 Mayer's haemalum (VWR International Ltd, Poole, UK) for 1 minute, drained, and washed in tap water for 5 minutes. Slides were then flooded with 1% eosin 81 82 aqueous solution (BDH, Poole, UK) (in distilled water) for 30 seconds, briefly washed in tap water and dehydrated through a series of isopropanol (Genta 83 Medical, York, UK) concentrations in distilled water and xylene, for 5 minutes 84 each. Following dehydration, the tissue sections were mounted under glass 85 coverslips (Cell Path Ltd, Newtown, UK) with Pertex mounting medium (Histolab 86 87 Products AB, Gothenburg, Sweden) and allowed to air dry. Glycosaminoglycan (GAG) content was assessed by metachromasia by flooding sections with 1% 88 89 aqueous toluidine blue (BDH) solution for 30 seconds and rinsed in tap water.

Slides were left to air dry before mounting under glass coverslips with Pertexmounting medium.

92 For collagen type II immunolocalisation, dewaxed and rehydrated sections 93 were pre-treated with 0.1% (w/v) hyaluronidase and 0.2% (w/v) trypsin (Sigma-Aldrich, Poole, UK) for 1 hour at 37°C. Sections were then washed in phosphate 94 95 buffered saline (PBS) and incubated for 2 hours at room temperature in a 96 humidified chamber with 10µg/ml of primary mouse monoclonal collagen type II 97 antibody (clone CIIC1, Developmental Studies Hybridoma Bank, University of 98 lowa) in PBS. Parallel sections were incubated with a non-specific, isotype-99 matched antibody (IgG1- Dako, Glostrup, Denmark) instead of the primary 100 antibody at the same concentration as a negative control. After incubation with the primary antibodies all sections were washed in PBS before incubation for 1 101 102 hour with a secondary biotinylated antibody at 50µg/ml (Vector ABC System, 103 Vector Laboratories, Peterborough, UK) according to manufacturers instructions. To eliminate endogenous peroxidase activity sections were blocked with 3% (v/v) 104 hydrogen peroxide (H₂O₂) in methanol (BDH) for 30 minutes. Collagen type II 105 106 immunopositivity was finally visualised by testing for bound peroxidise, which is 107 detected by incubation with a substrate of diaminobenzidine tetrahydrochloride 108 (DAB), activated by H_2O_2 . The sections were then dehydrated before mounting 109 under glass coverslips with Pertex mounting medium as described previously.

110 Chondrocyte and ACI repair cell isolation and culture

Approximately 300mg of MNC and ACI repair tissues were harvested from the medial femoral condyle (MFC) of Patient 1 and the femoral head (FH) of Patient

2. Cells were isolated and cultured as described previously for chondrocytes[13]. 113 MNC and ACI repair tissues were dissected into ~2mm³ pieces and placed into 114 25cm² tissue culture flasks (Falcon 250 ml Polvstvrene Tissue Culture Flask, BD 115 116 Biosciences, UK). The weight of each tissue type was recorded and cells 117 released by enzymic digestion. Dulbecco's Modified Eagle's Medium (DMEM)/F12 (Life Technologies, Paisley, UK) containing 0.8mg/ml type XI 118 collagenase (Sigma-Aldrich) was added to each of the flasks which were then 119 incubated at 5% (v/v) CO_2 for 20 hours at 37°C. 120

121 Following this incubation, each tissue digest was passed through a 70µm cell strainer (BD Biosciences); cells were recovered by centrifugation at 750g for 122 10 minutes to form a cell pellet. Cells were plated out in DMEM/F12, 123 supplemented with 10% foetal bovine serum (FBS) (Life Technologies), 50µg/ml 124 ascorbic acid (AA; Sigma-Aldrich) and 1% (v/v) penicillin and streptomycin (P/S; 125 Life Technologies) at a seeding density of 5 x 10^5 cells per cm². After 5 days, 126 non-adherent cells were removed and the adherent cell population was cultured 127 128 in monolayer in DMEM/F12 10% FBS medium supplemented with AA and P/S. Cells were routinely passaged at 70% confluence by trypsinisation (0.05% v/v 129 Trypsin-EDTA) and re-seeded at 5×10^3 cells/ cm². 130

131 Human bone marrow stem cell (MSC) isolation and culture

The underlying bone from the MFC of Patient 1 and the FH of Patient 2 was
perfused with DMEM/F12 (Life Technologies) supplemented with 10% FBS and
P/S. Mononuclear cells isolated and MSC cultured as described previously[14].
Mononuclear cells isolated by density gradient centrifugation (Lymphoprep,

Fresenius Kabi Norge, AS) were plated out in DMEM/F12, supplemented with 136 20% FBS and P/S at a seeding density of 20 x 10⁶ cells per 25cm² tissue culture 137 138 flask. After 24 hours, non-adherent cells were removed and the adherent cell population was cultured in monolayer in DMEM/F12 10% FBS medium 139 140 supplemented with P/S. Cells were routinely passaged at 70% confluence by trypsinisation (0.05% v/v Trypsin-EDTA) and re-seeded at 5 x 10³ cells/ cm². 141 Viability was assessed at each passage by trypan blue exclusion (Sigma-142 143 Aldrich).

144 Microscopy, image capture and analysis

Histological sections were viewed using bright light and polarized light 145 146 microscopy (Leitz Diaplan, Wetzlar, Germany) and digitized images were 147 captured with a Nikon digital camera (DS-Fi1, Nikon, Kingston-upon-Thames, 148 UK). Cultures were viewed using phase contrast microscopy (Nikon Eclipse 149 TS100) and digitized images were captured with a Hamamatsu digital camera 150 (C4742-95, Bridgewater, NJ). The mean cell area was determined from passage 151 0-3 for the three cell populations using IPLab software (Version 3.6, Nikon). For each cell type, results from at least 5 separate images per culture were 152 153 combined.

154 Growth kinetics

155 Culture doubling time was calculated for each cell population (from passage 0-3) 156 using the following formula: doubling time (DT) = $(t2-t1) \times \ln(2)/\ln(n2/n1)$, where 157 t1 = the time of cell seeding, t2 = the time of cell harvest and n = the matching 158 cell numbers at these time points.

159 Immunoprofiling

Immunoprofiling via flow cytometry was used to assess culture expanded cells (at 160 161 passage 2) using a FACScan flow cytometer (BD Biosciences). A profile typical of MSC was targeted [15]. In brief, cells were blocked for one hour in a buffer of 162 10% normal human Ig (Grifols, Cambridge, UK). Cells were then incubated with 163 164 mouse anti-human monoclonal primary antibodies against CD14, CD19, CD31, 165 CD34, CD45, CD73, CD90, CD105 and HLA-DR (all phycoerythrin-conjugated) 166 (Immunotools, Friesoythe, Germany) for 30 minutes. Matched cell populations were also exposed to isotype-matched IgG negative control antibodies (Sigma-167 Aldrich). Immunoprofiles were produced using Cell Quest software (BD 168 169 Biosciences).

170 Multipotency assays

Established protocols [16-18] were used to assay the differentiation potential of 171 172 cells at passage 2 for adipogenic, osteoblastic and chondrogenic lineages. In 173 brief, for 21 days, cell cultures were exposed to appropriate conditions for: (i) 174 adipogenic differentiation via monolayer culture in DMEM/F12 10% FCS, 1% ITS-175 dexamethasone, 3-isobutyl-1-methylxanthine X (Life Technologies), and 176 indomethacin (Sigma-Aldrich); (ii) osteoblastic differentiation via monolayer 177 culture in DMEM/F12 10% FCS, ascorbate 2-phosphate, dexamethasone and β -178 glycerophosphate (Sigma-Aldrich); (iii) chondrogenic differentiation via micro-179 mass pellet culture in DMEM/F12, 1% ITS-X, ascorbate 2-phosphate (Sigma-180 Aldrich), dexamethasone (Sigma-Aldrich) and TGF-B1 (PeproTech Ltd., London, 181 UK).

At the 21 day time point, adipogenic differentiation potential was examined via oil red-O visualisation of lipid formation, alkaline phosphatase activity was used to assess osteoblast differentiation and for chondrogenic differentiation, toluidine blue staining was used to detect the presence of GAGs in micro-mass pellets.

187 Statistical analysis

188 The Kruskal-Wallis non-parametric ANOVA and post-hoc Bonferroni pair-wise

189 comparison tests were used to assess significant differences between the size

190 (area coverage) of each cell type isolated from the same joint e.g. Chondrocytes,

191 ACI repair cells and MSC between passages 0-3.

193 **Results**

194 Histological analysis

195 Tissue sections from the region bridging MNC and ACI repair tissues (Figure 1A) 196 were examined via H&E staining (Figure 1B) which demonstrated a disorganized 197 cellular distribution throughout ACI repair tissues in both patients compared to 198 neighbouring MNC. There appeared to be good integration between the repair 199 cartilage and the neighbouring MNC and the underlying bone. The surface 200 integrity and smoothness of ACI repair tissues differed between patients; in the 201 knee (patient 1) the repair surface was rough, whereas the hip repair tissue 202 (patient 2) was smooth but undulating compared to neighbouring MNC which was 203 smooth and flat in both patients.

Viewing the same section under polarized light (Figure 1C) revealed illuminated areas of scattered light indicative of fibrocartilage throughout the repair region and in particular at the border with MNC. Toluidine blue staining (Figure 1D) indicated that the GAG content of ACI repair tissues was lower than that in neighbouring MNC, particularly in the surface zone. For both patients collagen type II staining (Figure 1E) was similarly weaker in the surface zone of ACI repair tissues.

211 Morphological analyses

For both patients freshly isolated chondrocytes and MSC had a uniform fibroblast-like morphology with their size (surface area coverage) increasing after expansion. MSC appeared consistently larger at each passage compared to chondrocytes. In addition, for MSC cultures numerous intracellular stress fibres were clearly visible at passage 3. In contrast, the morphologies of freshly isolated

ACI repair cells were more heterogeneous. ACI repair cells possessed a combination of early and later passage chondrocyte and MSC-like morphologies after initial seeding but became homogeneous after sub-culture. There were no obvious or significant differences between ACI and chondrocyte cultures (Figures 2A and 2B).

Growth kinetics

223 Chondrocytes and ACI repair cells proliferated at similar rates for both patients 224 (from passages 0-3), with DTs of 2 weeks between passages 0-1, decreasing to 225 4-6 days between passages 1-2 and 2-3. In contrast, the DT of MSC was 226 markedly higher at 8-14 days between passages 1-2 and 2-3 (Figure 3A). At 227 passage 3 chondrocyte cultures produced harvests of 1.4×10^8 , ACI repair cells 228 4.3×10^7 and MSC 9.0×10^6 (Figure 3B). Viability for all cells was >98% at each 229 passage.

230 Immunoprofiles

231 MSC from both patients' were CD14, CD19, CD31, CD34, CD45 and HLA-DR 232 negative and CD73, CD90 and CD105 positive; this matches previously published MSC immunoprofiles¹⁵. Chondrocytes and ACI repair cells from both 233 234 patients had MSC-like immunoprofiles apart from some positivity for CD14; for 235 patient 1 CD14 was detected on ~90% of chondrocytes and ~50% of ACI repair cells, whereas for patient 2 the reverse pattern of positivity was observed 236 between chondrocytes (~50% immunopositive) and ACI (~90% immunopositive) 237 238 (Figure 4).

239 **Differentiation potential**

Chondrocytes, ACI repair cells and MSC from both patients differentiated along 240 241 all three mesenchymal cell lineages tested but to varying degrees, as delineated 242 by lipid accumulation, alkaline phosphatase activity and toluidine blue GAG 243 staining. Chondrocytes from patient 1 produced a large frequency of clustered 244 globular lipids, whereas chondrocytes from patient 2 showed more diffuse staining of smaller lipids throughout. ACI and MSC staining for lipid accumulation 245 was similar in pattern for both patients, hence, a few unilocular lipid clusters 246 247 (which may be indicative of committed adipocytes) were seen in both (Figure 5A). 248 Chondrocytes from both patients showed intense uniform staining for alkaline 249 phosphatase activity, whereas ACI repair cells and MSC demonstrated a more 250 heterogeneous pattern of staining (Figure 5B). All pellet cultures showed the presence of some GAGs via toluidine blue staining. Chondrocyte fractions 251 252 showed the most intense toluidine blue staining, ACI repair cell and MSC pellets 253 showed weaker GAG staining for both patients (Figure 5C).

255 **Discussion**

256 These samples have provided a unique opportunity to study both the histology of 257 the complete area of ACI repair tissue compared to neighbouring MNC as well as 258 examining the phenotype of ACI repair cells compared to chondrocytes from 259 adjacent cartilage and MSC from the underlying subchondral bone, several years 260 post-ACI. For these analyses, tissues and cells were harvested from two former ACI patients at the time of arthroplasty. It is debatable whether the patients 261 262 included in this study and hence, the tissues and cells examined here should be 263 considered as ACI successes or failures. These individuals were 38 and 39 years 264 of age at the time that they received ACI and although their treated joints 265 ultimately failed, ACI prolonged the life of their natural joints for 11 and 5 years, respectively. For patients of such a young age this may have important 266 267 implications for later life as joint replacement is accompanied by some loss of 268 function which restricts activities [19]. In addition, joint replacements will probably 269 not last for the full life of younger patients. Hence, the initial surgery may need 270 revision which is a more complicated and expensive operation with lower 271 success rates than primary arthroplasty[20]. As such, ACI therapy which 272 increases the life-span of these young patients' joints may understandably be 273 considered a successful intervention even though their treated joints eventually 274 failed.

275 Histologically, we have shown that ACI repair tissues resembled those 276 previously described[8-10]; cells were numerous but disorganised and matrices 277 appeared predominantly fibrocartilaginous. Repair tissues were well integrated

with adjacent MNC and subchondral bone. The surface of repair tissue for patient 278 279 1 was rough compared to that of patient 2, perhaps because patient 1 received a 280 periosteal graft, which may exhibit hypertrophy, compared to the collagen 281 membrane which was used for patient 2[21]. There were no obvious histological 282 signs that may have explained joint failure in these patients. However, there is 283 limited evidence that histology is a reliable indicator of clinical outcome for ACI patients[22]. In this study we have some preliminary data for ACI repair cell 284 285 characterisation, which may represent a promising additional prognostic marker 286 in future analyses. An understanding of the ACI repair cell phenotype will help to 287 elucidate ACI repair tissue formation and remodelling processes. In addition, we 288 may be able to use this data to begin to 'unpick' the origin of ACI repair cells and 289 hence, to determine if ACI tissues contain any of those cells initially implanted at 290 ACI stage II. In preclinical studies transplanted cells are known to persist in ACI 291 repair zones for up to 14 weeks in large animal models[6]. Alternatively, the site 292 of ACI repair may contain a completely different cell type (e.g. synovium, bone, or 293 bone marrow-derived MSC) that have migrated and integrated into ACI zones 294 from surrounding tissues[23-25].

It is likely that the anatomical location (i.e. knee versus hip) and the patch used at ACI (i.e. periosteum versus Chondro-Gide®) will have influenced the quality and extent of the repair tissue observed for Patient 1 compared to Patient 2. However, there were no discernible differences observed in the phenotypes of ACI repair cells isolated from Patient 1 or Patient 2. This suggests that the dissimilarities between these examples of ACI i.e. the type of joint treated and

patch used might not have contributed significantly to the tissue regeneration 301 302 seen (or the cells involved) in these ACI treated joints. We have shown that 303 freshly isolated cells from ACI repair tissues appeared to contain a mixture of 304 chondrocyte and MSC morphologies, but that MSC-like cells disappear over time 305 in culture. Our growth kinetics data supports the theory that chondrocytes may 306 have outgrown MSC in vitro. In addition, the immunoprofiles of chondrocytes and 307 ACI repair cells were similar after sub-culture, both demonstrated some CD14 308 positivity, a marker found on freshly isolated chondrocytes[26,27] compared to a 309 complete absence of CD14 on MSC at the same passage. It is unlikely that the 310 source of CD14 positive cells that were cultured from ACI repair tissues 311 represent MSC which have migrated from surrounding tissues e.g. MSC from 312 synovium and bone or bone marrow as these MSC do not express CD14 in an 313 undifferentiated state[15,28]. It is conceivable that these cells instead either 314 represent a proportion of the chondrocytes that were originally transplanted at 315 ACI or that have migrated from adjacent cartilage. Alternatively, an MSC 316 population may have homed to the injured region and differentiated in vivo 317 towards a chondrogenic lineage[29,30].

Interestingly, our multipotency studies showed a marked contrast between passaged chondrocyte and ACI repair cell differentiation potential. Chondrocyte populations differentiated along adipogenic, osteogenic and chondrogenic lineages in a strongly positive and uniform manner as articular cartilage itself contains a multipotent progenitor cell population [31-35]. In contrast, ACI repair cells appeared heterogeneous and on the whole, differentiated along each

324 mesenchymal lineage tested to a lesser extent, akin to MSC isolated from the 325 same joint. However, we acknowledge that culture expansion may change the 326 characteristics of ACI repair cells. It is therefore difficult to ascertain exactly how the immunoprofile and differentiation potential of culture-expanded ACI repair 327 328 cells relate to their in vivo characteristics. None the less, this study demonstrates 329 that although cultured ACI repair cells possess many phenotypic characeristics of 330 chondrocytes e.g. similar morphologies, growth kinetics and immunoprofiles, they 331 do not possess the ability to differentiate, importantly in this setting, into 332 chondrocytes with physiologically relevant properties. For example, ACI repair 333 cells do not appear to synthesise GAG rich matrices to the same extent as 334 chondrocytes when chondrogenically induced in vitro. This finding coincides with 335 our in vivo histological evidence which clearly shows that ACI repair tissues 336 possess a lower GAG content in comparison to neighbouring MNC.

337 A limitation of this study is that it is based on only two human samples. 338 making the findings a little difficult to interpret. We hope to expand and corroborate the findings of these pilot experiments by increasing the sample size 339 of donors and expanding the molecules and markers to be investigated in longer 340 341 term studies. For example, immunohistochemical analyses to determine the 342 presence of type X collagen as a marker of hypertrophy in repair tissues and 343 gene expression studies of isolated repair cells for osteogenic and chondrogenic associated molecules using RT-qPCR would provide valuable additional data. 344 345 Nonetheless, this pilot study has provided a rare opportunity to carry out studies 346 which are normally only possible in animals, but even then at much shorter time

- 347 points. The information obtained is therefore completely novel and likely to be
- 348 more relevant to the human patient than results from animal studies.

350 Conclusions

This study presents the first data on human ACI repair cell phenotypes in culture, 351 352 several years after ACI treatment. Cells isolated from ACI repair tissue appeared 353 to contain a mixture of chondrocytes and MSC morphologies at initial seeding, 354 but became more like chondrocytes with regard to morphology, proliferation and 355 immunoprofile at later passage. However, the differentiation potential of 356 expanded ACI repair cells was reduced for each mesenchymal lineage tested 357 compared to chondrocytes and notably so for chondrogenic potential which is 358 considerably relevant in this setting. These findings indicate that ACI repair cells are composed of a mixture of cells with features resembling both chondrocyte 359 360 and MSC phenotypes. This suggests that ACI tissues contain both chondrocytes 361 (either originally implanted or integrated from surrounding cartilage in vivo) and 362 also MSC that have infiltrated the treated region from synovium or subchondral 363 bone. A better understanding of the source of cells which contribute to the repair 364 tissue in ACI, especially when associated with the best clinical outcome, will provide valuable information to help improve the ACI technique in the clinic. For 365 366 example, we may be able to select the most effective cells prior to implantation or 367 to augment the migration of desirable endogenous cells from the nearby tissues. 368 In this way we can make step changes and improvements in current cell therapy treatments of chondral defects. 369



371 Summary Points

372 **Patient Information**

• We have analysed tissues and cells isolated from regions of repair in two patients undergoing arthroplasty several years after cell therapy with autologous chondrocytes; these have been compared to tissues and cells derived from adjacent cartilage and bone.

377 Histological analysis

- ACI repair tissues were densely populated with cells but the extracellular
- 379 matrix was disorganised and contained little GAG or collagen type II in
- 380 surface zones compared to adjacent, macroscopically normal cartilage.

381 Morphological analyses

Cells which were isolated freshly from ACI repair tissues had a mixture of
 chondrocyte and MSC-like morphologies.

Growth kinetics, Immunoprofiles and Differentiation potential

Following culture expansion, cells isolated from ACI repair tissues
 resembled chondrocytes in terms of their growth and immunoprofile but
 their adipogenic, osteogenic and (importantly) chondrogenic differentiation
 capacity was markedly reduced in comparison to chondrocytes isolated
 from adjacent macroscopically normal cartilage.

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512 **Figure Legends**

513 Figure 1. Histology of the ACI: adjacent macroscopically normal cartilage (MNC) 514 interface. Dotted line shows the ACI treated areas. A, Samples received following 515 arthroplasty, showing ACI repair regions at the centre of the medial condyle 516 (patient 1, top) and femoral head (patient 2, bottom). Regions dissected for histology are indicated (boxed). B. Haematoxylin and eosin stained sections 517 showing a disorganized cell distribution in ACI repair tissues (left) compared to 518 519 typical hyaline cartilage cell organization (right). C, Polarized light illumination 520 showing fibrocartilage (bright scatter) in ACI repair tissues (left), particularly 521 evident at ACI: MNC interfaces (centre). D, Toluidine blue localization of GAGs 522 demonstrates that there are fewer GAGs present in ACI repair regions (left) 523 compared to MNC (right). A dramatic reduction in GAG staining intensity is seen 524 in the surface zone of ACI repair tissues. E, Collagen type II immunolocalization 525 demonstrates that ACI repair region matrices (left) contain less collagen type II 526 compared to MNC (right). All calibration bars = 1 mm.

527 Figure 2. Cell characterisation: Morphology. A. Freshly isolated cells are shown 528 (top panels) compared to cells at passage 3 (bottom panels). Chondrocytes (C) 529 and mesenchymal stromal cells (MSC) isolated from both patients have a uniform 530 fibroblast-like morphology which increase in size (area coverage) with subculture. Freshly isolated ACI repair cells (ACI) show a heterogeneous 531 532 morphology which becomes homogeneous at passage 3. All calibration bars = 200µm. B, Chondrocytes (black bars) and MSC (white bars) increase in size from 533 534 passages 0-3. Chondrocytes are significantly smaller than MSC at every passage

quantified. At initial seeding ACI repair cells (grey bars) are significantly larger than chondrocytes, but similar in size to MSC. At passage 3 ACI repair cells are significantly smaller than MSC, but similar in size to chondrocytes (means +/-SEM Kruskal-Wallis ANOVA and post-hoc Bonferroni ***p <0.001).

Figure 3. Cell characterisation: Growth kinetics. **A**, The doubling times (DT) of chondrocytes and ACI repair cells decrease similarly through passages 0-3. In contrast, MSC DT increase between passages 1-3 (means +/- SEM from pooled patient data). **B**, After harvest at passage 3 chondrocyte cultures produced 1.4X10⁸ cells, ACI repair cultures 4.3X10⁷ cells and MSC cultures 9.0X10⁶ cells (means +/- SEM from pooled patient data).

Figure 4. Phenotypic characterisation: Immunoprofile and differentiation capacity 545 546 (passage 2 cells). A, Flow cytometry analysis for CD-immunolabelling of 547 chondrocytes, ACI repair cells and MSC. All cultures tested are CD19-ve, CD31-548 ve, CD34-ve, CD45-ve, CD73+ve, CD90+ve, CD105+ve and HLA-DR-ve at 549 passage 2. C and ACI repair cells are also immunopositive for CD14 to varying 550 degrees (patient 1 chondrocytes are ~50%+ve and ACI are ~90%+ve, patient 2 chondrocytes are ~90%+ve and ACI are ~50%+ve), where as MSC are CD14-ve. 551 552 The green-lined histogram denotes immunopositivity (fluorescence intensity) for 553 each indicated marker, whilst the purple histogram denotes immunolabelling with an isotype-matched control antibody. 554

Figure 5. Phenotypic characterisation: Differentiation capacity (passage 2 cells). **A,** The presence of lipid vesicles is increased in chondrocytes treated with adipogenic stimuli (but a different type of staining pattern between patients can be observed, as revealed with oil red-O), compared with ACI repair cells or MSC

(which had similar localised unilocular lipid staining in patches). **B**, Alkaline phosphatase activity is markedly increased in chondrocyte cultures treated with osteogenic stimuli compared with ACI repair cells or MSC. **C**, Toluidine blue staining of chondrogenically induced pellet cultures shows more intense staining in chondrocyte pellets compared to ACI repair cell or MSC pellets. Calibration bars = 100μ m, inset calibration bar = 200μ m.

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	Patient 1	Patient 2
Time since ACI	135 months (~11 years)	59 months (~5 years)
Site of ACI	Medial Femoral Condyle	Lateral aspect of Femoral Head
Size of defect	25 x 15mm	12 mm diameter
Cells received	Knee chondrocytes	Hip chondrocytes
Patch received	Periosteum	Chondro-Gide®