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Fermor, HL, McLure, SWD, Taylor, SD et al. (4 more authors) (2015) Biological, biochemical and biomechanical characterisation of articular cartilage from the porcine, bovine and ovine hip and knee. Bio-Medical Materials and Engineering, 25 (4). 381 - 395. ISSN 0959-2989

https://doi.org/10.3233/BME-151533

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Biological, biochemical and biomechanical characterisation of articular cartilage from the porcine, bovine and ovine hip and knee

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Abbreviated title for use as a running headline: Biological, biochemical and biomechanical characterisation of articular cartilage

Keywords: Cartilage, Glycosaminoglycan, Collagen, Thickness, Water Content, Cellularity, Percentage Deformation, Permeability, Equilibrium Elastic Modulus

Abstract

This study aimed to determine the optimal starting material for the development of an acellular osteochondral graft. Osteochondral tissues from three different species were characterised; pig (6 months), cow (18 months) and two ages of sheep (8 - 12 months and > 4 year old). Tissues from the acetabulum and femoral head of the hip, and the groove, medial and lateral condyles and tibial plateau of the knee were assessed. Histological analysis of each tissue allowed for qualification of cartilage histoarchitecture, glycosaminoglycan (GAG) distribution, assessment of cellularity and cartilage thickness. Collagen and GAG content were quantified and cartilage water content was defined. Following biomechanical testing, the percentage deformation, permeability and equilibrium elastic modulus was determined. Results showed that porcine cartilage had the highest concentration of sulphated proteoglycans and that the condyles and groove of the knee showed higher GAG content than other joint areas. Cartilage from younger tissues (porcine and young ovine) had higher cell content and was thicker, reflecting the effects of age on cartilage structure. Cartilage from older sheep had a much higher elastic modulus and was less permeable than other species.

1 Introduction

2 Osteoarthritis (OA) is the most common disorder affecting joints [1]; this degenerative 3 disease causes progressive degradation of articular cartilage, bone and surrounding joint 4 tissues, resulting in pain and a loss of function [2]. In a proportion of patients OA has been 5 reported to originate from an initially minor cartilage defect [3], which, due to the avascular 6 and aneural nature of cartilage, is unable to heal and deteriorates with regular joint loading 7 and articulation [4]. However, OA may also result from changes in bone, soft tissues or 8 secondary to trauma or acute injury. Osteochondral lesions in themselves can cause pain, 9 swelling and mechanical problems, reducing the quality of life for patients [5]. Current 10 orthopaedic treatments for chondral defects, such as lavage and debridement [6, 7] and 11 marrow stimulation [8] have shown variable long term outcomes. Similarly, 12 perichondrial/periosteal grafting, mosaicplasty [9] and autologous chondrocyte implantation 13 [10] may result in donor site morbidity amongst other complications [5, 11]. Due to the 14 limitations of current treatments, chondral lesions present an ideal target for early 15 intervention tissue engineered therapies in the OA disease process. It is proposed here that an 16 acellular, xenogenic, osteochondral biological scaffold may provide the ideal graft for defect 17 repair.

18

Articular cartilage has a highly organised, specialised composition and structure, which allows it to function as a low friction, load-bearing surface [12]. The major tissue components are collagen type II, proteoglycans, water and cells. The collagen forms a characteristically oriented fibril network which provides the tensile strength of the tissue [13, 14].
Proteoglycans are dispersed throughout this network and bind water in the tissue, allowing it to resist compression [15]. Chondrocytes are responsible for cartilage matrix homeostasis

1

2 The aims of this study were to provide a comprehensive analysis of articular cartilage from 3 four animals as a potential source of acellular xenogeneic biological scaffold, assessing the 4 various matrix components and the resultant mechanical properties of each tissue. Cartilage 5 from the hip and knee joints of pigs, cows and two ages of sheep was assessed. Cellularity, 6 cell distribution, GAG content and distribution, collagen concentration, water content and 7 cartilage thickness was determined and the percentage deformation, permeability and 8 equilibrium elastic modulus of each tissue defined. The characterisation of these different 9 tissues will enable determination the most optimal starting material for the development of an 10 acellular xenogenic osteochondral biological scaffold.

11 Materials and Methods

12 Osteochondral samples

13 Legs of 9 month old pigs, 8 - 12 month old lambs, 4 - 8 year old ewes and femurs of 18 14 month old cows were obtained from a local accredited abattoir within 24-76 hr of slaughter. 15 Extra-skeletal tissues were removed using a scalpel and forceps to expose the joint surfaces. 16 Exposed cartilage was regularly moistened with phosphate buffered saline (PBS [Oxoid]) to 17 maintain hydration. Long bones were held in a specially designed clamp to immobilise them 18 while allowing access to the joint surfaces, acetabula were held in a bench-top vice. Using an 19 electric drill with a corer attachment, osteochondral plugs, 9 mm diameter and 12 mm deep, 20 were collected from the femoral head, patellofemoral groove, medial and lateral condyles and 21 medial and lateral tibial plateau of all species. Bovine acetabular tissue was not available and was collected from sheep and pigs. Plugs were washed in PBS before being stored at -20 °C 22 23 on PBS moistened filter paper. During dissection, macroscopic observations of the cartilage 24 from each species was made.

1 Histology

2 Osteochondral plugs (n=5) were fixed for 48 hours in 10 % (v/v) neutral buffered formalin 3 (NBF) [Bios Europe Ltd], before being decalcified in 12.5 % (w/v) diaminoethane tetraacetic 4 acid disodium salt [Na₂EDTA pH 7: Fisher Scientific] for 4 weeks or until soft enough to be 5 cut with a scalpel. Plugs were bisected before being dehydrated and paraffin wax embedded using an automated process [Lecia TP 1020, Lecia Microsystems]. Sections of 6 µm 6 7 thickness were cut through the cartilage surface into the bone, encompassing the different 8 cartilage zones and subchondral bone. Haematoxylin and eosin (H&E) [Bios Europe Ltd] 9 staining was used to assess tissue histoarchitecture. From the resultant digital images, Image J 10 software was used to delineate the superficial, middle and deep zones of cartilage (top 15 %, 11 middle 50 % and bottom 35 % of cartilage thickness respectively [17]) and cell shape analysis was performed on 500 μ m² regions from each zone, giving particle count, area and 12 13 roundness data. Roundness was calculated by the software as follows:

$$roundness = \frac{4 \times area}{\pi \times major \ axis^2}$$

with a measurement of 1.0 being perfectly round. One measurement for each zone was performed per image. GAG distribution within the tissue was visualised by staining with alcian blue [Bios Europe Ltd] (pH 2.5) and counterstaining with periodic acid Schiff and Mayers haematoxylin [Sigma]. This stain provided good definition of the cartilage-bone interface, so image software was used to measure the cartilage thickness. Measurements (n=6) were taken from the cartilage surface to the point at which alcian blue staining terminated, the mean of these measurements was calculated for each image.

21 **Biochemical Assays**

Sulphated sugar assay: Cartilage from each joint area (n=5) was cut away from the
 subchondral bone and lyophilised to a constant weight before being enzymatically digested

using papain solution (5 ml, 50 U.ml⁻¹ papain [Sigma] in PBS at pH 6 with 5mM L-cysteine 1 2 hydrochloride [Sigma] and 5mM Na₂EDTA [Fisher Scientific]) for 36-48 hours at 60 °C. 3 GAG content was quantified following a method adapted from Farndale et al. [18]. Briefly, 4 standard concentrations of chondroitin B sulphate [Sigma] were produced, 1,9-5 dimethylmethylene blue [Sigma] solution (250 µl) was added to each standard and sample 6 (40 µl) in a clear flat bottomed 96-well plate, the plate was agitated for 2 minutes before the 7 absorbance was read on a spectrophotometer [Multiscan Spectrum, Thermo Labsystems] at 8 525 nm. Interpolation of sample absorbance from the standard curve gave the GAG concentration of each digest which was then expressed in μ g.mg⁻¹ of dry weight of tissue. 9 10 11 Hydroxyproline assay: Cartilage was lyophilised, as above, before being hydrolysed in 6 M 12 hydrochloric acid [HCl: VWR] for 16 hours at 80 °C followed by neutralisation with 6 M 13 sodium hydroxide [NaOH: Fisher]. Hydroxyproline content was quantified following a 14 method adapted from Edwards & O'Brien [19]. Briefly, standard concentrations of trans-4-15 hydroxy-L-proline [Sigma] were produced, Chloramine-T oxidisation solution [Sigma] (100 16 μ l) was added to each standard and sample (50 μ l) in a clear flat bottomed 96-well plate, the 17 plate was agitated for 5 minutes. Ehrlich's reagent [Sigma] (100 µl) was then added and the 18 plate was incubated in a water bath at 60°C for 45 minutes before the absorbance was read at 19 570 nm on a spectrophotometer. Interpolation of sample absorbance from the standard curve gave the hyrdoxyproline concentration of each sample. This was then expressed as μ g.mg⁻¹ 20 21 dry weight of tissue.

22

Water content of samples (n=5) was determined during cartilage lyophilisation. Sample
weight was recorded prior to and post dehydration, the ratio of wet weight to dry weight gave
the water content of each cartilage sample.

1 **Biomechanical Testing**

2 *Compressive testing using an indenter:*

3 Methods used have been previously described by Pawaskar et al. [20] and Taylor et al. [21] to assess biomechanical properties of cartilage attached to bone. Briefly, osteochondral plugs 4 5 were compressed in a purpose built indentation rig using a 3 mm diameter, hemispherical, 6 stainless steel indenter under a load of 0.8 N. Plugs were submerged in PBS during testing to 7 maintain cartilage hydration. The deformation of cartilage was measured at a sampling 8 frequency of 5 Hz over one hour, after which all samples had reached equilibrium. Following 9 compression, pins were fully rehydrated in PBS before cartilage thickness was measured. A needle indenter was used to penetrate the cartilage, lowering at a rate of 4.5 mm.min⁻¹; the 10 11 resistance to motion was measured using a 500 N load cell [Instron 3365]. An increase in 12 load was recorded when the needle first contacted the cartilage surface and a second increase 13 when entering the bone, the distance between these two changes in load was taken as the 14 cartilage thickness. Deformation of cartilage was normalised to thickness to give percentage 15 deformation for each pin. The permeability and equilibrium elastic equilibrium modulus of 16 cartilage were derived from the cartilage thickness, deformation and water content using a 17 finite element (FE) method developed by Pawaskar et al. [20]. Briefly, an FE model of the 18 cartilage pin and indenter is produced, to which an initial permeability and equilibrium elastic 19 modulus are assigned. The FE model produces a deformation curve which is compared to the 20 experimental deformation curve. The input material properties are altered iteratively until the 21 modelled curve matches as closely as possible with the experimental curve. The accuracy of these modelled properties is assessed by the closeness-of-fit R² value for the curves, generally 22 23 acceptable over 0.6 for biological tissues.

1 Statistical analysis

Numerical data was analysed using Microsoft Excel. For each set of results the mean and
95% confidence limits were calculated. For data expressed as a ratio or percentage, values
underwent arcsine transformation before being statistically analysed, means and upper and
lower 95% confidence limits were then back transformed prior to presentation. Group means
were compared using the T-test following one-way or two-way (cell shape data) analysis of
variance (ANOVA). A p-value <0.05 was accepted as significant.

8 Results

9 Upon dissection, physical differences were observed between species. The length of the 10 femurs from femoral head to the front of the condyles was ~ 20 cm in both pigs and sheep; 11 however this was double in cows, ~ 40 cm. This size difference was reflected in the size of 12 the joints; the bovine femoral head measured ~ 8 cm from the terres ligament to femoral 13 neck, while the porcine was ~ 4 cm and ovine only ~ 2 cm. This pattern in joint size was seen 14 for both the hip and knee. Cartilage generally appeared white, glossy and healthy with 15 exception of ovine (> 4 yr) cartilage which had a yellowed appearance in most cases. Upon 16 drilling, the bone of porcine and young ovine was notably softer than that of bovine and older 17 ovine samples. Old ovine bone, despite being hard, was also found to be quite brittle when 18 cores were extracted from the joints.

19

H&E staining of osteochondral sections from each species showed characteristic chondrocyte
organisation (Figure 1), with linear columns of cells in the deep zone perpendicular to the
surface. This was less pronounced in porcine and young ovine, where the cell groups showed
a more rounded morphology.

1 Across most joint regions porcine and young ovine cartilage tended to have higher cellularity 2 than bovine and old ovine cartilage. This trend was observed within the superficial and deep 3 zones, but was less pronounced in the middle zone, where porcine cartilage showed lower 4 cellularity (Figure 2; p<0.05, ANOVA). There was no significant difference in the cell area 5 between species or joint region in the superficial or middle zones, cells of the deep zone 6 tended to be larger than those of the middle or superficial zones. Porcine cells were larger 7 than those of other species in the deep zone, significantly so in the acetabulum, femoral head, 8 femoral groove and medial condyle (Figure 3; p<0.05, ANOVA). Cells in the superficial and 9 middle cartilage zones in all species and joint regions tended to have a roundness value 10 around 0.75, however porcine cells in the deep zone were less round than those of other 11 species, significantly so in the acetabulum, femoral head, groove and medial condyle (Figure 12 4; p<0.05 ANOVA)

13

Staining of sections with alcian blue showed the distribution of GAGs within the cartilage tissues. All tissues showed characteristically less staining at the superficial zone becoming more intense through the middle and deep zones, then depleting in the calcified zone with the bone unstained.

18

Thickness of cartilage for each species was measured from digital images of the stained sections (Figure 5). Porcine tibial cartilage was significantly thinner than that of the groove and medial condyle, young ovine femoral head cartilage was significantly thinner than that of the medial tibia, no significant difference between joint areas was seen in bovine or old ovine (p<0.05 ANOVA), however there was a trend for cartilage of the femoral knee (groove and/or condyles) to be thicker than other joint areas.. Porcine and bovine cartilage was significantly thicker than ovine cartilage in a number of joint regions.

1

2	Following quantitative analysis it was found that porcine cartilage generally had significantly
3	higher GAG content than the other species, and old ovine cartilage generally had the lowest
4	(Figure 6). Across the joint regions a similar pattern was seen to thickness: femoral knee
5	cartilage tended to have higher GAG content. In the pig, cartilage of the groove and both
6	condyles had a significantly higher GAG content than that of the medial side of the tibial
7	plateau, only the medial condyle and groove showed higher concentration than the lateral
8	side. Cartilage from the medial condyle of the older sheep had a significantly higher GAG
9	content than that of the acetabulum, groove and tibia. Cartilage of the bovine lateral condyle
10	had significantly fewer GAGs than the medial tibia (p<0.05 ANOVA).
11	
12	The cartilage hydroxyproline content (representative of collagen) was quantified.
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between 70–82 % (Figure 8). There was no significant difference between any of the species
except for bovine, in which cartilage of the tibia had significantly higher water content than

24 the femoral head and medial groove, the lateral condyle also showed higher water content

than the femoral head. Across the species only bovine lateral tibial cartilage was significantly
higher in water content than old ovine (p<0.05 ANOVA).

3

The mechanical properties of cartilage from the different species and joint regions are shown in Figure 9 and Table (1), detailing the percentage deformation of cartilage after one hour, the permeability, elastic equilibrium modulus and thickness of each tissue. Properties could not be determined for ovine groove cartilage as the joints were too small to collect suitable pins.

9 There was a trend toward cartilage from the tibias and lateral condyles having higher 10 percentage deformation than cartilage of the hip. Porcine tibial cartilage showed significantly 11 greater percentage deformation than all other joint areas. Bovine femoral head cartilage, 12 medial condyle and medial tibia cartilage had significantly lower percentage deformation 13 than that of the lateral condyle and tibia. Young ovine acetabular cartilage deformed 14 significantly less than that of the femoral head. Cartilage of the old ovine acetabulum 15 deformed significantly less than that of the lateral condyle and tibial plateau. Old ovine 16 cartilage from the acetabulum deformed significantly less than other species. Young ovine 17 femoral head cartilage deformed significantly more than other species. Porcine lateral 18 condyle cartilage deformed significantly less than bovine and old ovine. Bovine medial tibial 19 cartilage deformed significantly less than other species (p<0.05 ANOVA).

20

Generally the equilibrium elastic modulus of porcine cartilage was lowest, increasing in
bovine, further increasing in young ovine cartilage and old ovine cartilage showed the highest
equilibrium elastic modulus in all cases. A trend was seen in all species across joint regions,
for cartilage of the tibia to have a lower modulus than areas of the femoral knee and hip.

25

Permeability showed an inverse relationship with equilibrium elastic modulus, with ovine
 being least permeable and porcine being most permeable. Permeability increased in the tibial
 plateau and was lowest in cartilage of the hip.

4

5 Discussion

6 As a starting point the xenogenic source material for decellularisation would be similar to 7 human cartilage in terms of composition, structure and mechanical properties. An additional 8 benefit could be to find a starting material with a higher GAG content than human cartilage, 9 as it is reported that decellularisation leads to a loss of GAGs within acellular scaffolds, and 10 therefore diminished mechanical properties [22-25]. This study looked at the biological, 11 biochemical and biomechanical properties of different osteochondral tissues. Various regions 12 of the porcine, bovine and ovine hip and knee were analysed to determine which would be 13 the best material for development of a decellularisation protocol.

14

From initial observations, cartilage of older sheep (>4 years) had a yellowed appearance, as seen in aged, osteoarthritic cartilage [26]. This would make it an inferior substitute for healthy young cartilage in vivo. The joint size has implications for the practicality of a species; more osteochondral samples can be harvested from a large bovine joint than porcine, and even fewer from an ovine joint. The apparent softness of porcine and to some extent the younger ovine bone highlighted the immaturity of these tissues.

21

H&E staining showed that cellular orientation was more defined in bovine and old ovine cartilage indicating a fully mature tissue. Cell shape analysis identified that the cells of the deep zones of porcine cartilage were larger and less round than those of the cartilage deep zones of other species, these observations of characteristically hypertrophic chondrocytes

1 suggested that the porcine tissues are immature and still undergoing growth. As in humans, 2 the cellularity of cartilage showed a trend of decreasing with age [27], with older ovine (> 4) 3 years) and bovine having fewest cells and porcine (6 months) having the most. The difference 4 in cellularity between joint regions in pigs related to the cartilage thickness. Cartilage from 5 the groove had significantly fewer cells than that of the tibia; however, tibial cartilage was 6 significantly thinner than that of the groove. Thicker cartilage has been reported to have 7 lower cell density than thin cartilage [28]. Comparison between cartilage zones indicated that 8 in thick porcine cartilages, there was higher cell density in the superficial and deep zones 9 compared the middle zone. For decellularisation, a starting material with fewer cells would 10 be preferable.

11

12 Cartilage thickness has been reported to increase as body weight increases [28]. This was not 13 necessarily supported in the present study. Although bovine and porcine cartilage were 14 thicker than that of ovine, despite the body mass of a cow being greater, porcine cartilage was 15 thicker. Julkunen et al. [29] showed that immature cartilage is thicker than that of mature 16 animals, and that thickness decreases with maturity. This was supported by comparison of the 17 two age groups of sheep, with the less mature ovine cartilage tending to be thicker than that 18 of the older sheep. Cartilage of the femoral condyles and groove showed a trend toward being 19 thicker than that of the hip and tibia in all species, this trend was also seen when comparing 20 the thicknesses for these joint regions in humans [30-32]. Simon et al. [33] have shown that 21 cartilage thickness is relative to joint congruence, so in joints with a high level of congruence, 22 such as the hip, cartilage is thinner. Human cartilage was on average thicker than that of any 23 species in this study, Shepherd et al. [30] report the thickness of human femoral head 24 cartilage to range between 1.08-2.40 mm where as ovine cartilage was only ~ 0.68 mm in this 25 area, porcine 1.14 mm and bovine 1.53 mm. Cartilage from the human groove ranged

between 1.76-2.59 mm, ovine cartilage is again much thinner; 0.86 mm in young sheep and 0.63 mm in old sheep. Cartilage from the porcine groove was 2.04 mm and bovine 1.52 mm, within or approaching the range of human cartilage thickness. The relative thickness of human cartilage probably reflects the increased loads placed on human knee cartilage due to the bipedal posture, as discussed by Stockwell [28]. It is likely that a decellularised xenogenic graft with cartilage of a similar thickness to the surrounding host tissue would integrate and function better, therefore porcine and bovine tissues would be preferred.

8

9 Porcine cartilage had the highest GAG content; this was likely due to the young age of the 10 tissue, as the concentration of proteoglycans is reportedly higher in immature cartilage, and 11 decreases with age [34]. Bolton et al. [35] suggest this is due to changes in chondrocyte 12 activity; aggrecan and link protein synthesis decreases with maturity. Except for cartilage of 13 the medial condyle, there was no significant difference in the GAG content of bovine and 14 ovine cartilage. Again, cartilage of the groove and condyles showed higher GAG content than 15 other joint regions. The relationship in these joint areas of increased GAG content alongside 16 increased thickness has been observed when tissues undergo increased loading or activity 17 [36]. However, forces acting on the healthy knee are reportedly lower than those of the hip 18 [37]. This trend across the joint regions may again be due to the congruence of the joints, 19 affecting the contact area and therefore stresses. Increased load on a very local scale may 20 cause specific remodelling at that site, to increase thickness and GAG concentration so to 21 decrease contact pressure. There is limited information in the literature regarding 22 concentrations of GAGs per dry weight of human cartilage. Various figures have been reported for cartilage of the femoral head, Vilim et al. [38] quote 54.7 and 38.6 mg.g⁻¹ GAG 23 24 per wet weight cartilage from two donors, while Hollander et al. [39] reported values ranging between $25 - 40 \,\mu\text{g.mg}^{-1}$ wet weight. In this study porcine femoral head cartilage was shown 25

to have $78 \pm 30 \ \mu g.mg^{-1}$ (n=5 ± 95 % CL) GAG per wet weight, bovine cartilage was 68 ± 8 µg.mg⁻¹, young ovine was $45 \pm 18 \ \mu g.mg^{-1}$ and old ovine was $62 \pm 16 \ \mu g.mg^{-1}$, all higher than values reported for human cartilage. GAG content was reported to be between 5-10 % total cartilage mass, as described for human cartilage by Mow *et al.* [14]. The comparatively high GAG content of cartilage found for all species indicates any could reasonably replace human cartilage, however, porcine having highest GAG concentration would be preferred as loss of GAGs is a limitation in decellularisation processes[22-25].

8

9 Hydroxyproline content showed an inverse relationship to GAG content. Porcine cartilage
10 had the lowest collagen content while old ovine had the highest. Muir *et al.* [40] assessed
11 collagen content of human cartilage and found great variation between individuals and joints.
12 They found no correlation between cartilage collagen content and age. This study only
13 quantified collagen content and did not look qualitatively at collagen fibre alignment,
14 although content was constant collagen orientation may differ.

15

On the whole, tissue water content did not vary significantly between species or joint area,
only bovine tibial cartilage showed significantly higher water content. The water content of
human cartilage varies between 68-85 % [14], the water content of animal tissues is currently
within the same range, although, this may differ once cartilage has undergone
decellularisation.

21

Within a species, percentage deformation of cartilage tended to be higher in the tibia and lower in the hip, this was reflected in the equilibrium elastic modulus derived for each type of tissue, with tibial cartilage showing a lower modulus. Permeability was also higher in the tibia than the hip as the GAG content of these tissues was lower, increased water flow

through the tissue would mean that load support from the cartilage fluid phase was less in
these tissues, therefore, the tissue deformed more. The relationship between material
properties and biological/biochemical composition was most pronounced in the bovine knee;
cartilage of the medial condyle and medial tibia was thicker and had higher GAG content
than its lateral counterpart, cartilage of the medial side of each joint also showed lower
permeability, and stiffer equilibrium elastic modulus (lower percentage deformation).

7

8 Between species, old ovine cartilage was stiffest and porcine was the least stiff. Old ovine 9 was stiffer than young ovine despite young ovine having higher GAG content, bovine was 10 also stiffer than porcine, despite porcine having a higher GAG content. This was likely due to 11 the quality of GAGs within the tissue, mature bovine and ovine cartilage may have more 12 effective proteoglycan structures which are better immobilised in the ECM compared to 13 porcine and young ovine, therefore these mature tissues had lower permeability and stiffer 14 cartilage. Old ovine cartilage was less permeable and stiffer than bovine cartilage, which may 15 be a result of further GAG maturity, or an effect of ovine cartilage being much thinner than 16 bovine. In this study, thinner cartilage tended to have a stiffer equilibrium elastic modulus.

17

The equilibrium elastic modulus of bovine lateral condyle cartilage was similar to the equilibrium aggregate modulus quoted for human (0.70 MPa) and bovine (0.89 MPa) previously [14, 31]. Porcine cartilage also had a similar elastic modulus, but was lower; this was possibly as a result of having softer subchondral bone. Old ovine cartilage had a much higher elastic modulus, so may be too stiff to replace human cartilage. Cartilage from all species was less permeable than values reported for human cartilage [14, 31].

1 Conclusion

2 A limitation of this study was that tissues from animals of different ages were compared; this 3 was due to the restricted availability of animals bred from the food chain for human 4 consumption. Cartilage properties were found to vary greatly between species and joint 5 region. For decellularisation, the optimal starting material should have properties as similar to 6 human cartilage as possible. Porcine cartilage from 6 month old animals was the thickest of 7 all the tissues studied and the GAG content was high, however the cartilage was highly 8 cellular and, did not have a fully defined structure due to the immaturity of the tissue. 9 Conversely bovine cartilage available from 18 month old animals was fully mature and less 10 cellular, the GAG content was lower than porcine cartilage, but in the range reported for 11 human. Bovine cartilage was also relatively thick. Both porcine and bovine cartilage had 12 equilibrium elastic moduli similar to the equilibrium aggregate modulus reported for human 13 cartilage. Ovine cartilage from 8-12 month and >4 year olds was very thin. Young ovine 14 cartilage, like porcine was not fully mature, with a higher cell density and less well defined 15 cellular distribution. Ovine cartilage again had a lower GAG content than porcine. 16 17 It has been found that porcine condylar cartilage and cartilage from the bovine groove and 18 condyle would make the most optimal starting materials on which to develop a

19 decellularisation protocol. Additionally, this study has highlighted the affect of species, age

20 and joint region on the physical properties of articular cartilage, allowing a more informed

21 selection of animal model for fundamental studies of cartilage biotribology.

1 Acknowledgements

This work was funded by the EPSRC and partially funded through WELMEC, a Centre for
Excellence in Medical Engineering funded by the Wellcome Trust and EPSRC, under grant
number WT 088908/Z/09/Z. J Fisher is an NIHR senior investigator. Additionally the
principle investigators J Fisher and E Ingham are supported in part by the NIHR LMBRU
Leeds Musculoskeletal Biomedical Research Unit and the ERC. The authors would like to
acknowledge Dr. Sainath Pawaskar for his assistance with the derivation of cartilage
mechanical properties.

9

10 Statement of Interests and declaration for healthcare compliance

E Ingham and J Fisher are academic founders of Tissue Regenix and are shareholders and
advisers to Tissue Regenix Group PLC.

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Equilibrium elastic modulus	Porcine		Bovine		Ovine (8 month)	8-12	Ovine (4 years)
(MPa)	Mean	\mathbf{R}^2	Mean	\mathbf{R}^2	Mean	\mathbf{R}^2	Mean	\mathbf{R}^2
(<i>MI u</i>)	1 30	0.00	Wiedii	K	1 60	0.08	1 3 2	0.01
Econoral hand	a1.12	0.99	^a 1 50	0.01	1.09	0.90	a2 52	0.91
Crease	1.15	0.97	0.96	0.91	1.22	0.94	2.32	0.95
Groove Maliata and the	1.04	0.95	0.80	0.94	-	-	b2 0.4	-
Medial condyle	⁵ 0.74	0.63	^{-1.1} /	0.80	1.31	0.99	2.04	0.89
Lateral condyle	0.66	0.67	0.81	0.81	1.42	0.98	2.02	0.86
Medial tibia	^b 0.88	0.86	^b 0.85	0.73	0.95	0.99	^b 2.02	0.87
Lateral tibia	^b 0.70	0.86	^b 0.81	0.85	1.19	1.00	^b 1.96	0.82
			Ovin	e (8-12				
Permeability, k	Porcine		Bovine		month)		Ovine (> 4 years)	
$x10^{-16}$ (m4/Ns)	Mean	\mathbf{R}^2	Mean	\mathbb{R}^2	Mean	\mathbf{R}^2	Mean	R^2
Acetabulum	3.04	0.99	-	-	1.12	0.98	0.37	0.91
Femoral head	5.70	0.97	1.78	0.91	2.99	0.94	0.76	0.95
Groove	4.49	0.93	6.33	0.94	-	-	-	-
Medial condyle	7.18	0.63	4.49	0.80	2.91	0.99	1.64	0.89
Lateral condyle	6.77	0.67	5.45	0.81	2.73	0.98	1.59	0.86
Medial tibia	8.09	0.86	4.52	0.73	3.67	0.99	1.30	0.87
Lateral tibia	10.29	0.86	5.75	0.85	2.87	1.00	1.37	0.82
					Ovin	le (8-12		
Thickness	Por	rcine	Bo	ovine	m	onth)	Ovine (> 4 years)
<i>(mm)</i>	Mean	95% CL	Mean	95% CL	Mean	95% CL	Mean	95% CL
Acetabulum	1.12	0.15	-	-	0.85	0.12	1.06	0.45
Femoral head	1.22	0.05	1.32	0.13	0.48	0.07	0.52	0.10

1.70

2.23

2.08

0.87

0.82

Groove

Medial condyle

Lateral condyle

Medial tibia

Lateral tibia

0.53

0.20

0.23

0.15

0.08

1.36

1.28

0.93

1.70

1.02

Table 1. Material properties of cartilage from various joint regions of the pig, cow and sheep.

Thickness data is expressed as the mean (ovine 8-12 months, acetabulum and groove n=5; all other n=6) \pm 95 % confidence limits, Elastic equilibrium modulus and permeability data was modelled on mean deformation data and R² values for modelled and experimental deformation curve agreement are presented. ^aFrom Taylor et al. [21], ^b from McLure et al. [41], all other data from the present study.

0.10

0.17

0.15

0.36

0.21

0.87

0.80

0.76

0.70

-

0.12

0.22

0.23

0.28

_

0.71

0.55

0.61

0.59

0.24

0.18

0.24

0.27

Figure Captions



Fig. 1. H&E staining of osteochondral pins from the medial condyle. A – pig, B- cow, C 8-12 month old sheep and D – > 4 year old sheep. Size bars are 500 μ m.



Fig. 2. Cellularity of superficial, middle and deep zone cartilage from various joint regions of the pig, cow and sheep. Data is expressed as the mean $(n=5) \pm 95$ % confidence limits. Data was analysed by two-way analysis of variance. This showed that for superficial zone cartilage, there was no significant variation in cell number between joint regions (p= 0.276). The effect of species on the cell number was dependent on the joint region (species x joint region interaction p= 2.74×10^{-8}). For the middle zone cartilage, there was no significant variation in cell number between joint region interaction p= 0.276. For the deep zone cartilage there was no significant variation in the cell number between joint region interaction p= 0.035). For the deep zone cartilage there was no significant variation in the cell number between joint regions but there was highly significant variation in the cell number between

species ($p=3.39 \times 10^{-12}$) with no significant interaction between species and joint region (p=0.384). Perusal of the data indicated that this was due to the higher cellularity of the porcine deep zone cartilage compared to the bovine and ovine cartilages.



Fig. 3. Size of chondrocytes in deep zone cartilage from various joint regions of the pig, cow and sheep. Data is expressed as the mean $(n=5) \pm 95$ % confidence limits. Data was analysed by two-way analysis of variance which revealed no significant variation in the size of the chondrocytes between joint regions (p=0.387). The effect of species on the size of the chondrocytes in the deep zone (p=1.13 x 10⁻²⁶) was dependent on joint region (species x joint region p= 9.47 x 10⁻⁵). Perusal of the data indicated that, although the chondrocytes in the deep zone of acetabular and femoral joint regions of the porcine cartilage were larger than in the ovine and bovine cartilages, this was not the case for the tibial cartilages.



Fig. 4. Roundness of chondrocytes in deep zone cartilage from various joint regions of the pig, cow and sheep. Data is expressed as the mean $(n=5) \pm 95$ % confidence limits. Data was analysed by two-way analysis of variance which revealed no significant variation in the roundness of the chondrocytes in the deep zone between joint regions (p=0.1) but a highly significant variation in the roundness of the chondrocytes in the deep zone between species (p=5.38 x 10⁻¹⁵) with no significant interaction between species and joint region (p=0.344). Perusal of the data indicated that this effect was due to the chondrocytes in the deep region of porcine cartilages being less round than the chondrocytes in the deep regions of bovine and ovine cartilages.



Fig. 5. Thickness of cartilage from various joint regions of the pig cow and sheep. Data is expressed as the mean $(n=5) \pm 95$ % confidence limits. Data for a given tissue site was

analysed by one-way analysis of variance and individual means compared by the T-method .

* indicates p<0.05.



Fig. 6. GAG content of cartilage from various joint regions of the pig cow and sheep.

Data is expressed as the mean $(n=5) \pm 95$ % confidence limits. Data for a given tissue site was analysed by one-way analysis of variance and individual means compared by the T-method . * indicates p<0.05.



Fig. 7. Hydroxyproline content of cartilage from various joint regions of the pig cow

and sheep. Data is expressed as the mean $(n=5) \pm 95$ % confidence limits. Data for a given tissue site was analysed by one-way analysis of variance and individual means compared by the T-method . * indicates p<0.05.



Fig. 8. Water content of cartilage from various joint regions of the pig cow and sheep.

Data is expressed as the back transformed mean $(n=5) \pm 95$ % confidence limits. Data for a given tissue site was arcsine transformed then analysed by one-way analysis of variance and individual means compared by the T-method . * indicates p<0.05.



Fig. 9. Percentage deformation of cartilage from various joint regions of the pig cow and sheep. Data is expressed as the back transformed mean (ovine 8-12 months, acetabulum and groove n=5; all other n=6) \pm 95 % confidence limits. Data for a given tissue site was arcsine transformed then analysed by one-way analysis of variance and individual means compared by the T-method . * indicates p<0.05. ^aFrom Taylor *et al.* [21], ^b from McLure *et al.* [41], all other data from the present study.