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Distinct horizontal patterns in the spatial organization of superficial zone chondrocytes of human joints

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Abstract

A better understanding of the unique cellular and functional properties of the superficial zone of articular cartilage may aid current strategies in tissue engineering which attempts a layered design for the repair of cartilage lesions to avert or postpone the onset of osteoarthritis. However, data pertaining to the cellular organization of non-degenerated superficial zone of articular cartilage is not available for most human joints. The present study analyzed the arrangement of chondrocytes of non-degenerated human joints (shoulder, elbow, knee, and ankle) by using fluorescence microscopy of the superficial zone in a top-down view. The resulting horizontal chondrocyte arrangements were tested for randomness, homogeneity or a significant grouping via point pattern analysis and were correlated with the joint type in which they occurred. The present study demonstrated that human superficial chondrocytes occurred in four distinct patterns of strings, clusters, pairs or single chondrocytes. Those patterns represented a significant grouping (p<0.001) with horizontal alignment. Each articular joint surface was dominated by only one of these four patterns (p<0.001). Specific patterns correlated with specific diarthrodial joint types (p<0.001). Further studies need to establish whether these organizational patterns are a consequence of their surrounding environment or whether they are linked to a functional purpose.

Keywords

Superficial zone; articular cartilage; human; chondrocyte; cluster; clustering; chondron; point pattern; knee; ankle; joint type; cell organization

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Introduction

Articular cartilage is an avascular, aneural, alymphatic, and viscoelastic connective tissue that functions hydrodynamically to bear loads and provide almost friction-free movement of diarthrodial joints (Kuettner et al., 1991). Its histomorphology is dominated by an extracellular matrix (ECM) consisting mainly of proteoglycan and type II collagen. Although cells of articular cartilage are represented by relatively sparsely distributed chondrocytes (Kuettner et al., 1991; Stockwell, 1971), articular chondrocytes and their pericellular microenvironment, termed chondrons, function as the primary metabolic units of articular cartilage to maintain matrix homeostasis via a synchronized balance between anabolism and catabolism (Kuettner et al., 1991; Poole, 1997).

Four major cartilage layers exist in articular cartilage: the superficial, transitional, radial, and calcified zones. Each can be distinguished based on differences in morphology (Aydelotte and Kuettner, 1988; Jadin et al., 2005; Kuettner et al., 1991), content of glycosaminoglycans (Venn and Maroudas, 1977), collagen (Nieminen et al., 2001), water (Maroudas and Venn, 1977), and mineral (Burr, 2004). The zonal differences in organization and biochemical content are directly associated with depth-dependant changes in biomechanical and bioelectrical tissue properties (Chen et al., 2001; Klein et al., 2007; Krishnan et al., 2003; Schinagl et al., 1997). This association is relevant both scientifically and clinically because current strategies for repair of traumatic cartilage lesions attempt to utilize the depth-dependent organization of articular cartilage (Klein et al., 2007). Additionally, engineering of layered tissues raises the hope to improve the wear characteristics of surrogate tissues and avert or postpone the frequent development of secondary osteoarthritis (OA) (Davis et al., 1989).

The superficial zone of articular cartilage in particular may be important for engineering of layered tissues due to the unique role that superficial chondrocytes play in joint lubrication (Jay et al., 2001; Schumacher et al., 1994), its biomechanical function as a tension resisting diaphragm (Meachim and Stockwell, 1979) and its distinctive proliferative and biosynthetic response to mechanical stimuli (Lee et al., 1998). Studies that were undertaken to further clarify the unique role of the superficial zone thus far have concentrated on cell shape (Hunziker et al., 2002; Jadin et al., 2005), cell density (Bywaters, 1937; Hunziker et al., 2002; Jadin et al., 2005; Quinn et al., 2005; Stockwell, 1971), cytoskeletal organization (Kim and Spector, 2000), growth-related changes in the bovine chondrocyte organization (Jadin et al., 2007), single cell mechanical properties (Jurvelin et al., 1996), gene expression (Darling et al., 2004; Jay et al., 2001; Khan et al., 2001; Schumacher et al., 1994), and metabolism (Eger et al., 2002). While it is widely accepted that the organization of human chondrocytes of the radial zones occur in vertical columns and the transitional zone in small "random" groups (Aydelotte and Kuettner, 1988; Brighton et al., 1984; Kuettner et al., 1991), it remains unclear how the human chondrocytes of the superficial zone are organized.

Studies investigating the cellular organization of the superficial zone of human articular cartilage are rare. Few groups have access to non-degenerative human donor tissue in which the superficial zone is still intact. Furthermore, despite an abundant availability of surgical specimens, most of these specimens are often degenerated and therefore have partially lost their superficial zone. However, organizational information on non-degenerative human tissue may prove vital to modern cartilage repair strategies that attempt to engineer the layered structure of articular cartilage. An additional potential problem is that conventional histological analyses on tissue sections are looked at in the vertical plane. When viewed vertically, superficial chondrocytes appear as single cells (Hunziker et al., 2002). However, we recently showed that chondrocytes within the superficial zone of normal human ankle cartilage were arranged in horizontal clusters parallel to the articular surface (Schumacher et al., 2002). Although this study suggests that human superficial chondrocytes, in a top-down view, may

appear in horizontally oriented groups, particular patterns in the superficial chondrocyte organization have not been investigated.

To further elucidate the spatial arrangement of superficial chondrocytes, the present study quantitatively analyzed fluorescent microscopic images of the entire intact superficial zone of four major human joints (shoulder, elbow, knee, and ankle) that were photographed from top-down views. Further vertical or horizontal tissue sectioning was avoided to prevent any potential loss of three-dimensional information. The resulting arrangements of the chondrocytes were tested for randomness, homogeneity or a significant grouping in the context of a point pattern analysis (Diggle et al., 2000). The presence of specific chondrocyte patterns were also correlated with the joint type in which they occurred.

Methods

Human Articular Cartilage

Articular cartilages from joints of the upper (elbow and shoulder joints) and lower (knee and ankle joints) extremities were obtained with institutional approval from healthy adult human tissue donors within 24 hours after death through the Gift of Hope Organ and Tissue Donor Network (Illinois, USA) and graded according to a 5-point scale by a modified Collins grading (Muehleman et al., 1997), using the following criteria: grade 0 (normal cartilage without signs of degeneration), grade 1 (minor surface roughening), grade 2 (fibrillations and fissuring), grade 3 (full defects covering less than 30% of the articular surface), and grade 4 (full defects covering more than 30% of the articular surface). Only joints with the grades 0–1 were included into this study. No other pathology other than articular degeneration was observed in any of the joints.

Tissues from eight human donors (age 65.9 ± 5.4 years, range 52-85 years) were obtained from the proximal humerus (center of the humeral head articular surface facing the glenoid, n = 6, grade 0: n = 4, grade 1: n = 2), distal humerus (capitulum center, n = 4, all grade 0), ulna (semilunar notch, convex medial portion, n = 4, all grade 0), radius (radial head, concave fossa, n = 4, all grade 0), distal femur (condyles, weight bearing region in knee joint extension; patellofemoral groove, medial part, n = 7, grade 0: n = 5, grade 1: n = 2), proximal tibia (edge of the medial plateau, that is covered by the medial meniscus body, n = 5, grade 0: n = 3, grade 1: n = 2), distal tibia (center, n = 4, all grade 0), and talus (dome center, n = 8, grade 0: n = 5, grade 1: n = 3).

Fluorescence Microscopy

In order to visualize the arrangement of the superficial chondrocytes through the depth of the superficial zone cartilage, samples of approximately 200 μ m thickness and 1 cm² area were obtained by manual horizontal dissection from the weight-bearing area of each joint surface and not further sectioned to preserve intact chondrocyte patterns. The orientation of the cartilage slices was maintained by cutting one border of the sample slightly angulated producing an arrowhead shape. The samples were directly fixed in 4% paraformaldehyde and stained as previously described (Schumacher et al., 2002). Briefly, samples were washed with 1% Triton X-100 in phosphate buffered saline (PBS) and treated with 5% aqueous CaCl₂ for 30 minutes. All samples were stained for 60 minutes with propidium iodine (10 μ g/ml in PBS) and washed for 20 minutes with Tris buffered saline (1 M Tris, 15 M NaCl, pH 7.5). The samples were viewed perpendicular to the articular surface, providing a top-down view. Two-dimensional images were digitally recorded for each articulating surface (Nikon Eclipse TE200, Melles-Griot 43 series ion laser) at the sample center as a single topographical location beneath the surface when cells first came into focus and at successive depths when further cells came into focus within the superficial zone.

Quantitative Image Analysis

Images of the superficial zone were imported into Adobe Photoshop (San Jose, CA). The images were digitally filtered by using the contour threshold function of Fovea Pro in Photoshop (Reindeer Graphics, Asheville, NC) to exclude cells that were out of focus and therefore in different planes. Based on proximity, the depicted nuclei were then grouped numerically into single cells or groups of 2, 3, 4, 5, 6 to 8, or 9 to 12. Based on their spatial arrangement, groups were classified as singles, pairs, clusters, and strings. Singles were defined as single cells, pairs as two neighboring chondrocytes, clusters as a circular arrangement of chondrocytes, and strings as chondrocytes that were aligned in a line.

Point Pattern Analysis

To determine if the spatial arrangement of cells was random, homogeneous or significantly grouped in the context of a point pattern analysis (Diggle et al., 2000), the nearest neighbor method was used (Skellam, 1951) for which grouped data was defined as a sub-collection of data that is proximate based on distance (Rousseeuw, 1987). Briefly, nuclear centroids (one pixel marking the geometrical nucleus center) were determined with Fovea Pro in Photoshop (Reindeer Graphics, Asheville, NC), and the mean nearest neighbor distance between the centroids of two neighboring chondrocytes was calculated. The ratio of observed values of the distance to the nearest neighbor and their expected values in an assumed homogeneous distribution was used to determine the presence or absence of grouping. To quantify the departure from random, the appropriate test of significance for this method was used (Clark and Evans, 1954).

Correlation of Chondrocyte Patterns with Diarthrodial Joint Types

To test the correlation between specific chondrocyte patterns and the particular diarthrodial joint type (Gray, 1995) in which they occurred, the articular surfaces were coded according to their joint function (1 = uni-axial hinge joint [talus, distal tibia]; 2 = bi-axial joint with a secondary rotational axis [condyles, humeral head, distal humerus capitulum]; and 3 = concave surface [patellofemoral groove, radial head]).

Cell Density and Split Lines

To separately analyze the cell densities of the superficial and deeper zones, the entire articular surface was dissected to a depth of 200 µm and defined as the superficial zone. The remaining non-calcified cartilage was dissected and defined as deeper zones. The wet weights of both the superficial and deeper zones were determined. The chondrocytes were isolated using an established method (Aydelotte and Kuettner, 1988). Briefly, the superficial and deeper zones were digested in 0.2% pronase (Calbiochem, La Jolla, CA) in Dulbecco's Modified Eagle's Medium (DMEM) containing 10 % fetal bovine serum (FBS; HyClone, Logan, UT) for 90 min at 37°C, followed by overnight digestion with 0.025% collagenase-P (Boehringer Mannheim, Indianapolis, IN) in DMEM containing 10 % FBS.

The cell numbers were determined for each joint surface and for the superficial and deeper zones by averaging three manual counts of the released chondrocytes. Before counting, the complete digestion of tissues was ensured microscopically for each group. The cell densities were calculated as the number of released chondrocytes per gram wet weight of the dissected cartilages. It has been suggested that the yield of released cells, especially from human cartilage, is relatively low and perhaps depends on the amount of damage near sliced edges. Due to methodological differences, direct comparisons with other studies are difficult. However, to compare our results to previous studies, we calculated the ratio between the densities of the superficial zone and the deeper zones and found that it was comparable to those of other studies (Hunziker et al., 2002; Quinn et al., 2005; Temple et al., 2007).

On selected distal femora and human tali, cartilage were removed manually from the weight bearing areas of the articular surfaces, pinpricked and stained with Indian ink to reveal splitline patterns as previously described (Below et al., 2002; Meachim et al., 1974).

Statistical Analysis

All data is presented as mean \pm SEM (standard error of the mean) of *n*. All data were subjected to one way ANOVA or ANOVA on Ranks, followed by Holm-Sidak or Dunn's Test post-hoctests depending on normal or non-normal data distribution, respectively. A Spearman Rank Order Correlation was performed to test for significant correlations between the percentage of cells that were situated in a specific pattern and the diarthrodial joint type of the articular surface in which this particular pattern occurred. Differences were considered significant at $p \le 0.05$. All tests were performed using SigmaStat 3.1 software (Systat Software Inc., San Jose, CA, USA).

Results

Fluorescence Microscopy

To visualize the spatial arrangement of chondrocytes of the superficial zone, fluorescent microscopic images of the entire depth of the uncut, intact superficial cartilage of human shoulder, elbow, knee, and ankle joints were photographed from top-down views. Chondrocytes of the superficial zone were neither homogeneously nor randomly distributed. As shown in Figure 1, they were organized into groups that were aligned parallel to the articular surface. Throughout the depth of the superficial zone, chondrocyte groups were situated in distinctly different focal planes. With increasing tissue depth, chondrocyte groups came in and out of focus and were separated from each other by extracellular matrix both vertically and horizontally. Superficial chondrocytes were organized in four distinctly different spatial patterns that we classified as strings, containing three or more cells in a straight line (Figure 2, A and B), clusters containing three or more cells in a circle or ellipse (Figure 2, C and D), pairs containing two cells in close proximity of one another (Figure 2, E and F), or single chondrocytes that were not in proximity to one another. The chondrocyte patterns were not aligned with the orientation of the cartilage sample within the joint surface.

Quantitative Image Analysis

Although in all articular surfaces four chondrocyte patterns such as strings, clusters, pairs, and singles were found, most articular joint surfaces were dominated by only one of these four patterns. Most chondrocytes of the articular surfaces of the ankle joint were situated in pairs; $67.08 \pm 1.96\%$ of talus dome chondrocytes and $62.14 \pm 1.32\%$ of distal tibia chondrocytes were paired (Figure 3). In contrast, strings of chondrocytes were found in half of the superficial chondrocytes of the femoral condyles of the knee joint $(51.25 \pm 4.94 \%)$, whereas circular clusters were observed in two thirds of the chondrocytes in the patellofemoral groove of the femur (68.72 \pm 5.46 %). In the third articulating surface of the knee, the meniscus-covered medial tibia plateau, we found that $44.5 \pm 3.81\%$ of the chondrocytes were paired and $7.0 \pm$ 2.1% had formed strings. In the shoulder and elbow joints, all articular surfaces were also dominated by a particular cellular pattern. Thus, almost half of all cells of the distal humerus were located in strings (42.83 ± 6.14 %), whereas more than two thirds of cells of the radial head were positioned in clusters (79.24 \pm 1.02%). Half of the superficial chondrocytes of the humeral head (47.05 \pm 6.33 %) and almost two thirds of the cells of the proximal ulna (64.11 \pm 6.7 %) were situated in pairs. Overall, the predominant pattern varied across joints, and, depending on the surface, consisted either of pairs, clusters, or strings but not single chondrocytes. Furthermore, the percentage of cells situated in the predominant cell pattern of each given articular surface was consistently and significantly higher than the percentage of cells in the remaining, non-predominant cell patterns (p < 0.001). Significant differences

between the various examined anatomical locations are summarized in Table I. In addition, we looked at the femoral border between condyles and patellofemoral groove and noticed a transitional zone, in which the predominant patterns of the two bordering regions were intertwined. However, we did not further quantify those images.

In some articular surfaces there was, in addition to a predominant pattern, a predominant number of superficial chondrocytes per group (Figure 4). In the patellofemoral groove, the groups consisted predominantly of 6 to 8 chondrocytes (25.12 ± 9.19 %), whereas the superficial zones of the talar dome and distal tibia consisted largely of groups of two cells $(48.72 \pm 2.47 \%$ and $56.97 \pm 2.34 \%$ of cells were paired). In other articular surfaces such as the femoral condyles, there was no predominant number of chondrocytes per group (Figure 4). These findings were significantly different from those of the patellofemoral groove (p < 0.008) or the talus (p < 0.001). In the upper extremity, two predominant cell number groupings were found. Within the articular surface of the radial head, 35.71 ± 6.07 % of chondrocytes were situated in groups of five cells, whereas in the proximal ulna, the chondrocytes were primarily grouped in pairs (63.92 ± 3.72 %, p < 0.001). While single cells were never the predominant organization, the highest percentages of single cells were found in the proximal humerus and the distal tibia (Figure 4). The lowest percentage of single cells were found in the distal humerus $(5.94 \pm 1.62\%)$ and the radial head $(2.38 \pm 1.86\%)$. Significantly higher numbers of superficial chondrocytes per cell group were also found when data of the upper extremity was compared with data of the lower extremity (p = 0.002) and when the knee joint was compared with the ankle joint (p < 0.001). No differences in cell numbers per group between shoulder and elbow joints were noted.

Point Pattern Analysis

To test if the spatial arrangement of cells was random, homogeneous or grouped, a point pattern analysis (Diggle et al., 2000) was performed by calculating the ratio (R) of observed values of the distance to the nearest neighbor and their expected values in an assumed homogeneous distribution. Briefly, a ratio = 1 describes a random distribution, a ratio < 1 describes a grouped distribution, while a ratio > 1 defines a homogeneous distribution. The ratio (R) was R = 0.794 for chondrocytes forming strings, R = 0.625 for clusters, and R = 0.836 for pairs, strongly indicating a significantly grouped and not a random or homogeneous cellular distribution within normal cartilage (strings, p < 0.001, clusters, p < 0.0001, pairs, p < 0.05).

To determine differences in the proximity of chondrocytes, the mean nearest neighbor distances of superficial chondrocytes were calculated. Those of knee cartilages were significantly smaller than those of the ankle joint (p < 0.01). This finding was based on the mean nearest neighbor distances between cells of the condyles and the talus (p < 0.05) but not on the patellofemoral groove and the talus, suggesting a closer proximity of condylar than talar superficial cells. No differences were noted when comparing the mean nearest neighbor distances of superficial chondrocytes of the two ankle joint surfaces (talus and tibia) or those of the upper with those of the lower extremity or of the shoulder compared with those of the elbow joint (data not shown).

Correlation of Chondrocyte Patterns with Joint Types

To determine if there was a significant correlation between specific chondrocyte patterns and the joint type of the articular surface in which they were situated, the articular surfaces were coded as convex surfaces of a uni-axial hinge joint or biaxial joint with a secondary rotational axis, or as concave surfaces. A Spearman Rank Order Correlation was performed, which revealed significant correlations between single chondrocytes and a uni-axial hinge joint type (p < 0.001, correlation coefficient = -0.829), between paired chondrocytes and a uni-axial hinge joint type (p < 0.001, correlation coefficient = -0.828), and between clusters and a

concave articular surface shape (p < 0.001, correlation coefficient = 0.831). For chondrocyte strings, significant correlations were not found. However, the percentage of cells that were situated in strings of a joint type with a rotational axis was significantly higher than the percentage of cells that was located in strings of joints without a rotational axis (p < 0.001).

Cell Density and Split Lines

To determine if differences in the numerical, spatial and proximity based chondrocyte organization may or may not affect the bulk zonal cell density, for each surface the cell densities were analyzed. Interestingly, our results showed no differences between the superficial densities of the different articular surfaces that were analyzed (Figure 5), suggesting that the observed spatial, numerical and proximity based differences were local occurrences whose presence did not affect the bulk cell density. As shown previously for articular cartilage (Hunziker et al., 2002;Quinn et al., 2005;Stockwell, 1971), all cartilages of the knee and ankle joints showed significantly higher cell densities in their superficial zones than in their corresponding deeper zones (p < 0.05). In the shoulder and elbow joints, a trend to higher superficial than deeper zones cell densities was observed.

Split line patterns are well described in the distal femur and talar dome (Below et al., 2002; Meachim et al., 1974), however, their significance is not entirely clear. While some authors assume a correlation between split lines and collagen fibril orientation, ultrastructural studies revealed little evidence that the split-line direction correlated with any preferred alignment of fibrils (Kamalanathan and Broom, 1993). We observed split line patterns of the distal femur and the talar dome similar to those reported (Below et al., 2002; Meachim et al., 1974). However, we were unable to establish any relationship between the split lines and superficial chondrocyte groups, which were not aligned in any identifiable pattern with neighboring or distant cell groups (data not shown).

Discussion

The present study compared the organizational structure of superficial chondrocytes in nondegenerated articular cartilages obtained from different human joints in the upper and lower extremities. Perhaps the most important observation of the present study was that superficial chondrocytes, when viewed in a top-down view, were grouped in patterns that were oriented parallel to the articular surface. We identified four distinctly different patterns: strings, clusters, pairs and single chondrocytes. The observed patterns were not due to a random distribution of chondrocytes but rather represented a significant grouping of cells in the context of a statistical analysis of point patterns. Although a few studies of the superficial zone reported clusters in human cartilage (Schumacher et al., 2002), pairs in bovine cartilage (Jadin et al., 2005) and incidentally symmetrical rows in rabbit cartilage (Clark and Rudd, 1991), to the author's knowledge, the present study is the first to identify complex patterns in the arrangement of human chondrocyte groups in the superficial zone of non-degenerated cartilage.

That chondrocytes of the middle and deep zones can be grouped in single, double or multiple chondron columns with interconnecting segments between adjacent chondrons has been reported for non-degenerative tibial cartilage (Poole, 1997). It has been suggested that chondrocyte groups may have formed during skeletal development (Morrison et al., 2000), and their pairing may have arisen after cell division followed by an incomplete separation within the chondron microenvironment (Chi et al., 2004). The position of daughter cells can be determined by positioning of the cell division axis and is influenced by the spatial distribution of the ECM (Thery et al., 2005). In fact, an electron microscopy study of the articular surface reported associations between ECM surface depressions and superficial cell groups in rabbit cartilage (Clark and Rudd, 1991). In addition, the local zonal collagen architecture appear to influence the chondron shape and orientation (Youn et al., 2006). Thus, specific patterns in the

organization of superficial chondrocytes may be associated with the local articular surface structure.

In fact, data from the present study suggests that a specific chondrocyte pattern is correlated with the diarthrodial joint type of the articular surface in which they occur. All articular surfaces, which were predominated by chondrocyte pairs, such as both surfaces of the ankle joint and the humero-ulnar compartment of the elbow joint, function mainly as an uni-axial hinge joint (Gray, 2004). The surfaces in which chondrocyte strings were predominantly observed (the femoral condyles of the knee joint and the capitulum of the distal humerus) function in bi-axial joints with a secondary rotational axis (Gray, 2004). To a lesser extent, strings were found in the humeral head, which is part of a ball-and-socket joint that also allows rotatory movements (Gray, 2004). Surfaces with a concave anatomy were associated with chondrocyte clusters. Concave surfaces differ from a convex anatomy in terms of joint functioning; when a concave surface moves on a convex surface, rolling and sliding occur in the same direction (Gray, 2004), but when a convex surface moves on a concave surface, rolling and sliding movements occur in opposite directions (Gray, 2004). Clusters were found in the shallow concave patellofemoral groove of the distal femur and the concave depression of the radial head. Although we found significant correlations between chondrocyte patterns and diarthrodial joint types, further studies will be required to determine if specific chondrocyte patterns are causally linked to specific biomechanical functions.

The present study showed numerical, spatial and proximity based differences in the superficial cellular organization when comparing the articular surfaces of different joint types. Numerical differences may occur with alterations in the cellularity of human articular cartilage due to aging. Those age-dependent alterations occur in joints such as the hip and knee, but are less likely to occur in the humeral head (Meachim and Collins, 1962; Stockwell, 1967; Vignon et al., 1976). If present, the principal age-related changes are seen in the superficial zone (Stockwell, 1978). Therefore, it may be possible that age-related changes in the cellularity affect the observed superficial chondrocyte patterns, possibly by changing the pattern type or by decreasing the presence of a certain pattern. However, in the present study we did not observe such age-related changes, perhaps because young donors were not included in the present study. Our numerical, spatial and proximity based data may be relevant to tissue engineering with different seeding densities of chondrocytes that are used to increase material properties such as compressive and dynamic stiffness (Mauck et al., 2003). However, our data showed that the cell densities of the superficial zones were similar when comparing the articular surfaces examined. Thus, the observed differences in the numerical, spatial and proximity based chondrocyte organization did not affect the zonal cell density and appear to be local variations within the chondrocyte microcosm.

The question arises whether the presented cellular orientation was dependent on collagen fiber orientation or on the established split line pattern found in the distal femur and talar dome (Below et al., 2002; Meachim et al., 1974). We were unable to show a split line and cellular correlation. Because we examined only the cellular, but not the collagen organization, the question concerning the alignment of collagen and cells cannot be answered directly. However, an electron microscopy study showed that rows of rabbit chondrocytes run parallel to adjacent collagen fibrils within the same superficial plane (Clark and Rudd, 1991), suggesting a spatial relationship between collagen fibers and the chondrocyte organization. An ultrastructural study revealed little evidence that the split-line direction correlated with any preferred alignment of collagen fibrils (Kamalanathan and Broom, 1993). Our data excluded a close association between the presented chondrocyte orientation and the split lines of the femur and tibia.

Biomechanical forces and their mechanotransduction are an important modulator in the development (Heegaard et al., 1999; Wong and Carter, 2003), differentiation (Brama et al.,

2000; Little and Ghosh, 1997), and maintenance of articular cartilage (Grodzinsky et al., 2000; Lane Smith et al., 2000; Smith et al., 1995). During skeletal development, the morphogenesis of articular cartilage is regulated by mechanical forces, resulting in the formation of the four cartilage zones (Heegaard et al., 1999; Wong and Carter, 2003). The biochemical content of these zones is further modulated by a functional adaptation of the extracellular matrix to weight bearing (Brama et al., 2000). Mechanical forces also modulate the chondrocyte phenotypic expression during development (Little and Ghosh, 1997). In adulthood, mechanical loading directly influences cartilage homeostasis, which is important for maintaining a functional matrix (Lane Smith et al., 2000; Smith et al., 1995). Biomechanical forces play a major role in the genesis of articular cartilage zonal heterogeneity. They may also contribute to the unique superficial zone heterogeneities in the organizational patterns of human chondrocytes across different joint types as observed in the present study. Although a functional explanation for these organizational chondrocyte patterns has not yet been established, their correlation with a specific joint type may suggest an association with the occurring biomechanical forces and perhaps a role in mechano-sensing and transduction. Additional bovine and human articular cartilage studies are ongoing in our laboratory to further elucidate a possible metabolic or biomechanical role of the superficial chondrocyte organization in articular functioning and repair. Although the hypothesis of such roles is clearly intriguing, it remains to be established whether these organizational patterns are a passive consequence of their surrounding anatomical and biomechanical environment or whether they actively serve a functional purpose. The present study has uncovered a new dimension of complexity to an already intricate tissue that has not been appreciated previously.

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Figure 1. Different focal planes of patellofemoral groove clusters

Representative images of the most superficial (A, approximately a few micrometers beneath the tissue surface) and increased tissue depths (B–D, approximately 30 micrometer steps deeper into the tissue) within the superficial zone. The square, circle and triangle represent different groups of cells situated in different focal planes. Scale bar, $100 \,\mu$ m.



Figure 2. Chondrocyte cell patterns of the superficial zone

Fluorescent microscopic representative images showing cells stained with propidium iodide of strings (A–B), clusters (C–D) and pairs (E–F). Single cells were dispersed among the other patterns. Scale bar, 100 µm.



Figure 3. Percentage of lower and upper extremity chondrocytes that were grouped in strings, clusters, pairs or as singles

Values are expressed as the mean \pm standard error of the mean (SEM). p values are listed in Table I.



Figure 4. Percentage of chondrocytes that appear as single cells or in groups of 2, 3, 4, 5, 6 to 8, or 9 to 12 cells

Values are expressed as the mean \pm SEM.





Table I

Significant differences between articular surfaces

Pattern	Comparison	р
Strings	Condyles vs. Patellofemoral Groove	<0.001
	Condyles vs. Talar Dome	< 0.001
	Condyles vs. Proximal Tibia	< 0.002
	Condyles vs. Humeral Head	< 0.001
	Condyles vs. Semilunar Notch	< 0.001
	Condyles vs. Radial Head	< 0.001
	Distal Humeral vs. Humeral Head	< 0.002
	Distal Humerus vs. Semilunar Notch	< 0.002
	Distal Humerus vs. Radial Head	< 0.001
	Distal Humerus vs. Patellofemoral Groove	< 0.001
	Distal Humerus vs. Talar Dome	< 0.001
	Distal Humerus vs. Proximal Tibia	< 0.002
Cluster	Patellofemoral Groove vs. Condyles	< 0.001
	Patellofemoral Groove vs. Talar Dome	< 0.001
	Patellofemoral Groove vs. Humeral Head	< 0.002
	Patellofemoral Groove vs. Semilunar Notch	< 0.001
	Patellofemoral Groove vs. Distal Humerus	< 0.002
	Radial Head vs. Talar Dome	< 0.001
	Radial Head vs. Condyles	< 0.001
	Radial Head vs. Humeral Head	< 0.001
	Radial Head vs. Semilunar Notch	< 0.001
	Radial Head vs. Distal Humerus	< 0.002
	Proximal Tibia vs. Talar Dome	< 0.001
	Proximal Tibia vs. Semilunar Notch	< 0.001
	Proximal Tibia vs. Condyles	< 0.001
	Proximal Tibia vs. Humeral Head	< 0.002
	Proximal Tibia vs. Distal Humerus	< 0.002
Pairs	Talar Dome vs. Condyles	< 0.001
	Talar Dome vs. Patellofemoral Groove	< 0.001
	Talar Dome vs. Proximal Tibia	< 0.001
	Talar Dome vs. Radial Head	< 0.001
	Talar Dome vs. Distal Humerus	< 0.001
	Semilunar Notch vs. Condyles	< 0.002
	Semilunar Notch vs. Patellofemoral Groove	< 0.002
	Semilunar Notch vs. Proximal Tibia	< 0.001
	Semilunar Notch vs. Radial Head	< 0.001
	Semilunar Notch vs. Distal Humerus	< 0.002
	Humeral Head vs. Proximal Tibia	< 0.002
	Humeral Head vs. Radial Head	< 0.002
Singles	Talar Dome vs. Condyles	< 0.05
	Distal Tibia vs. Patallofamoral Groova	<0.05