

# Upregulation of insulin-like growth factor I gene expression in the lesions of osteoarthritic human articular cartilage

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## Abstract

**Expression of insulin-like growth factor I (IGF-I) mRNA and IGF-I protein was studied in human osteoarthritic and young articular cartilage by in situ hybridisation and immunohistochemistry. In situ hybridisation showed that relatively low amounts of IGF-I mRNA signal were present in anatomically normal regions of osteoarthritic and young cartilage. In fibrillated osteoarthritic cartilage, however, the signal intensity was significantly higher than in non-fibrillated cartilage. Particularly high levels of IGF-I mRNA were present in the surface cell clones of more advanced lesions, the amount of signal being about four to five times greater than in anatomically normal cartilage. The amount of message varied with cartilage depth. In young cartilage there was less IGF-I mRNA in the superficial zone than in the middle and deep zones. In fibrillated regions of osteoarthritic joints the amount of message in surface cells was greater than in deeper regions. A specific human IGF-I antibody was used to show the presence intracellularly of IGF-I protein in osteoarthritic and young cartilage.**

**Raised levels of IGF-I message in osteoarthritic chondrocytes may represent an attempt at increased matrix repair, operating by an autocrine/paracrine mechanism.**

Articular cartilage consists of chondrocytes embedded in an extensive extracellular matrix. In normal tissue the chondrocytes actively effect a stable equilibrium between the synthesis and degradation of matrix components, so that a constant concentration of these components is maintained. In degenerative joint disease, such as osteoarthritis, the stable equilibrium is disrupted and the rate of loss of proteoglycan from the matrix eventually exceeds the rate of deposition of newly synthesised molecules.<sup>1 2</sup> It is of interest, therefore, to understand the influence of growth factors on this equilibrium.

Insulin-like growth factors (IGFs) or somatomedins are polypeptides with structural and functional homology to proinsulin.<sup>3</sup> In man, two major IGFs, IGF-I (somatomedin C) and IGF-II, have been characterised by amino acid sequence analysis of peptides purified from serum<sup>4 5</sup> and by isolation of cDNAs encoding precursor forms of IGFs.<sup>6 7</sup> Insulin-like growth factors play a part in the growth and development of many tissues during fetal and adult life,<sup>3 8</sup> and are implicated in tissue hypertrophy and repair.<sup>9</sup> Insulin-like growth factor I is known to enhance matrix synthesis in normal

cartilage in vivo<sup>10</sup> and in vitro.<sup>11 12</sup> In addition, specific IGF-I receptors have been demonstrated on rabbit and bovine articular chondrocytes.<sup>13 14</sup>

Traditionally, IGF-I was considered to be produced primarily in the liver in response to a systemic signal and transported by the circulation to act on target tissues at distant sites (endocrine function).<sup>15</sup> There is now evidence, however, that IGF-I is synthesised by cells of mesenchymal origin in many different tissues. Immunoreactive IGF-I protein has been detected in extracts of human<sup>16</sup> and rat organs.<sup>17</sup> mRNA encoding IGF-I can be detected in a number of fetal and adult rat tissues<sup>18</sup> and human fetal tissues.<sup>19</sup> The concentrations of immunoreactive IGF-II and abundance of IGF-I mRNA are regulated by growth hormone in a variety of adult rat organs.<sup>17 20</sup> Insulin-like growth factor I is secreted by cultured explants of rodent tissues<sup>21 22</sup> and by monolayers of human fibroblasts and osteoblast-like cells.<sup>23 24</sup> These data suggest that IGFs can act locally by autocrine or paracrine mechanisms, or both.

The aim of our study was to determine whether the IGF-I gene is expressed by human articular chondrocytes. We asked the questions: Does articular cartilage have the potential to synthesise a growth factor that may be involved in matrix repair? Could chondrocytes respond to progressive matrix loss, as occurs in osteoarthritis, by producing more growth factor?

## Materials and methods

### TISSUE SOURCE AND PREPARATION

Twenty six blocks of osteoarthritic femoral articular cartilage were taken from seven subjects, aged 64-77 years, at total joint replacement operations. Osteoarthritic blocks were sampled from the superior surface of femoral heads adjoining regions of total cartilage loss and from the inferior surface that was macroscopically intact. Nine anatomically normal samples from the weightbearing regions of femoral articular cartilage were obtained from young subjects, aged 12-24 years, at osteosarcoma operations. Blocks were embedded in Tissue Tek II OCT compound (Miles Laboratories, Naparavill, IL, USA), frozen in liquid nitrogen, and stored at -70°C. The time between surgical operation and freezing of the tissue was not more than six hours.

### PREPARATION OF PROBES

In situ hybridisation was performed using in vitro synthesised RNA labelled with [<sup>35</sup>S]uridine triphosphate. Human cDNA encoding IGF-I was developed by Jansen *et al*<sup>6</sup> (Department of

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Paediatrics, State University of Utrecht, the Netherlands). This sequence encodes the entire IGF-I precursor, and is complementary to human IGF-I mRNA on northern blots with multiple transcript sizes of 0.7, 5.3, and 8.0 kilobases.<sup>25</sup> Insulin-like growth factor I cDNA (560 base pairs) was subcloned in opposite orientations into pSP64 and pSP65 vectors (Promega Biotec, Madison, WI, USA). After linearisation of the plasmids, antisense and control sense <sup>35</sup>S labelled RNA transcripts were synthesised using SP6 promoters. The template DNA was removed by RNase-free DNase I (Pharmacia, Freiburg, FRG), and the probes were purified by centrifugation chromatography

(Sephadex-G50; Pharmacia).<sup>26</sup> Full length transcripts were degraded to 50–150 bases by limited alkaline hydrolysis (40 mM NaHCO<sub>3</sub>, 60 mM Na<sub>2</sub>CO<sub>3</sub> (pH 10.2), 75 minutes, 60°C) for optimal hybridisation efficiency.<sup>27</sup>

IN SITU HYBRIDISATION

These procedures were modifications of protocols described by Brahic and Haase,<sup>28</sup> Cox *et al.*,<sup>27</sup> and Funa *et al.*<sup>29</sup> Cryostat sections, 6 µm thick, were cut at -40°C and thaw mounted onto slides treated with 3-aminopropyltriethoxysilane, prepared as described by Tourtellotte *et al.*<sup>30</sup> Sections were fixed in

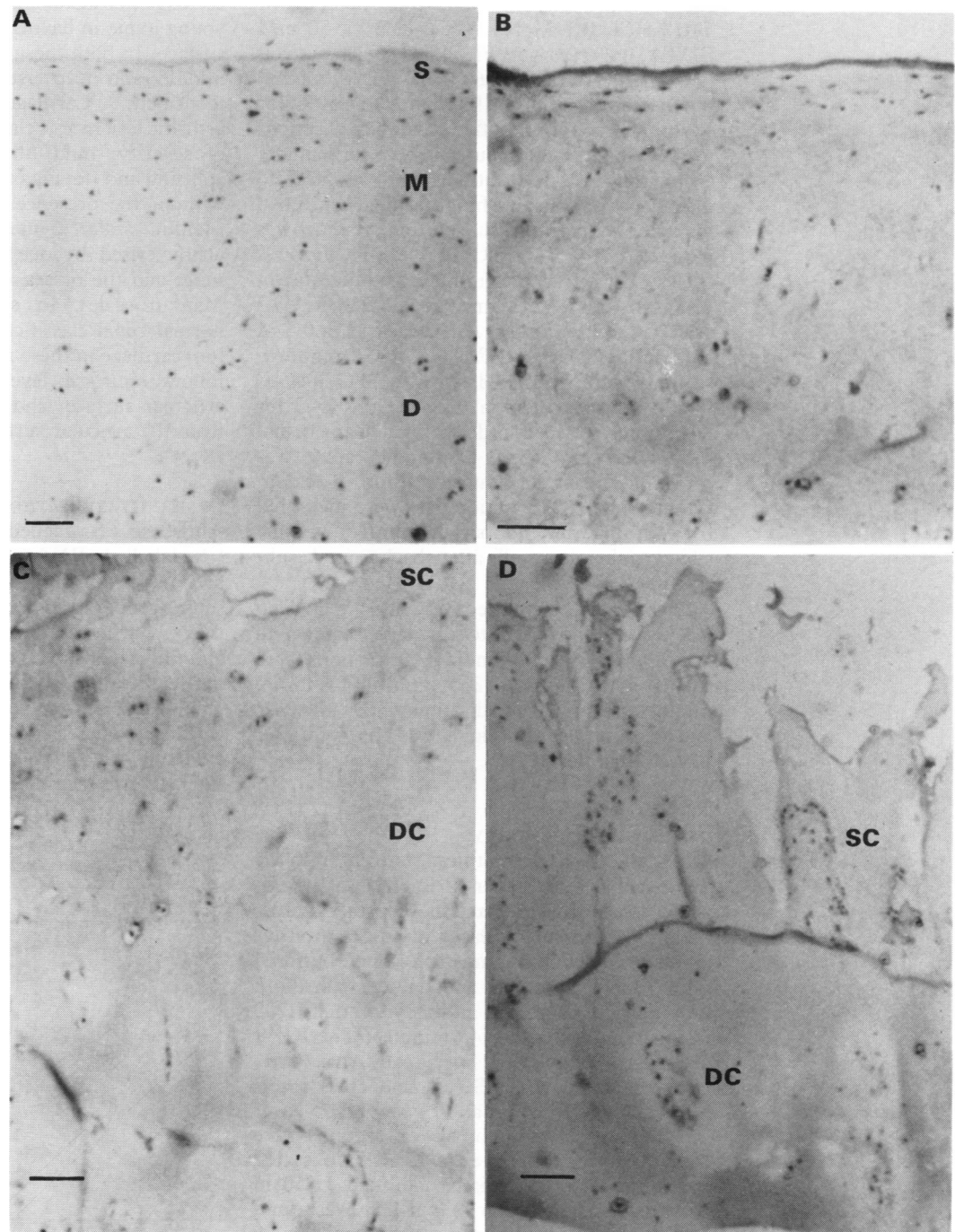


Figure 1 Representative low magnification photomicrographs of human articular cartilage. (A) Cartilage from young subjects showing a smooth intact articular surface. S=superficial layer; M=middle layer; D=deep layer. (B) Cartilage from macroscopically intact regions of osteoarthritic joints has the same anatomical features as A. (C) Mild fibrillation from an osteoarthritic joint with surface disruption. (D) More advanced fibrillation characterised by deeper clefts and cell clones. SC=surface cells (<150 µm deep); DC=deeper cells (>150 µm deep). Haematoxylin and eosin. Bar=100 µm.

4% paraformaldehyde, permeabilised in 0.1% Triton X-100, and dehydrated in a graded alcohol series. Slides were then air dried and stored at 4°C until used. Tissue sections were treated with 0.2 M HCl for 20 minutes to remove basic proteins, and acetylated for 10 minutes in freshly prepared 0.1 M triethanolamine (pH 8.0) containing 0.25% (vol/vol) acetic anhydride. Sections were dehydrated in an ethanol series and air dried. <sup>35</sup>S labelled RNA was heated to 80°C for 30 seconds then cooled on ice, and hybridisation was performed under siliconised glass coverslips in 40% formamide, 10% (wt/vol) dextran sulphate, 200 µg total yeast RNA/ml, 8 mM dithiothreitol, <sup>35</sup>S labelled RNA (3.5 × 10<sup>6</sup> cpm/section for antisense and sense probes), and 'salts' (3 M NaCl, 0.1 M TRIS-HCl, 0.1 M Na<sub>2</sub>PO<sub>4</sub> (pH 6.8), 50 mM EDTA, 0.2% (wt/vol) Ficoll 400, 0.2% (wt/vol) polyvinylpyrrolidone 360 000, 0.2% (wt/vol) bovine serum albumin fraction V) at 50°C for 18 hours in a humidified chamber. Sections were washed for 60 minutes in 50% formamide, 'salts', and 10 mM dithiothreitol at 50°C to dislodge coverslips; washing was then continued for two hours in fresh solution. Tissue was incubated in 0.5 M NaCl in TRIS-HCl/EDTA (pH 7.6) at 37°C for 15 minutes and digested in 20 µg/ml RNase A (Sigma, Poole, Dorset, UK) at 37°C for 30 minutes. After washing in 0.5 M NaCl in TRIS-HCl/EDTA (37°C for 30 minutes), sections were incubated twice in SSC at 45°C for 30 minutes then in 0.2 × SSC at 45°C for 30 minutes. Tissue was dehydrated in an ethanol series and air dried. Autoradiography was for 21 days using K5 emulsion (Ilford, London, UK). In situ hybridisation data were measured by counting the number of autoradiographic silver grains per cell after background subtraction.

Two controls were used: (a) sense <sup>35</sup>S labelled RNA was hybridised to cartilage sections to indicate non-specific binding; (b) sections were treated with RNase A (200 µg/ml in TE for 30 minutes) before hybridisation with antisense and sense probes. No signal was observed after RNase treatment.

#### IMMUNOHISTOCHEMISTRY

Cryostat sections (6 µm thick) were cut from the same cartilage as that used for in situ hybridisation. Sections were fixed in 4% paraformaldehyde for 30 minutes, placed in blocking buffer (1% bovine serum albumin in phosphate buffered saline) for 15 minutes, and treated with 25 µl rabbit antihuman IGF-I antibody (diluted 1:200 in blocking buffer) for 45 minutes. The IGF-I antibody binds IGF-I specifically, has 0.5% cross reactivity with IGF-II, and cross reacts minimally with insulin at 10<sup>-6</sup> mol/l.<sup>31 32</sup> After washing and blocking as above, sections were incubated for 30 minutes in 25 µl biotinylated goat antirabbit IgG (Sigma), diluted 1:250 in blocking buffer. After a further wash and blocking step, sections were treated for 30 minutes with 2.5 µg/ml streptavidin-peroxidase (Sigma) in 0.1 M TRIS-HCl (pH 7.6) containing 1% bovine serum albumin. The sections were washed with 0.1 M TRIS-HCl, pH 7.6, for

two minutes and then treated with substrate (0.5 mg/ml diaminobenzidine in TRIS-HCl and 0.01% H<sub>2</sub>O<sub>2</sub>) for 15 minutes. After a final wash with TRIS-HCl followed by distilled water, sections were mounted in Apathy's medium and viewed by Nomarski interference microscopy.

## Results

### TISSUE SAMPLING

Figure 1 shows low magnification photomicrographs of representative cartilage samples, indicating the type of tissue used for this study. Articular cartilage from macroscopically normal regions of osteoarthritic joints served as controls to cartilage from osteoarthritic lesions. This cartilage was histologically similar to tissue from young joints in having a smooth intact articular surface. In both these cartilages the superficial, middle, and deep layers (or zones) were easily identified.<sup>33</sup> Cartilage sampled from osteoarthritic lesions was classified into two categories of severity: mild fibrillation with superficial splitting and fraying of the articular surface, yet without the presence of cell clones (4 on the Mankin scale<sup>34</sup>); more advanced fibrillation characterised by deeper clefts, cartilage thinning, and the presence of cell clones (8 on the Mankin scale). In osteoarthritic lesions the normal zonal classification is disrupted; therefore cartilage in these regions was categorised as follows: surface layer, <150 µm from the articular surface, and deeper layer, >150 µm from the articular surface.

### IN SITU HYBRIDISATION

Antisense <sup>35</sup>S labelled RNA probes and sense <sup>35</sup>S labelled RNA probes were applied to sections of human articular cartilage. The antisense probe hybridises specifically to IGF-I mRNA, whereas the sense probe is a control and indicates non-specific binding. With the antisense probe, IGF-I mRNA was detected in chondrocytes from osteoarthritic and young cartilage, whereas only a minimal signal was evident with the sense probe (fig 2). Signal intensity was measured by counting the number

Table 1 Background measurements over cartilage matrix

Cartilage layer*	Antisense	Sense
<b>A</b>		
Superficial	2 (0.2)†	2 (0.3)
Middle	2 (0.2)	1 (0.2)
Deep	1 (0.1)	1 (0.2)
<b>B</b>		
Superficial	2 (0.1)	3 (0.2)
Middle	1 (0.2)	2 (0.3)
Deep	3 (0.2)	2 (0.3)
<b>C</b>		
Surface	2 (0.3)	1 (0.3)
Deep	2 (0.2)	1 (0.2)
<b>D</b>		
Surface	4 (0.4)	3 (0.3)
Deep	4 (0.3)	6 (0.4)

\*A=young normal cartilage; B=anatomically normal cartilage from osteoarthritic joints; C=mildly fibrillated cartilage from osteoarthritic joints; D=more advanced lesions from osteoarthritic joints.

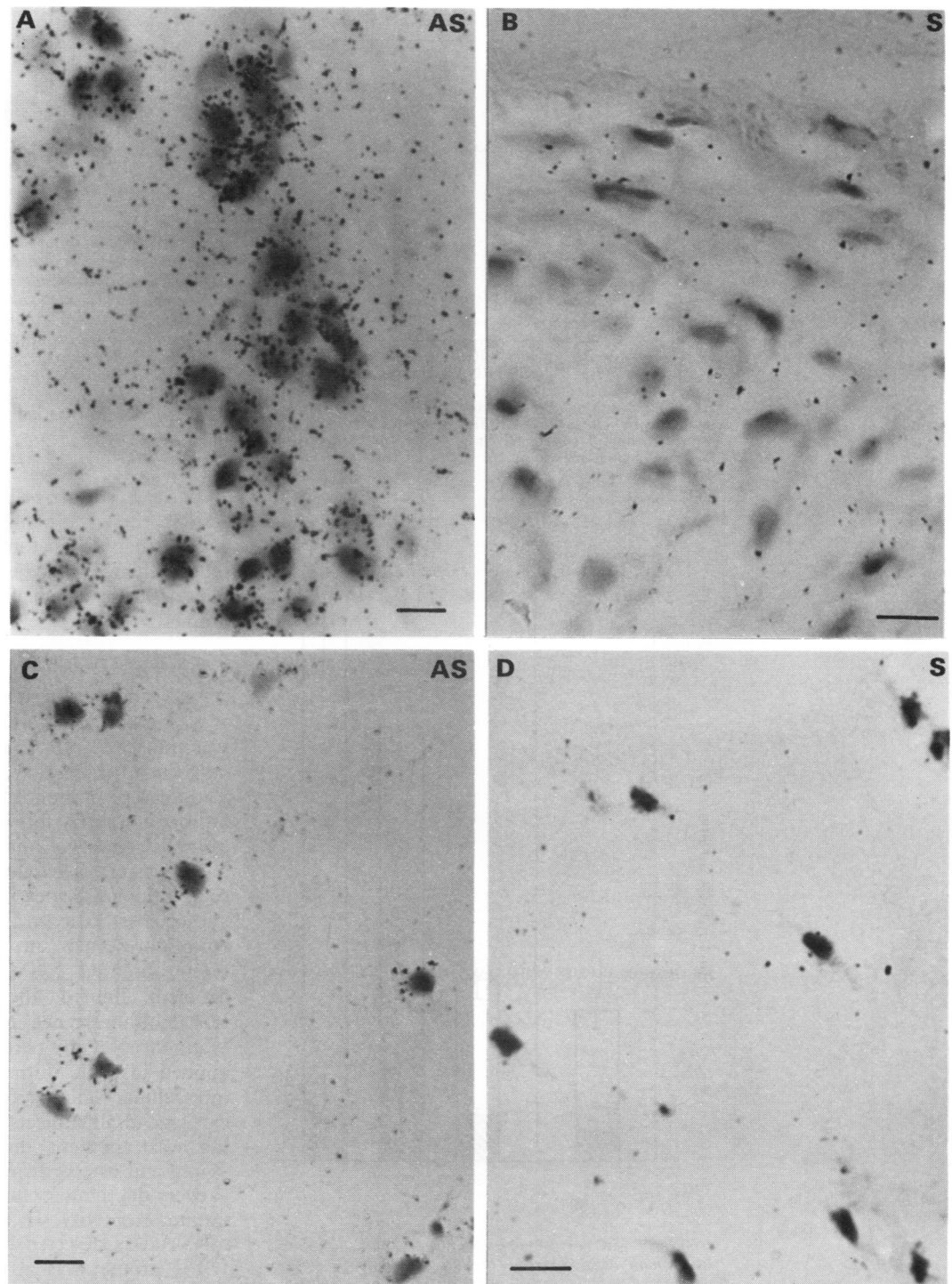
†All values represent means (SE) and indicate number of grains per 80 µm<sup>2</sup>, the average area of a human chondrocyte on a histological section. Total area sampled per value=3500 µm<sup>2</sup>.

of autoradiographic silver grains overlying a cell after background subtraction. Table 1 gives background measurements for the different cartilage samples. Chondrocytes were scored positive for IGF-I message when the signal intensity using the antisense probe was greater than that using the sense probe. A high proportion of osteoarthritic and young chondrocytes, between 70 and 100%, were positive for IGF-I mRNA, as shown in table 2.

The amount of IGF-I message in chondrocytes was estimated (fig 3). In young cartilage the

IGF-I mRNA signal intensity was relatively low and did not differ from anatomically normal cartilage from osteoarthritic joints. In fibrillated osteoarthritic cartilage the signal intensity was significantly higher than in anatomically normal cartilage. There were only slightly more silver grains/cell in more advanced lesions than in mild lesions, and this difference was not significant.

Several differences in signal intensity were apparent when data were expressed as variation with cartilage depth (fig 4). In young cartilage



**Figure 2** Photomicrographs of *in situ* hybridisation of insulin-like growth factor I (IGF-I) mRNA in human articular cartilage using  $^{35}\text{S}$  labelled antisense (AS) and sense (S) RNA probes. With the antisense probe IGF-I message is present in most chondrocytes, whereas with the control sense probe there is minimal signal over cells. (A and B) Cell clones from more advanced lesions of osteoarthritic cartilage (see fig 1D for low magnification histology). (C and D) Chondrocytes from anatomically normal young cartilage (see fig 1A for low magnification histology). Haematoxylin and eosin. Bar = 10  $\mu\text{m}$ .

there were less grains/cell in the superficial layer than in the middle and deep layers. Anatomically normal cartilage of osteoarthritic joints differed from young cartilage with significantly more signal in the middle layer than in superficial and deep layers. In both mild and more advanced fibrillation the number of grains/cell was significantly higher in surface cells than in deeper cells. There was a particularly high signal

Table 2 Percentages of chondrocytes containing insulin-like growth factor I (IGF-I) message

Cartilage layer <sup>a</sup>	Cells examined (n)	Chondrocytes containing IGF-I message (%)
A		
Superficial	60	73
Middle	60	85
Deep	60	88
B		
Superficial	15	80
Middle	15	100
Deep	15	93
C		
Surface	15	100
Deep	15	90
D		
Surface	66	97
Deep	66	72

Cells were scored positive for IGF-I mRNA when the signal using the antisense <sup>35</sup>S labelled RNA probe was greater than that using the control sense probe.

A=young cartilage; B=anatomically normal cartilage from osteoarthritic joints; C=mildly fibrillated cartilage from osteoarthritic joints; D=more advanced lesion from osteoarthritic joints containing cell clones. In C and D, surface=<150 µm; deep=>150 µm.

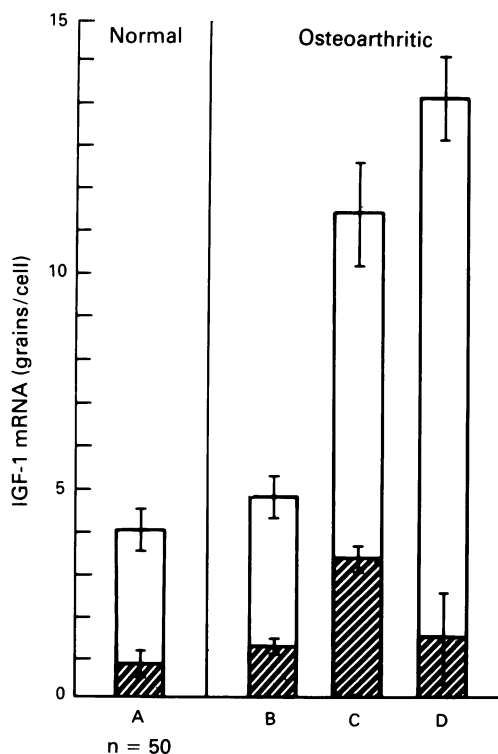


Figure 3 Measurement of in situ hybridisation of insulin-like growth factor I (IGF-I) mRNA in human articular cartilage. Unshaded values are results using <sup>35</sup>S labelled antisense RNA probes and shaded values using <sup>35</sup>S labelled control sense RNA probes. (A) Anatomically normal young cartilage; (B) anatomically normal cartilage from osteoarthritic joints; (C) mild fibrillation from osteoarthritic joints; (D) more advanced fibrillation with cell clones from osteoarthritic joints (see fig 1 for histology).

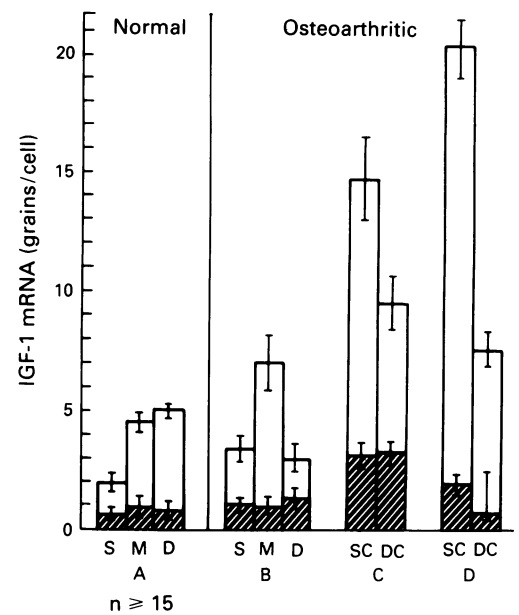
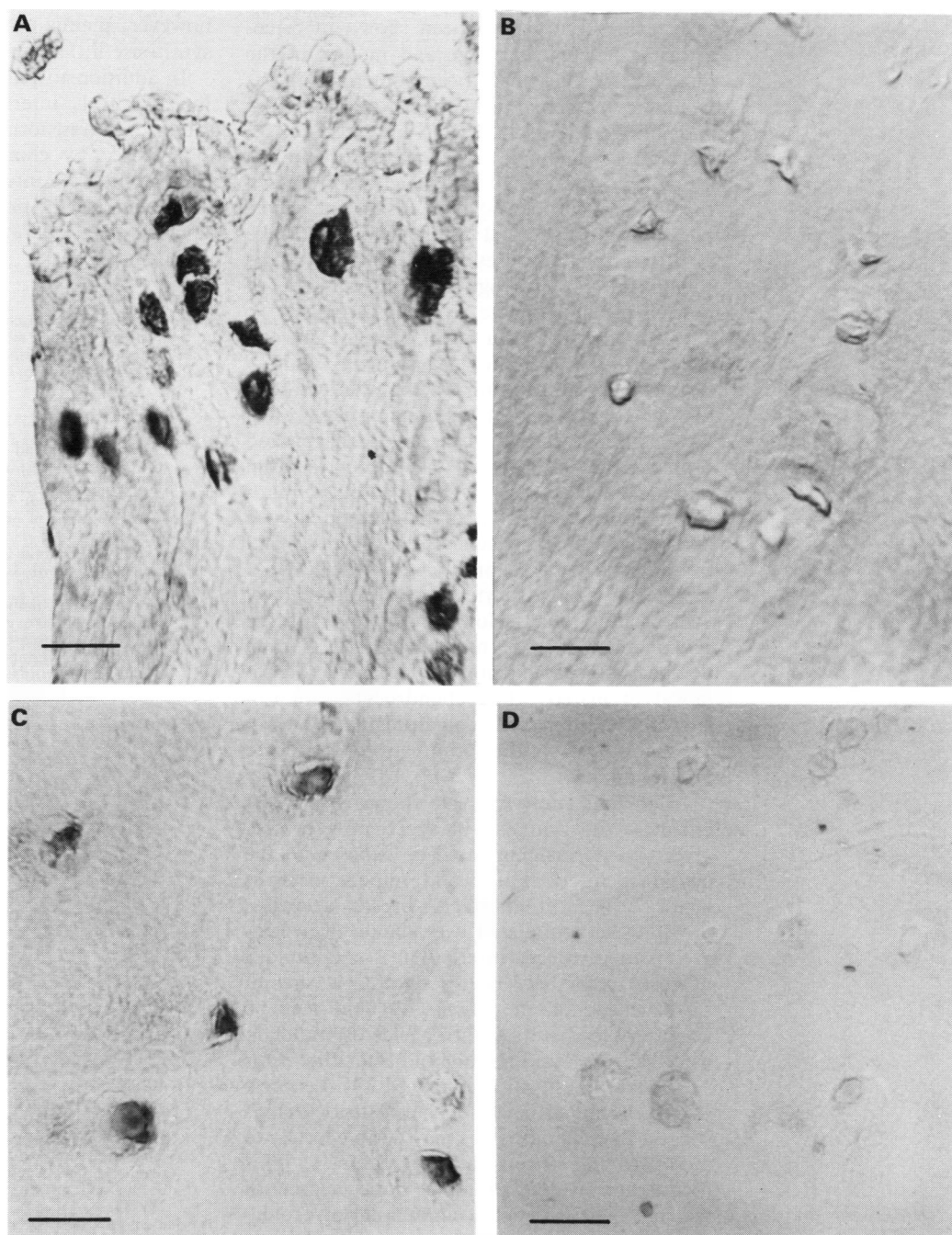


Figure 4 Measurement of in situ hybridisation of insulin-like growth factor I (IGF-I) mRNA in human articular cartilage. Variation with cartilage depth. Unshaded values are results using <sup>35</sup>S labelled antisense RNA probes and shaded values using <sup>35</sup>S labelled control sense RNA probes. (A) Anatomically normal cartilage; (B) anatomically normal cartilage from osteoarthritic joints; (C) mild fibrillation from osteoarthritic joints; (D) more advanced fibrillation with cell clones from osteoarthritic joints. S=surface layer; M=middle layer; D=deep layer; SC=surface cells; DC=deep cells (see fig 1 for histology).

intensity in surface cell clones of more advanced lesions, the levels being about four to five times higher than in anatomically normal cartilage. These changes were evident in the lesions of osteoarthritic cartilage from all the subjects studied.

#### IMMUNOHISTOCHEMISTRY

It was important to establish that IGF-I mRNA was translated into protein. Therefore sections adjacent to those processed for in situ hybridisation were immunolabelled with a polyclonal antibody specific for IGF-I.<sup>31 32</sup> Initially, immunofluorescence microscopy was used and a cytoplasmic distribution of IGF-I protein was seen in chondrocytes from all depths of young and osteoarthritic cartilage (results not shown). Autofluorescence in osteoarthritic cartilage often obscured IGF-I fluorescence and this problem limited the effectiveness of the immunofluorescence technique. Therefore, a biotin-streptavidin-peroxidase technique was applied instead. With this method an intense intracellular staining reaction was seen in the cartilage of all subjects examined (fig 5). Staining was apparent throughout the depth of young and osteoarthritic articular cartilage and at every depth most chondrocytes showed IGF-I immunoreactivity. In all regions where IGF-I mRNA was detected by in situ hybridisation, IGF-I protein was also detected by immunohistochemistry. Although it was not possible to measure the amount of IGF-I protein in chondrocytes, qualitative examination indicated particularly intense staining reactions in cell clones of more advanced osteoarthritic lesions.



**Figure 5** Immunohistochemical localisation of insulin-like growth factor I (IGF-I) protein in human articular cartilage, showing IGF-I distributed intracellularly in a cluster of chondrocytes close to the articular surface of an osteoarthritic lesion (A) and in mid-zone chondrocytes of an anatomically normal region of osteoarthritic cartilage (C). (B) and (D) are controls, in the absence of primary antibody, of cells from an osteoarthritic lesion and from anatomically normal osteoarthritic cartilage respectively. Nomarski interference microscopy. Bar=20  $\mu$ m.

Staining was absent in control experiments where the primary antibody was omitted.

#### Discussion

Our study shows relatively low levels of IGF-I message in anatomically normal osteoarthritic cartilage. In fibrillated osteoarthritic cartilage, however, the message levels were increased and were proportional to the severity of the lesion. This upregulation in osteoarthritic lesions probably represents an attempt at increased matrix replacement and reflects the local mechanical and biochemical demands made by the microenvironment on the chondrocytes.

Thus the particularly high levels of IGF-I mRNA in surface cell clones may indicate a situation where matrix loss is most acute and the demand for matrix repair highest. Our results are in accord with studies on kidney and skeletal muscle, where locally produced IGF-I may also be particularly important in hypertrophy and repair.<sup>8</sup>

Increased synthesis of IGF-I at osteoarthritic lesions may lead to one or more of the following effects: stimulation of proteoglycan production, decreased degradation of proteoglycan, stimulation of collagen production, and increased chondrocyte replication. Increased proteoglycan and collagen production and enhanced

mitogenic activity have been shown in osteoarthritic articular cartilage and models of the disease.<sup>34-36</sup> There have been numerous studies in animal systems showing the matrix promoting and mitogenic effects of IGF-I, but there is a lack of information on osteoarthritic human articular cartilage. It has long been known that IGFs could stimulate [<sup>35</sup>S]sulphate incorporation into rat costal cartilage.<sup>37</sup> Insulin-like growth factor I increases synthesis and decreases catabolism of proteoglycans in cultures of normal pig and bovine articular cartilage.<sup>38-39</sup> The growth factor is a major mediator in fetal calf serum<sup>12</sup> and human synovial fluid,<sup>40</sup> responsible for increasing proteoglycan synthesis in bovine cartilage explants. Insulin-like growth factor I increases the synthesis of collagen and proteoglycan in cultured chondrocytes from rabbit articular cartilage.<sup>11-41</sup> In addition, mitogenic actions of IGF-I are demonstrated in vitro by normal human articular chondrocytes,<sup>42</sup> and by rabbit articular chondrocytes, where IGF-I secreted by chondrocytes may stimulate their own proliferation.<sup>43</sup> However, these mitogenic actions of IGF-I are not apparent in normal bovine cartilage explants.<sup>38</sup> Possibly, the removal of matrix when chondrocytes are isolated for culture results in upregulated IGF-I production, which then stimulates chondrocyte proliferation.

In another study we have shown that IGF-I enhances the synthesis of aggregating proteoglycan in normal pig cartilage exposed to the cytokines interleukin 1 and tumour necrosis factor.<sup>39</sup> In addition, IGF-I directly decreases the cytokine stimulated degradation of proteoglycan in cartilage. Interleukin 1 and tumour necrosis factor have been identified at sites of cartilage erosion in patients<sup>44-45</sup> and may be important mediators of cartilage deterioration in osteoarthritis<sup>46</sup> and rheumatoid arthritis.<sup>47</sup> Our present results showing enhanced IGF-I expression in the chondrocytes of osteoarthritic lesions suggest that chondrocytes may be able locally to modulate the actions of interleukin 1. This modulation would promote some degree of matrix repair and net accumulation of proteoglycan.

A high proportion of chondrocytes, about 70-100%, contain IGF-I message, depending on cartilage source and tissue depth. These data support the concept of an autocrine/paracrine mode of action of IGF-I, acting locally on the same cell or nearby cells. Insulin-like growth factor I mRNA and immunoreactive IGF-I have also been demonstrated in epiphyseal chondrocytes of the rat growth plate by in situ hybridisation and immunofluorescence microscopy<sup>48-49</sup>; these observations suggest that locally produced IGF-I has a specific role in the clonal expansion of differentiated epiphyseal chondrocytes. Variable results have been reported for IGF-I production by chondrocytes in vitro, depending on the culture system used. Pelvic cartilage from chick embryos produces IGF-I peptides in serum free organ culture where cartilage growth is mediated by IGF-I.<sup>50</sup> Rabbit articular chondrocytes synthesise IGF-I in serum free medium supplemented with fibroblast growth factor.<sup>43</sup> Bovine articular cartilage explants,

however, maintained in serum free conditions synthesise little or no IGF-I.<sup>38</sup>

In addition to IGF-I, evidence suggests that interleukin 1, interleukin 6, tumour necrosis factor, and transforming growth factor  $\beta$  are all synthesised by chondrocytes.<sup>51-54</sup> Thus there may be intricate interactions between chondrocytes and locally produced agonists and antagonists that operate by autocrine/paracrine mechanisms.

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