HIV INFECTION OF HUMAN CARTILAGE

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Infection of human cartilage with HIV in vivo has not previously been reported. Specimens of articular cartilage taken at postmortem from ten patients who were HIV-positive were examined. Two had AIDS and eight were believed to have stage-2 disease.

The standard polymerase chain reaction (PCR) protocol was modified to allow semiquantitative analysis of the samples. Oligonucleotide primers labelled with $^{32}$P gamma-ATP were used to detect a segment of HIV DNA and a control DNA gene segment (HLA genome) to estimate the ratio of infected cells. The $^{32}$P-labelled PCR products were separated on acrylamide gels and visualised directly by autoradiography and computer densitometry.

Infection of human cartilage in vivo was demonstrated in nine of the ten samples in which the PCR analysis was positive. The other did not react sufficiently to produce detectable radiolabelled PCR product despite repeated DNA digestion and extraction. Cartilage infected with HIV could be a potential source of HIV when used in operations.


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Infection of human cartilage with HIV in vivo has not been described. Our aim was to examine cartilage from patients infected with HIV for evidence of infection with this virus.

MATERIALS AND METHODS

Specimens were received from ten postmortems. All the patients were HIV-antibody-positive; two had AIDS and eight were believed to have stage-2 disease. One patella was obtained from each patient. Donor blood was washed from the sample during collection. The articular cartilage of all specimens was macroscopically normal.

The standard polymerase chain reaction (PCR) protocol was modified to allow semiquantitative analysis of samples. Simultaneous amplification and detection of a single-copy human gene and an HIV sequence have been developed to allow a semiquantitative assay of HIV infection when the cell number or DNA input is unknown (Lee et al 1991). This method uses an internal control to validate the efficiency of the PCR reaction and to determine the HIV copy number relative to that of the DNA of the input cell.

A further modification of the PCR uses $^{32}$P 5' end-labelled synthetic oligonucleotide primers so that the transfer and hybridisation steps used in other methods are
omitted and the number of PCR cycles is therefore decreased. This is then followed by direct autoradiography of gel-resolved products (Arrigo et al 1989).

Details of methods. One gram of articular cartilage was crushed with a mortar and pestle and incubated for 72 to 78 hours at 37°C in a lysis solution consisting of 100 mM NaCl, 10 mM TRIS.Cl (pH 8), 25 mM EDTA (pH 8), 0.5% sodium dodecylsulphate (SDS) and 0.1 mg/ml proteinase-K. The solution was extracted three times with phenyl-chloroform-isooamylalcohol (25:24:1) and the DNA was precipitated by ethanol sodium acetate. The pellet was rinsed with 70% ethanol, dried and resuspended in 50 µl TE (10 mM Tris pH 7 and 1 mM EDTA). The DNA content was measured spectrophotometrically.

A control curve for the quantitative PCR method was constructed using dilutions of HIV from infected human lymphocytes. The chronically HIV-infected cell line H3B (Li and Burrell 1992) was spiked with non-infected HUT-78 cells (NIH AIDS Research and Reference Reagent Programme, ERC Bioservices Corp, Rockville, Maryland). Chromosomal DNA extracted from mixed HUT-78 and H3B cells, with tenfold dilutions of 10^{-2} to 10^{-5} H3B cells, was analysed in duplicate with each PCR analysis of clinical samples, which were assessed in triplicate. A fresh PCR reaction master mix and control cell DNA were used for each repeat analysis. A new control curve was constructed for each new reaction mixture or radiolabelled probe pair. A negative control derived from HUT-78 cell chromosomal DNA was included in each PCR reaction.

Oligonucleotide probes GH26 (Saiki et al 1985) and SK38 (Kellogg and Kwok 1990) were independently end-labelled on the 5' terminus with ^32P gamma-ATP (1 to 5 × 10^6 cpm). Labelled nucleotides were used immediately or stored overnight at −20°C in lead pigs.

The PCR reaction mixture of 100 µl contained 5 µl 10 × Thermus Aquaticus (Tag) buffer (Bresatec, Adelaide, South Australia), 2.25 mmol MgCl₂, 1 mmol of each of the four deoxynucleoside triphosphates (Perkin Elmer Cetus, Roche Molecular Systems, New Jersey), 0.04 units of Taq polymerase (Bresatec), and 5 µl of each primer dilution. The hot-start technique was used with the reaction mixture with less enzyme and DNA, overlaid with ampliwax PCR gem 100's (Perkin Elmer) and heated to 60°C for 5 to 10 minutes. The reaction tubes were cooled to room temperature and then an upper reaction mixture consisting of 0.4 µl Taq polymerase (5.5 units/µl; Bresatec), water and 5 µg DNA was added using a positive displacement pipette (Bresatec) to aliquot the DNA.

Each amplification cycle consisted of 1.5 minutes at 95°C, one minute at 56°C and two minutes at 72°C followed by ten minutes at 72°C. Twenty-five cycles were used in all experiments.

Ten microlitres of each sample and a ^32P gamma-ATP radiolabelled molecular weight marker (pUC 19 DNA restricted with Hpa II; Bresatec) were added to 2 µl of loading buffer. The samples and marker were subjected to 8% polyacrylamide gel electrophoresis at 100 V for 60 to 120 minutes. After running, the gel was dried at 80°C under vacuum for one hour and exposed to autoradiographic film (XAR-5, Kodak, Rochester, New York) without enhancing screen for 90 to 180 minutes at room temperature.

A clean phosphor screen (Molecular Dynamics, Sunnyvale, California) was exposed to the radiolabelled gels for 18 hours at room temperature. The phosphor screen was scanned with a PhosphorImager (ImageQuant, model 400B; Molecular Dynamics) using Molecular Dynamics ImageQuant version 3.0 software. Net optical densities for the specific HIV and HLA bands were determined by volume integration with manual background subtraction for each sample. The results of HIV and HLA band intensities (in OD/mm²) for each sample were analysed.

HIV and HLA band intensities for the control cell dilutions were plotted for each H3B/HUT-78 cell dilution. CA-cricket graph computer software (1990 Computer Associates International Inc, San Diego, California) was used to construct a simple regression curve which determined the ratio of HIV-infected cells from the radiolabelled PCR product of clinical samples.

RESULTS

Autoradiography showed delineation of the 115 base pair HIV-1-gag region and 242 base pair segment of the HLA-DQ-α genome with no interfering bands (Fig. 1). HUT-78 cell DNA included as a negative control was amplified in all experiments and the 115 base pair HIV-1-gag region

![Fig. 1](image-url)

PCR coamplification of cartilage from ten HIV infected patients. Detection of the 115 base pair HIV segment indicates HIV infection with incorporation of HIV into the chromosomal DNA of the host cells. A 242 base pair segment of the HLA-DQ-α genome is included as an internal control of PCR efficiency and to determine the HIV copy number relative to cell copy number.
was not detected.

The two primer pairs detected from cartilage specimens and H3B cell dilutions had HIV and HLA specific bands that were successfully amplified. Detection of the HIV-1 \( \text{gag} \) region from nine samples indicates in vivo infection of human cartilage with HIV.

The PCR product was not detected from case 6 after three PCRs and repeated DNA digestion and extraction. Samples were examined in triplicate but the PCR product was obtained from cases 1, 2 and 4 in only one experiment after repeated DNA extraction.

<table>
<thead>
<tr>
<th>Case</th>
<th>HIV infected cells ( \log_{10} )</th>
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<tr>
<td>1</td>
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<td>3</td>
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<tr>
<td>10</td>
<td>-2</td>
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The results of semiquantitative PCR analysis are summarised in Table I. The calculated ratio of infected cells to uninfected cells was \( 10^{-2} \) to \(<10^{-5} \).

**DISCUSSION**

In this study HIV was detected in all samples of human cartilage in which PCR analysis was achieved. One of the ten samples did not react sufficiently to produce a detectable radiolabelled PCR product despite repeated DNA digestion and extraction. This may have been related to the quality of the extracted DNA or to the presence of inhibitors.

Human cartilage is avascular and the specimens examined were intact and macroscopically normal. They were thoroughly washed and were not contaminated with blood or other body fluids. It is therefore assumed that the HIV DNA detected was present in infected chondrocytes. In nine of the ten samples in which the PCR product was obtained HIV infection was identified in 1.0% to 0.001% of cells. In our study a finding of up to 1% HIV infection of chondrocytes in vivo is consistent with in vitro findings of HIV infection in cells of mesenchymal origin (Ikeuchi et al 1990; Mellert et al 1990).

The mechanism of HIV infection of chondrocytes in vivo remains unknown. The mean ‘pore’ size of human cartilage is approximately the size of the serum albumin molecule (Maroudas 1979). Thus, cell-to-cell HIV transmission is prevented but cell-free infection is possible. HIV infection independent of the CD4 receptor by phagocytosis and endocytosis has been reported (Clapham et al 1989; Harouse et al 1989; Tateno, Gonzalez-Scarano and Levy 1989) and it may be that this mechanism is effective in vivo in human cartilage. The differing results of investigation of in vitro infection of human chondrocytes (Ikeuchi et al 1990; Bujia et al 1993a) are not explained by our study which suggests that chondrocytes can be infected under appropriate conditions.

Our results are consistent with the finding of the presence of HIV in synovial fluid in patients infected with this virus (Withrington et al 1987). HIV arthropathy has been described (Forster et al 1988) and chondrocyte infection may be related to this. We did not study the function and microscopic appearance of the chondrocytes and therefore the clinical role of chondrocytes infected with HIV remains unknown. Cartilage may be a potential site of systemic HIV infection. Although HIV infection from cartilage allografts has not been reported, we do not agree with Bujia et al (1993a) who suggest that they may be less likely to transmit HIV than other allografts. Cartilage allografts from donors of unknown status should be vigilantly screened for the possible presence of HIV.

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No benefits in any form have been received or will be received from a commercial party related directly or indirectly to the subject of this article.

**REFERENCES**


