

Complement C1s, a classical enzyme with novel functions at the endochondral ossification center: immunohistochemical staining of activated C1s with a neoantigen-specific antibody

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Abstract. The secondary ossification center of 14- to 16-day-old hamster tibiae was examined immunohistochemically with active and inactive C1s-specific antibodies, RK5 and RK4, respectively. At the ossification center, chondrocytes differentiate from proliferating and hypertrophic to degenerating stages, and their site is occupied by the bone marrow. C1s was strongly immunostained in hypertrophic chondrocytes. In order to discover whether C1s is activated at a particular site, the cartilage was immunostained with RK5 and RK4. RK5 mainly reacted with degrading matrix around invading vessels. In contrast, RK4 strongly stained hypertrophic chondrocytes. Immunoelectron microscopy revealed C1s on degrading fragments of chondrocytes and fibers of cartilage matrix. Decorin, one of the major matrix proteoglycans, was dose and time dependently degraded by C1s. Type II collagen and type I gelatin were also degraded. Articular cartilage from patients with rheumatoid arthritis was positively immunostained (11/12 cases) with an anti-C1s monoclonal antibody (mAb) PG11, whereas normal articular cartilage (5/5 cases) was negative, suggesting C1s participation in the etiology of rheumatoid arthritis.

Key words: Complement C1s – Neoantigen – Cartilage – Hypertrophic chondrocyte – Immunohistochemistry – Rheumatoid arthritis – Decorin – Syrian hamster – Human

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Abbreviations: *mAb*, Monoclonal antibody; *PLP*, periodate-lysine-paraformaldehyde; *SDS-PAGE*, sodium dodecyl sulfate-polyacrylamide gel electrophoresis; *HE*, hematoxylin and eosin; *DFP*, diisopropyl fluorophosphate; *MMP-9*, matrix metalloproteinase 9

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Introduction

In the classical complement cascade, C1s activated by C1r cleaves C2 and C4 which have been known as sole substrates of C1s. Recently, however, we have demonstrated that C1s degrades type II collagen (Yamaguchi et al. 1990), a major constituent of cartilage matrix and gelatin, and activates zymogen of matrix metalloproteinase 9 (MMP-9) (Sakiyama et al. 1994). MMP-9 is known to be synthesized by chondrocytes in vivo (Toyoguchi et al. 1996) and in vitro (Nakagawa et al., 1997), and colocalized with C1s at the primary ossification center of human femur (Sakiyama et al. 1994). Therefore, C1s is thought to have a role in cartilage degradation.

Cartilage consists of chondrocytes and extracellular matrix, which contains several types of collagen and proteoglycans. During development and growth of long bones, chondrocytes pass through resting, proliferating, maturing, hypertrophic and degenerating stages. This naturally occurring cell death aids in tissue remodeling. Degenerated chondrocytes disappear, leaving enlarged lacunae and surrounding matrix which is eventually degraded by monocytes (Hunziker et al. 1984) and chondroclasts (Lewinson and Silberman 1992), and the site is replaced by bone marrow (Cancedda et al. 1995; Caplan and Pechak 1987). Programmed cell death of chondrocytes is accompanied by an expression of an *s-myc* protein (Asai et al. 1994) and tissue transglutaminase (Aeshimann et al. 1993; Fesus et al. 1991). In most tissues, dead cells are usually cleaned away by phagocytic cells. In cartilage, an avascular tissue, however, degenerating cells disappear before phagocytic cell invasion. Apoptotic signaling is known to induce various kinds of proteolytic enzymes (Nicholson et al. 1995; Seamus and Green 1995). Degrading enzymes may play a major role in the cleaning mechanisms of cartilage. We have shown that C1s synthesis increases in accordance with chondrocyte differentiation both in vivo (Toyoguchi et al. 1996)

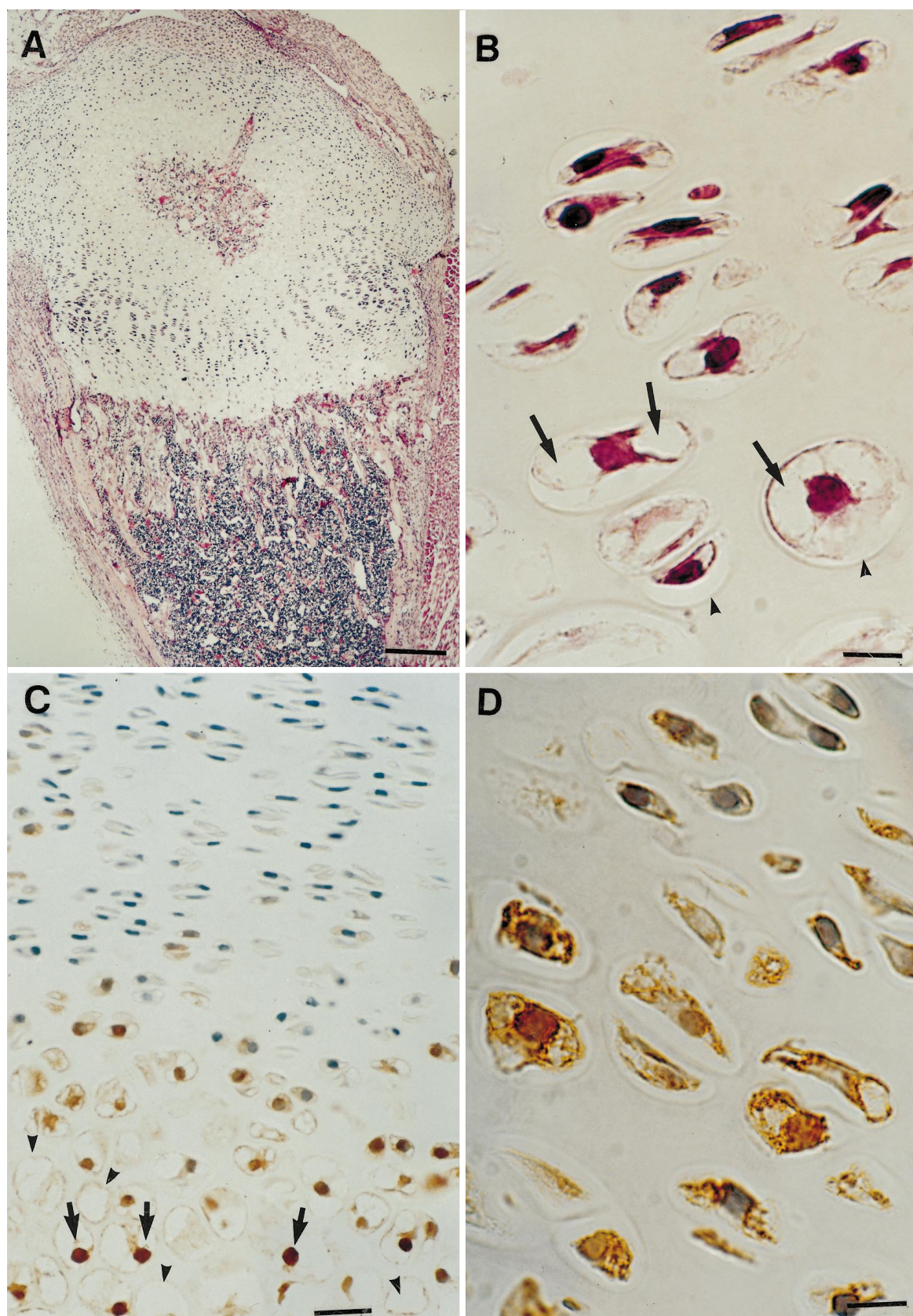


Fig. 1A-D. Morphology and immunostaining of CIs at the secondary ossification site of hamster tibia. Proximal tibia of 14-day-old hamster fixed with PLP and frozen-sectioned. **A** Staining with HE. *Bar:* 270 μm ; $\times 37$. **B** Higher magnification of **A**. *Bar:* 10 μm ; $\times 925$. **C** Section stained with anti-hamster CIs mAb PG11 (20 $\mu\text{g/ml}$), post-stained with hematoxylin. Proliferating chondrocytes

negative in the staining with PG11, but degenerating hypertrophic chondrocytes strongly positive (nuclei and cytoplasm). *Bar:* 54 μm ; $\times 185$. **D** Higher magnification of hypertrophic chondrocytes. Degrading edges of cytoplasmic holes were stained predominantly. *Bar:* 10 μm ; $\times 925$

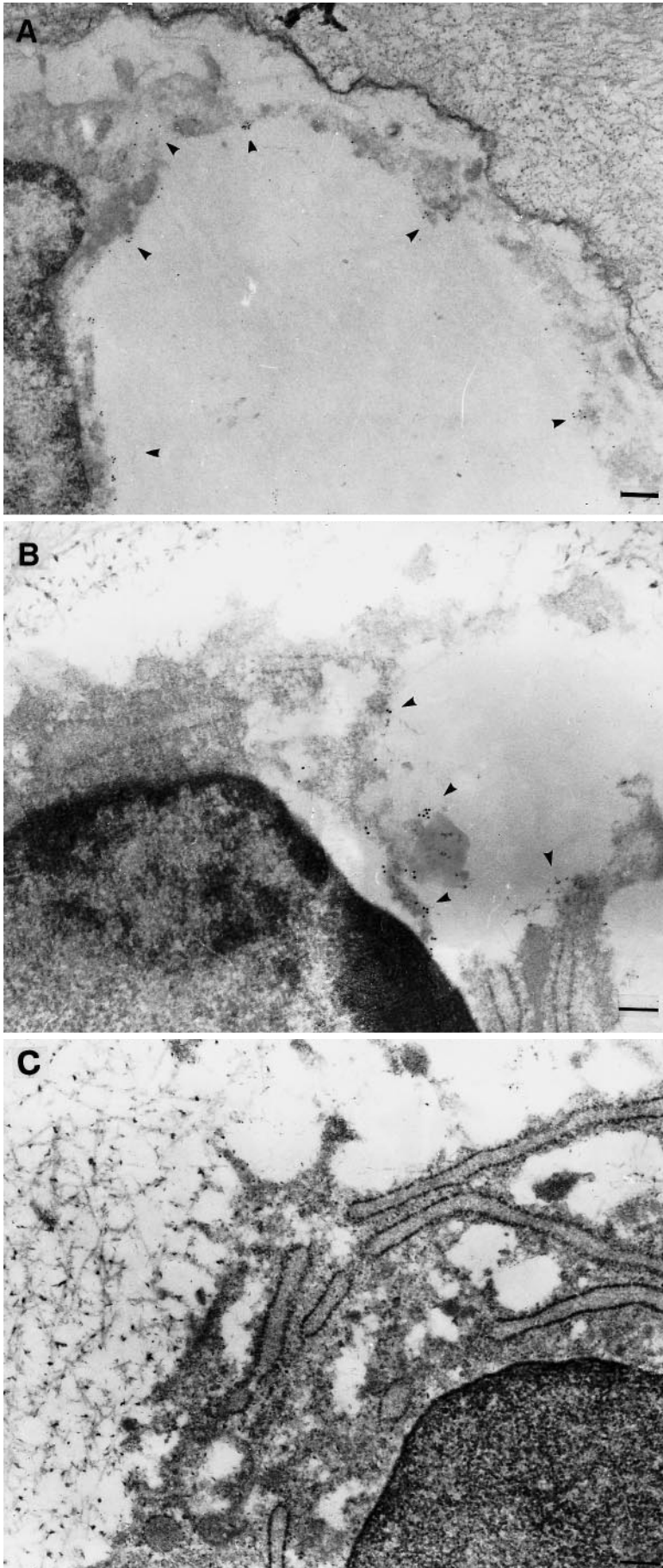


Fig. 2A–C. Immunoelectron microscopy of hypertrophic chondrocytes. Epiphyseal cartilage of 14-day-old hamster tibia fixed with PLP and immunostained with PG11 (**A, B**) or M241 (**C**), followed by biotin-anti-mouse IgG and streptavidin-gold as described in the text. Gold particles (*arrowheads*) in **A** and **B** are seen on degrading cellular fragments. **A** Bar: 267 nm, $\times 30000$; **B, C** bar: 200 nm, $\times 40000$

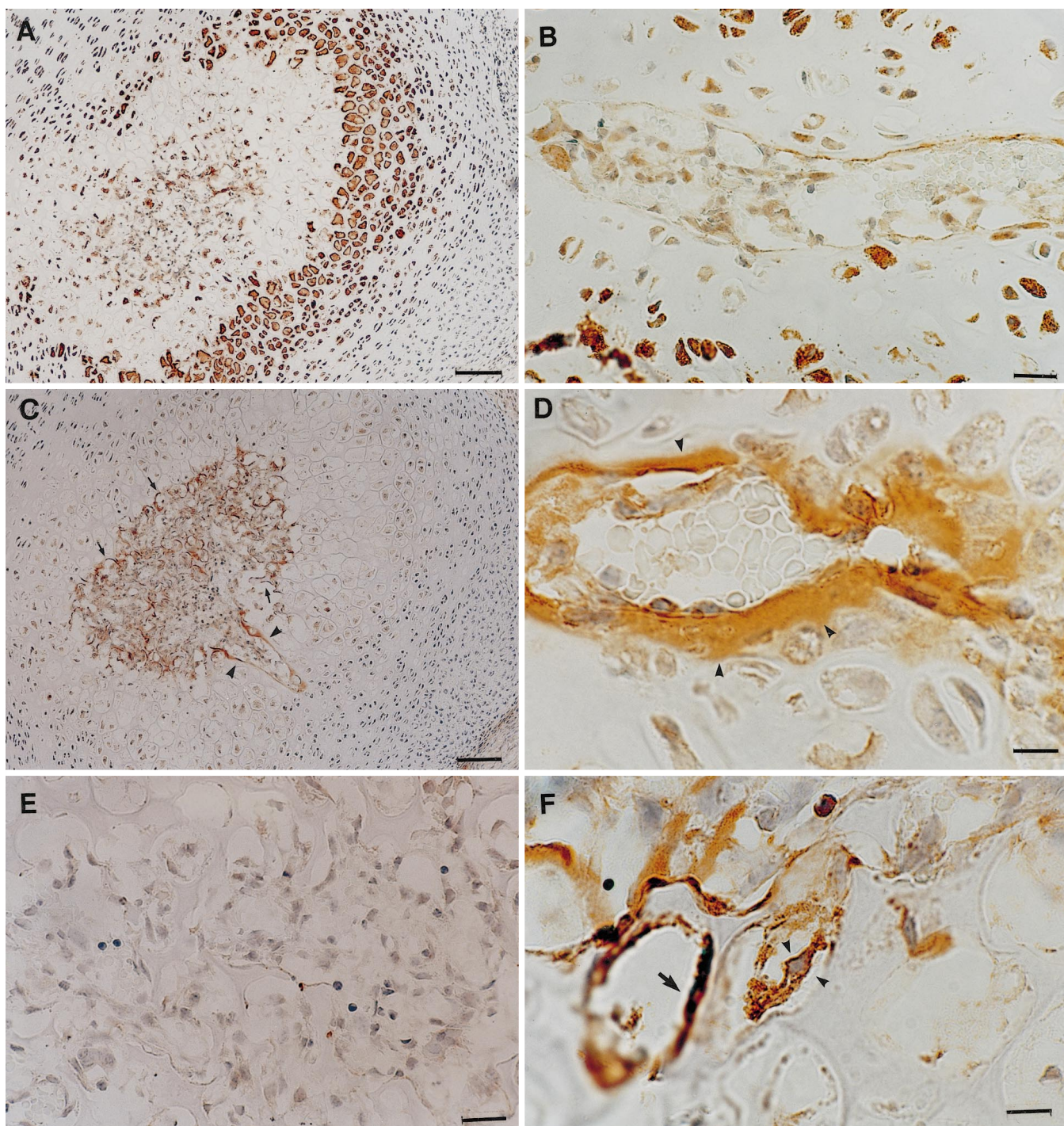


Fig. 3A–F. Differential immunostaining of active and inactive Cls. Epiphyseal cartilage fixed and frozen-sectioned as described in the text. The samples were incubated with either RK4 (20 $\mu\text{g}/\text{ml}$; **A, B**), RK5 (20 $\mu\text{g}/\text{ml}$; **C, D, F**) or normal rabbit IgG (20 $\mu\text{g}/\text{ml}$; **E**) and post-stained with biotin-anti-rabbit IgG F(ab)₂ and HRP-

streptavidin. Inactive and active Cls were found mainly in hypertrophic chondrocytes (**A, B**) and extracellular matrix around invading vessels (**C, D, F**). **A, C** Bar: 108 μm , $\times 65$; **B** bar: 27 μm , $\times 260$; **D, F** bar: 10 μm , $\times 700$; **E** bar: 54 μm , $\times 130$

Hamster tibial secondary ossification center was immunostained with anti-hamster-Cl mAb PG11 (Fig. 1C, D). In the hypertrophic zone, edges of lacunae and cytoplasmic holes of hydrated chondrocytes and some, but not all, nuclei were strongly immunostained (Fig. 1C, D), whereas proliferating chondrocytes were nega-

tive (Fig. 1C). The strongly stained cells in Fig. 1C (arrows) may correspond to the dark condensed cells reported by Farnum and Wilsman (1987). M241, a mAb specific to human activated Cls, did not react even with degenerating hypertrophic chondrocytes (data not shown). Immunoelectron micrographs revealed that Cls

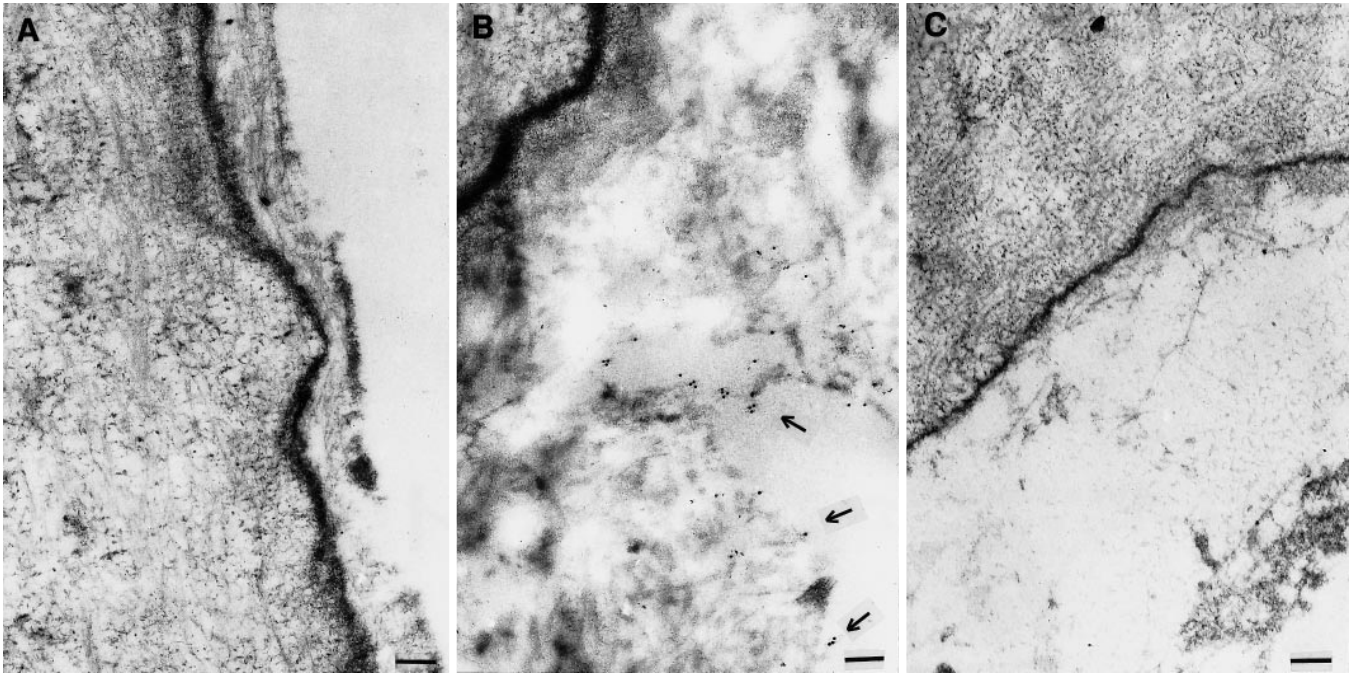


Fig. 4A–C. Immunoelectron micrograph of matrix around invading marrow. The sections were incubated with RK4 (**A**), RK5 (**B**) or normal rabbit IgG (**C**) and then biotin-anti-rabbit IgG F(ab)₂,

followed by streptavidin-gold. Gold particles (*arrows*) were observed on amorphous mat-like structures at the degrading side of the matrix. *Bar*: 200 nm; $\times 40000$

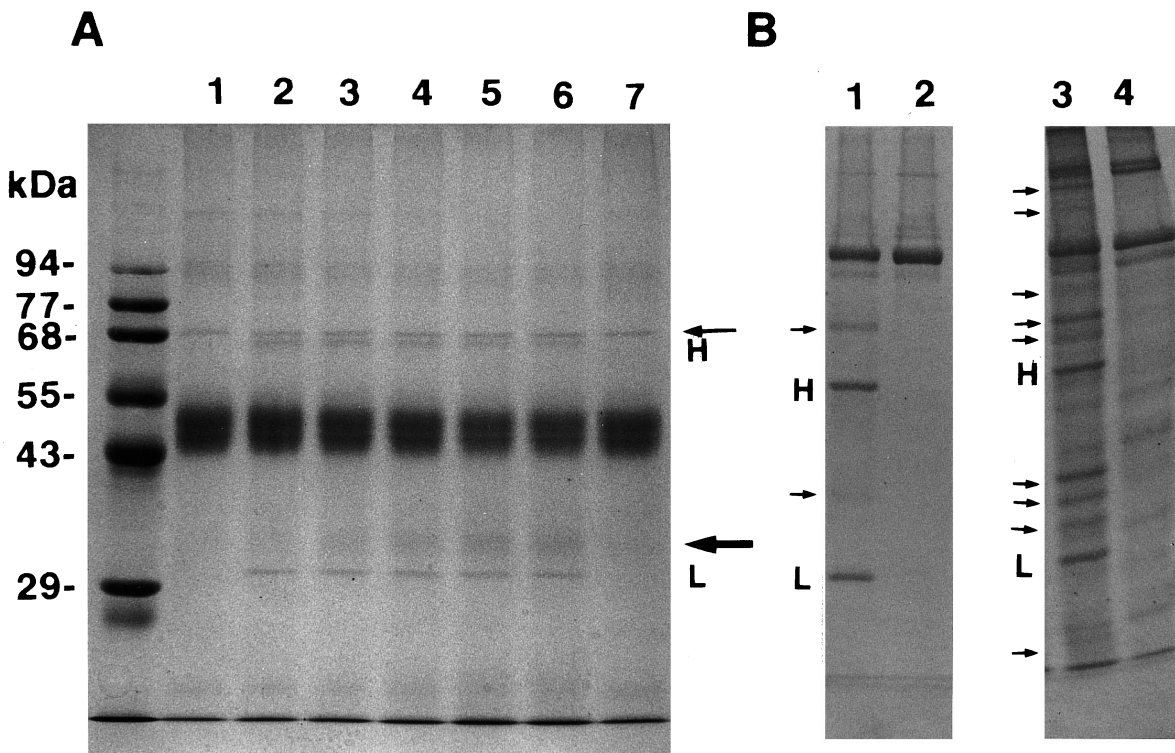


Fig. 5A, B. Digestion of matrix components by Cls. **A** Time-dependent degradation of decorin by Cls. Decorin (8 μg) was incubated with (*lanes 2–6*) or without (*lanes 1, 7*) Cls at a Cls-to-decorin ratio of 1:25 in PBS for 0 h (*lane 1*), 2 h (*lane 2*), 4 h (*lane 3*), 6 h (*lane 4*), 12 h (*lane 5*) and 24 h (*lanes 6, 7*) as described in the text. After the incubation, samples were analyzed on SDS-PAGE under reducing conditions and stained with Coomassie blue. *Large arrow*, Degradation product of decorin; *small*

arrow, chondroitinase ABC; *H, L, H* and *L* chains of Cls, respectively. **B** Degradation of type II collagen (*lanes 1, 2*) and type I gelatin (*lanes 3, 4*) by Cls. Five micrograms of each protein was incubated with (1 μg ; *lanes 1, 3*) or without (*lanes 2, 4*) Cls in TRIS-HCl buffer (pH 7.4) overnight at 37° C. The incubation mixtures were analyzed on SDS-PAGE as described in the text. *Small arrows*, Digestion products of the substrates; *H, L, H* and *L* chains of Cls

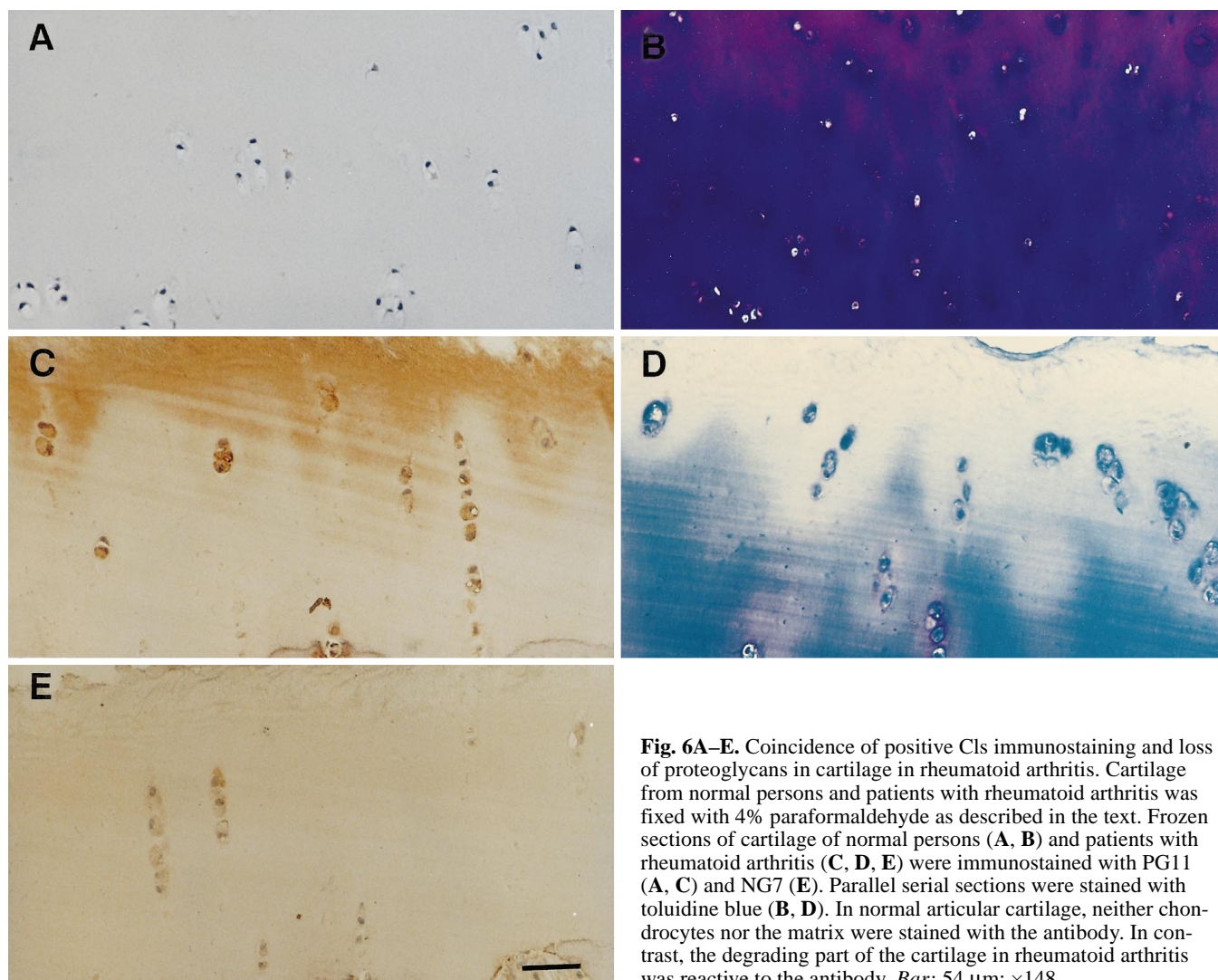


Fig. 6A–E. Coincidence of positive Cls immunostaining and loss of proteoglycans in cartilage in rheumatoid arthritis. Cartilage from normal persons and patients with rheumatoid arthritis was fixed with 4% paraformaldehyde as described in the text. Frozen sections of cartilage of normal persons (**A, B**) and patients with rheumatoid arthritis (**C, D, E**) were immunostained with PG11 (**A, C**) and NG7 (**E**). Parallel serial sections were stained with toluidine blue (**B, D**). In normal articular cartilage, neither chondrocytes nor the matrix were stained with the antibody. In contrast, the degrading part of the cartilage in rheumatoid arthritis was reactive to the antibody. *Bar:* 54 μm ; $\times 148$

was localized at degrading edges of cytoplasmic vacuoles and cellular fragments (Fig. 2A, B). No gold particles were visible when M241 was used (Fig. 2C).

Differential immunostaining of active and inactive Cls in situ

Since localization of Cls in the late hypertrophic zone suggested its involvement in cartilage degradation, we further examined whether Cls is activated at the degrading sites. When the specimen was treated with RK4, which reacts only with inactive Cls, hypertrophic chondrocytes (Fig. 3A, B) were positively stained. In contrast, RK5, the antibody specific to the neopeptide of activated Cls, stained Cls in the extracellular matrix around the invading blood vessels (Fig. 3C, D, arrowheads), the edges of the degrading matrix (Fig. 3C, F, arrows) and degenerative chondrocytes (Fig. 3F, arrowheads). These staining patterns displayed a striking contrast in the location of active and inactive Cls. Control rabbit IgG showed negligible staining (Fig. 3E).

Immunoelectron-microscopic study around the marrow revealed that activated Cls was localized on the fine filamentous mat-like structure at the degrading side of the demarcation line. No definite labeling was seen in the samples stained either with RK4 or non-immune Rig (Fig. 4A, C).

Digestion of matrix components by Cls

Since Cls was found to be activated at the ossification center, we examined whether it degraded matrix proteins. Decorin is known to stabilize collagen fibrils by binding to the molecules (Scott 1988; Hedbom and Heinegard 1989; Svensson et al. 1995). The susceptibility of decorin to Cls was also examined in the present study. When decorin was incubated with active Cls at enzyme-to-substrate ratios of 1:50, 1:40, 1:30, 1:25, 1:20, 1:15, 1:10 and 1:5, it was degraded in concentration-dependent manner (data not shown). Time-dependent degradation of decorin is shown in Fig. 5A. A major decorin degradation fragment of 33 kDa molecular mass was

generated after 2 h incubation with Cls (1:25) and increased with incubation time up to 24 h (Fig. 5A, large arrow). At the same time the bands of decorin became thin.

Cls split type II collagen into 2:1 fragments (Fig. 5B, lane 1, arrows), which were different from digestion fragments of collagenase (Harper 1980). Gelatin was degraded into several pieces by Cls (Fig. 5B, lane 3, arrows). The activities of Cls were inhibited by DFP and M241 (data not shown). These results confirmed our previous observations (Yamaguchi et al. 1990).

Cls immunostaining in human articular cartilage

Articular cartilage in rheumatoid arthritis (11/12 cases) was positively immunostained in both matrix and chondrocytes with PG11 (Fig. 6C), whereas normal cartilage (5/5 cases) was negative (Fig. 6A). The staining pattern of Cls in cartilage with rheumatoid arthritis coincided with the loss of proteoglycans, which was detected by toluidine-blue staining (Fig. 6D). Cartilage in rheumatoid arthritis showed poor reactivity to NG7 (Fig. 6E), a mAb against hamster Cls (Sakiyama et al. 1991), which is poorly cross-reactive to human Cls.

Discussion

In the present work we have shown by immunohistochemistry that Cls was activated at the secondary ossification center of hamster tibia. At the central part of the ossification center, degenerating late hypertrophic chondrocytes disappeared before bone marrow cell invasion. Since no phagocytic cells were seen in the lacunae at this stage, chondrocytes might be digested by their own degrading enzymes. It is possible that Cls is one of the degrading enzymes, because (1) Cls synthesis increases with hypertrophic changes of chondrocytes and (2) active Cls-specific antibody RK5 immunostains the edges of degrading fragments of chondrocytes. Immune complex-independent activation of Cls is induced by exposure to various cellular components such as cell membranes (Storrs et al. 1983), intermediate filaments (Linder et al. 1979), DNA (Ziccardi 1982), cardiolipin (a mitochondrial inner membrane component) (Kovacsovic et al. 1987) and polyanionic substances (Rent et al. 1975). These cellular components may be available for Cls binding at the ossification center where chondrocyte degradation proceeds. Proteoglycans (belonging to the polyanionic molecules) are also abundant in the cartilage matrix. Cls around blood vessels may be permeated from vessels and/or secreted from endothelial cells, since endothelial cells are reported to synthesize Cls (Gulati et al. 1993) and to be immunostained by PG11 (Sakiyama et al. 1994). Cls reaching cartilage matrix can be activated rapidly, because Cls at this site was reactive to RK5, but not to RK4.

Our previous studies have shown that Cls has the ability to degrade collagens and activate proMMP-9 (Sakiyama et al. 1994). The present study further dem-

onstrates that decorin is also a substrate of this enzyme, indicating that Cls has a broad range of substrates.

In hamster articular cartilage, Cls is not detected by either immunostaining or in situ hybridization (Toyoguchi et al. 1996). Consistent with these observations, normal human articular cartilage displayed no immunostaining with anti-Cls mAb PG11. However, since strong Cls immunostaining is seen in cartilage with rheumatoid arthritis, it is suggested that Cls may participate in pathological cartilage matrix degradation as well as in the physiological turnover observed in the ossification center.

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