

Fibrogenesis and Fibrolysis in Collagenous Colitis

Patterns of Procollagen Types I and IV, Matrix-Metalloproteinase-1 and -13, and TIMP-1 Gene Expression

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Collagenous colitis is characterized by the deposition of a superficial subepithelial collagenous layer, the pathogenesis of which is unknown. Because the excess matrix deposition is potentially reversible, a labile imbalance between fibrogenesis and fibrolysis may be suspected. Expression of procollagen $\alpha 1(I)$ and $\alpha 1(IV)$, matrix-metalloproteinase (MMP)-1 and -13, and tissue inhibitor of metalloproteinase (TIMP)-1 genes was semiquantitated by *in situ* hybridization on serial biopsies of 12 patients with collagenous colitis and compared to controls. Collagen types I, III, IV, and VI, tenascin, undulin/collagen XIV, and α -actin were localized by immunohistology. The superficial collagen layer stained strongly for collagen types I, III, and VI, and particularly for tenascin, but not for undulin. Elevated procollagen $\alpha 1(I)$, procollagen $\alpha 1(IV)$, and TIMP-1 transcript levels were found in α -actin-positive cells with linear distribution underneath the superficial collagenous layer, whereas MMP-1 RNA expression was variable and restricted to cell clusters. MMP-13 expression was undetectable. The patterns of procollagen $\alpha 1(I)$ - and $\alpha 1(IV)$ -specific labeling, combined with an intense tenascin- but absent undulin-specific staining, indicate deposition of an immature interstitial matrix that may be susceptible to degradation. The restricted MMP-1 RNA expression, counteracted by increased TIMP-1 expression, suggests locally impaired fibrolysis as a relevant factor in the pathogenesis of collagenous colitis. (*Am J Pathol* 1999, 155:493–503)

Collagenous colitis (CC), initially described by Lindström in 1976, is clinically characterized by watery diarrhea in the absence of significant endoscopic and radiological findings.¹ Histologically, CC typically displays a band-like linear deposition of extracellular matrix (ECM) underneath the surface epithelium, with a minimal thickness of 10 μ m required for diagnosis.^{2–5} This ECM layer does not extend along crypts and is mainly composed of collagens type I and III as well as fibronectin.^{6,7} The surface epithelium is often flattened and atrophic and may be infiltrated by lymphocytes, similar to the condition described as lymphocytic colitis, which, however, lacks the subepithelial collagen band.^{3,8,9} The etiology of CC is still unknown and has been related to autoimmune phenomena, to the use of nonsteroidal anti-inflammatory drugs,¹⁰ and to other exogenous, possibly bacterial, agents.¹¹ Today many authors favor a combination of inflammatory and autoimmune mechanisms as cause of the disease.⁹ The pathogenic events leading to the formation of the subepithelial collagenous band are similarly enigmatic.

Recent studies on patients subjected to surgical fecal stream diversion¹² as well as anecdotal evidence¹³ point to a potentially rapid reversibility of the excess collagenous deposits. Thus, these deposits seem to reflect a local disturbance of ECM turnover resulting in the formation of a provisional ECM. Moreover, this suggests that not only the formation, but also the degradation of excess ECM may be relevant in the pathogenesis of CC.

Among ECM degrading enzymes, matrix-metalloproteinases (MMPs) have a central role. MMPs are a family of zinc-dependent neutral proteinases with overlapping but distinct substrate spectra. Four subfamilies of MMPs are known: collagenases, gelatinases, stromelysins, and membrane-MMPs.^{14–16} Together with MMP-8 (collagenase-2, neutrophil collagenase), MMP-1 (collagenase-1, interstitial collagenase) and MMP-13 (collagenase-3) form the MMP-subfamily of collagenases, which are instrumental for the degradation of native interstitial colla-

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gens, in particular collagen types I, II, and III.¹⁴⁻¹⁶ Because of their destructive potential, MMPs are regulated at three levels, namely transcription, zymogen activation, and inhibition of the active enzyme. *In vitro*, stromelysin-1 (MMP-3) may enhance the activity of MMP-1 by a factor of 5 to 12 by additional N-terminal proteolysis, whereas MMP-13 may be activated by gelatinase A (MMP-2) and the membrane type MT1-MMP (MMP-14).¹⁶⁻¹⁸ The details of MMP-1 and -13 activation *in vivo*, however, are largely unknown. Inhibition of active MMP-1 is a major function of tissue inhibitor of metalloproteinases (TIMP)-1.¹⁴⁻¹⁶ In body fluids, MMP-1 may also be inhibited by α_2 -macroglobulin.¹⁹

CC is diagnosed by histology and, unlike ulcerative colitis, does not require continuous histological re-evaluation. Thus, in a common clinical setting, CC biopsies are usually available only as single time-point formalin-fixed specimens. Because of their size, extractive procedures cannot be performed, making analysis of enzyme activities impossible. The study of ECM and MMPs in CC biopsies depends, therefore, on localizing a limited number of relevant gene products.

These considerations and restrictions led us to use immunohistology and semiquantitative *in situ* hybridization to establish the expression patterns of procollagens type I and IV as representative molecules for interstitial and basement membrane ECM, respectively. Furthermore, immunostaining for tenascin and undulin/collagen XIV was performed to assess the maturity of the matrix deposits. Expression of MMP-1, MMP-13, and TIMP-1 as pivotal elements in collagen type I degradation was assessed by *in situ* hybridization to obtain indicators for the local fibrolytic potential.

Materials and Methods

Tissues

Paraffin blocks of mucosal biopsies from 12 patients with a clinical history of chronic watery diarrhea were drawn from the files of the University Institutes of Pathology in Berlin, Hamburg, and Freiburg. Biopsies had been immersed in formalin immediately after removal and submitted for histopathological diagnosis. Criteria for the diagnosis of CC were as recently described.⁷ The thickness of the superficial collagen layer was measured with an ocular micrometer on well-oriented tissue sections. For each biopsy the minimal, maximal, and mean thickness of the superficial ECM layer were recorded along a range of 5 to 10 consecutive intercryptal spaces. The patient data, the biopsy sites, and the histopathological characteristics are listed in Table 1. As controls, we used colonic tissues with regular histology (at a distance of >10 cm from the focal lesions, $n = 7$), with ulcerative colitis ($n = 4$), and with Crohn's disease ($n = 3$), and squamous cell carcinomas of the head and neck region ($n = 9$).

Tissue Sectioning

Sections (5 μ m) were collected onto 3-aminopropyl triethoxysilane-coated slides, dried for several hours at 37°C, dewaxed in xylene, rehydrated in graded alcohols, and immediately processed.

Immunohistology

Collagens type I, III, and VI were detected using affinity-purified rabbit antibodies specific for monkey collagen type I and monkey procollagen type III aminoterminal propeptide²⁰ as well as a rabbit antibody against human collagen type VI.²¹ Tenascin was localized with a monoclonal antibody specific for human tenascin (MAB1927, Chemikon, Temecula, CA). Immunohistology for undulin was performed with a monoclonal antibody against human undulin (clone 15/III/D6).²² Collagen type IV and smooth-muscle α -actin were detected with the antibodies CIV22 and 1A4, respectively (Dako, Glostrup, Denmark). The monoclonal antibodies required pretreatment of paraffin sections with pronase (Boehringer Mannheim, Mannheim, Germany) at a concentration of 0.5 mg/ml in phosphate buffered saline, pH 7.2, for 10 minutes at 37°C. The immunohistochemical detection of MMP-1 was performed with affinity-purified polyclonal rabbit antibodies specific for human MMP-1 (Quartett, Berlin, Germany). The alkaline phosphatase anti-alkaline phosphatase (APAAP) method was used for the detection of immobilized antibodies, using affinity-purified mouse anti-rabbit immunoglobulin serum (Dianova, Hamburg, Germany), affinity-purified rabbit anti-mouse immunoglobulin serum (Dako), and APAAP complex (1:20 dilution, Dako). Alkaline phosphatase was developed with new fuchsin.

Preparation and Labeling of Probes

pGEM1 (Promega, Madison, WI)-based run-off transcription expression vectors were constructed by subcloning of the 1300-bp Pst fragment of $\alpha 1$ R1 corresponding to the triple helical region of rat $\alpha 1$ (I) procollagen,²³ the 500-bp BamHI fragment of pHT21 corresponding to the triple helical region of human $\alpha 1$ (IV) procollagen,²⁴ as well as the 735 bp SstI-EcoRI and the 730 bp EcoRI-SstI fragments of pCilasel (#57685, American Type Culture Collection, Manassas, VA) harboring a human MMP-1 cDNA.²⁵ The TIMP-1 probe was a reverse transcription-polymerase chain reaction-derived cDNA fragment.²⁶ All of these probes had been extensively characterized in previous studies.²⁶⁻²⁹ The MMP-13 probe was generated by reverse transcription and subsequent amplification of positions 24 to 495 in the reported sequence³⁰ using placental RNA as well as the primers 5'-CTG CCT TCC TCT TCT TGA GCT GG and 5'-ATG ATG TCA GCA ATG CCA TCG. All constructs were subjected to nucleic acid sequence analysis and conformed to the published sequences. After linearization of the plasmids with either HindIII or EcoRI restriction endonuclease, T7 or SP6 RNA-polymerase (Gibco-BRL, Eggenstein, Germany), respectively, were employed to obtain run-off transcripts of ei-

Table 1. Summary of Clinical Information, Histopathological Diagnoses, and Morphometric Data

Case	Age/sex	Diagnosis	Location	Collagenous band (μm)	Inflammation	IEL			
1	43 F	CC	1 Ascending	30.3 (7.5–50.0)	1	<5			
			2 Hepatic flexure	38.0 (12.5–42.5)	1	<5			
			3 Transverse	29.3 (7.5–40.0)	1	<5			
			4 Splenic flexure	27.3 (15.0–40.0)	1	<5			
			5 Descending	24.5 (7.5–50.0)	1	<5			
			6 Sigmoid	36.7 (17.5–42.5)	1	<5			
			7 Rectum	26.7 (10.0–50.0)	1	<5			
2	81 F	CC	Splenic flexure	12.7 (5.0–37.5)	0–1	<5			
			3	52 F	CC	Sigmoid	18.8 (7.5–57.5)	1	<5
			4	56 M	CC	Rectum	16.7 (5.0–37.5)	1	<5
5	52 F	CC	1 Appendix	16.9 (10.0–75.0)	1	<5			
			2 Sigmoid	11.1 (5.0–50.0)	1	<5			
			3 Rectum	11.3 (5.0–12.5)	1	<5			
6	51 F	CC	1 Colon	36.3 (5.0–50.0)	1	<5			
			2 Colon	35.3 (12.5–42.5)	1	<5			
			3 Colon	22.9 (10.0–45.0)	1	5–10			
			4 Colon	33.8 (17.5–50.0)	1	5–10			
7	55 F	CC	1 Cecum	38.3 (2.5–82.5)	1	<5			
			2 Ascending	23.8 (2.5–50.0)	1	<5			
			3 Transverse	44.2 (20.0–65.0)	1	<5			
			4 Descending	37.7 (12.5–50.0)	1	<5			
			5 Sigmoid	36.2 (5.0–55.0)	1	<5			
			6 Rectum	17.1 (10.0–52.5)	1	<5			
8	55 F	CC	1 Descending	16.8 (5.0–27.5)	1	5–10			
			2 Sigmoid	14.5 (7.5–37.5)	1	<5			
			3 Rectum	15.4 (5.0–35.0)	1	<5			
9	68 M	CC	Cecum	26.3 (5.0–82.5)	1	<5			
10	38 F	CC	1 Ascending	38.2 (7.5–100.0)	1	<5			
			2 Transverse	25.8 (5.0–42.5)	1	<5			
11	85 F	CC	1 Colon	35.0 (12.5–75.0)	1	<5			
			2 Colon	24.3 (7.5–65.0)	1	<5			
12	75 F	CC	1 Rectosigmoid	30.5 (12.5–137.5)	1	<5			
			2 Transverse	20.8 (5.0–75.0)	1	<5			
13	60 M	Adenoma	Colon*	6.0 (2.5–12.5)	0	<5			
14	68 M	Carcinoma	Colon*	7.8 (2.5–12.5)	0	<5			
15	84 M	Carcinoma	Colon*	8.0 (2.5–12.5)	0	<5			
16	74 M	Carcinoma	Colon*	4.0 (1.5–7.5)	0	<5			
17	64 M	Carcinoma	Colon*	3.8 (1.5–5.0)	0	<5			
18	40 F	Carcinoma	Colon*	6.5 (2.5–10.0)	0	<5			
19	73 M	Diverticulitis	Colon*	7.5 (1.5–12.5)	0	<5			
20	30 F	UC	Colon	2.5 (1.5–5.0)	2	<5			
21	40 F	UC	Colon	3.0 (1.5–5.0)	2–3	<5			
22	70 F	UC (rem.)	Colon	1.5 (0.0–5.0)	1	<5			
23	50 M	UC (rem.)	Colon	2.0 (0.0–2.5)	1	<5			
24	38 F	CD	Colon	5.5 (1.5–7.5)	2	<5			
25	26 F	CD	Colon	2.5 (1.5–4.0)	2	<5			
26	31 F	CD	Colon	2.0 (1.5–2.5)	2	<5			
27	64 M	CD	Colon	2.2 (2.0–2.4)	3	<5			

CC, collagenous colitis; UC, ulcerative colitis; UC (rem.), ulcerative colitis in remission; CD, Crohn's disease; Colon*, mucosa distant to focal lesion; IEL, intraepithelial lymphocytes per 100 epithelial cells.
 Inflammation: density of lymphoplasmacytic infiltrate graded as absent (0), mild (1), moderate (2), or severe (3).

ther the anti-sense (complementary to mRNA), or sense (anti-complementary, negative control) strands. Transcription and labeling of RNA probes were performed as described previously using [³⁵S]-uridine-5'-(α -thio)-triphosphate (1250 Ci/mmol, New England Nuclear, Dreieich, Germany).^{26,27} The specific activity routinely obtained was $1.2\text{--}1.4 \times 10^9$ cpm/ μg .

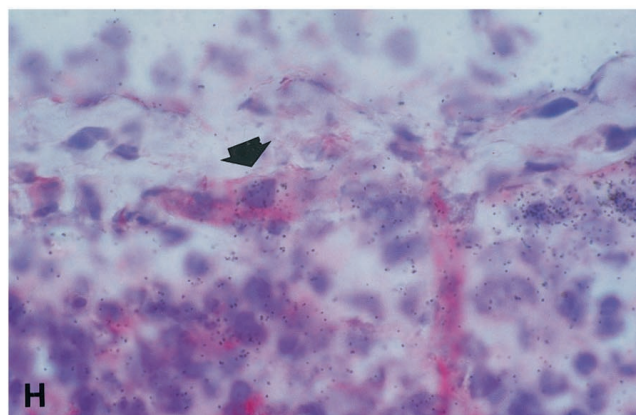
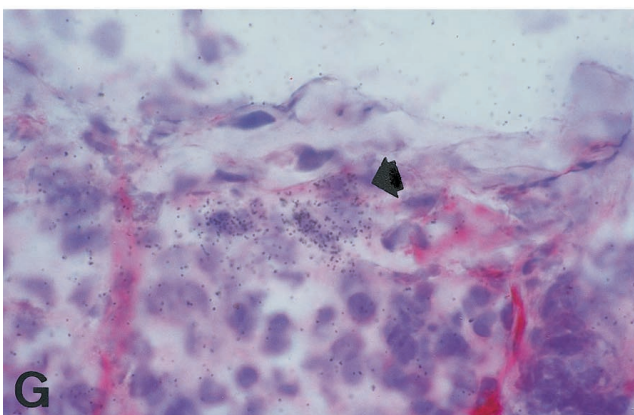
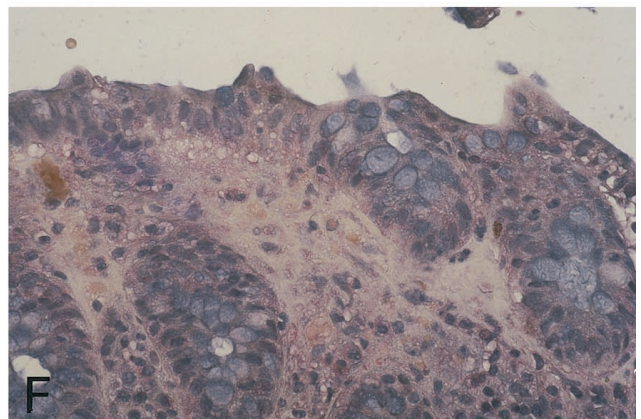
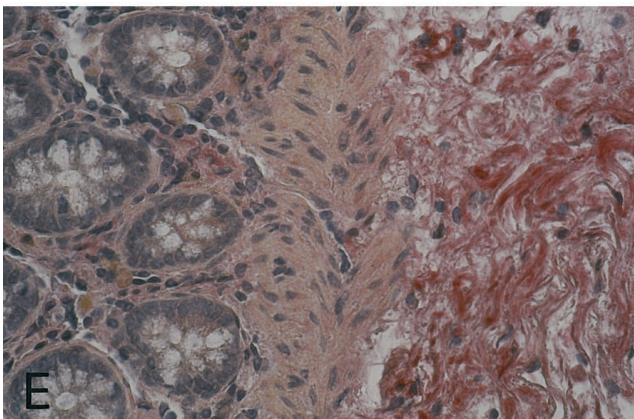
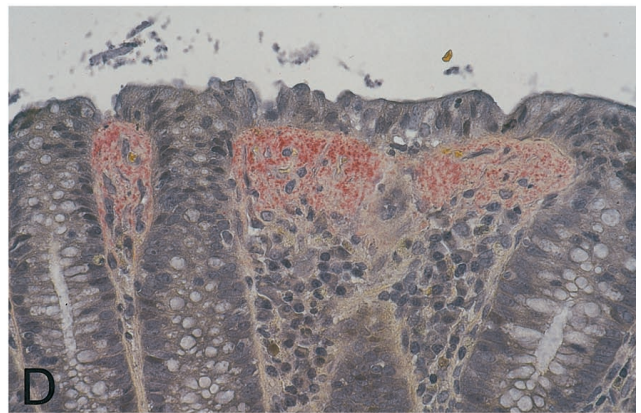
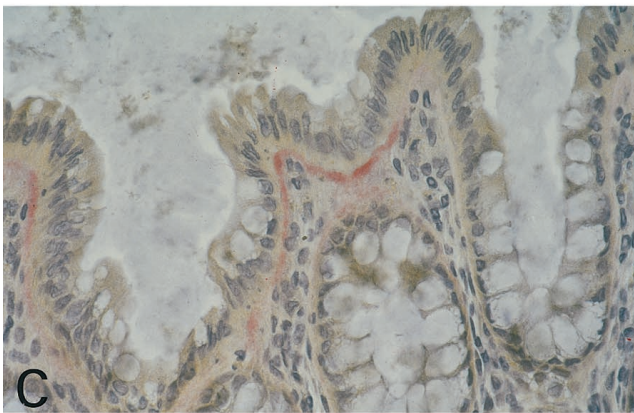
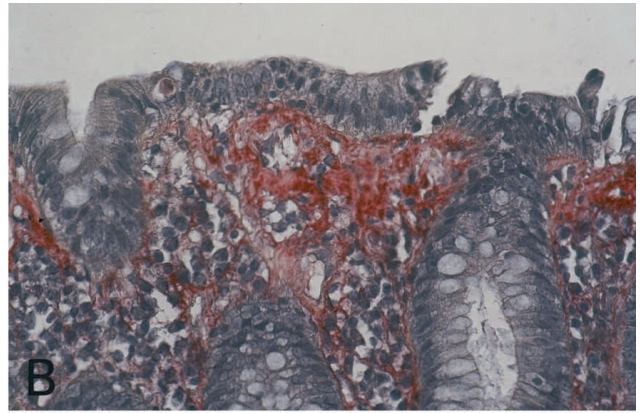
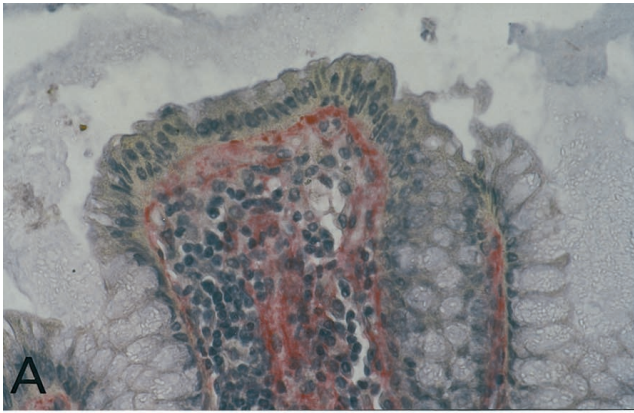
In Situ Hybridization

Pre-hybridization, hybridization, washing procedures including removal of non-specifically bound probe by ribonuclease A digestion, and autoradiography of slides, as

well as the procedure for combined immunohistology and radioactive *in situ* hybridization, were as described in detail.^{26,27}

Statistical Evaluation

Autoradiographic signals were evaluated by counting silver grains attributable to individual cells. The background labeling was determined on adjacent serial sections subjected to sense control hybridization. In each tissue section the number of positively labeled cells and the relative transcript levels, ie, grains per cell, were determined in 10 microscopic fields at a magnification of 1:400. For this



purpose the lamina propria was divided into three compartments: the superficial subepithelial myofibroblast (sSEMF) sheet, the pericryptal subepithelial myofibroblast (pcSEMF) sheet, and the remainder of the lamina propria (LP). The sSEMF sheet was defined as the double cell layer immediately under the superficial collagenous band extending to the crypt opening. The pcSEMF sheet consisted of the double cell layer immediately adjacent to the necks and bases of crypts. The standard error of the mean (SE) was calculated and statistical evaluation was performed using the Mann-Whitney U-test.

Results

Conventional Histology

Colon biopsies diagnosed as CC displayed a characteristic sharply delineated, band-like linear deposition of ECM immediately underneath the surface epithelium without extension along the crypts. The subepithelial layer showed an average thickness between 11.1 and 44.2 μm . In many cases, the surface epithelium appeared flattened. The number of intraepithelial lymphocytes was small. These findings were accompanied by a mild, uncharacteristic inflammatory infiltrate of the lamina propria, composed mainly of small lymphocytes, plasma cells, and few eosinophils (Table 1).

Immunohistology

In all cases immunostaining for procollagen type III (Figure 1, A and B), and collagens type I and VI was found in the extracellular space of the lamina propria, around blood vessels, and under the surface and crypt epithelium. In CC, staining with these antibodies was most intense in the superficial collagenous layer. In ulcerative colitis and Crohn's disease, a moderate and partially intense staining for procollagen type III and collagens type I and VI was seen in the submucosal stroma. Collagen type IV was localized in basement membranes of epithelium and blood vessels but did not accumulate in the superficial collagenous layer. Tenascin-specific staining was largely restricted to a thin layer underneath the surface epithelium in normal mucosa (Figure 1C). In CC biopsies, tenascin-specific staining was most prominent within the subepithelial layer in its entire thickness (Figure 1D). In ulcerative colitis, staining with this antibody was seen in areas with a dense inflammatory infiltrate of the lamina propria, particularly around blood vessels. The undulin-specific antibody displayed a weak staining in the lamina propria, decorating densely packed fiber bundles in the submucosal stroma (Figure 1E). The subepi-

thelial ECM in normal controls and, remarkably, in CC specimens did not display undulin-specific staining (Figure 1F). In contrast, areas of transmural fibrosis in Crohn's disease showed intense staining for undulin but not for tenascin. Some of the stromal cells beneath the superficial collagenous layer stained for smooth-muscle α -actin, indicating a myofibroblastic phenotype of these cells. Immunostaining for MMP-1 displayed a weak diffuse staining within the lamina propria with accentuation around scattered mononuclear cells.

In Situ Hybridization

Procollagen $\alpha 1(\text{I})$ transcripts were detectable within the lamina propria of all specimens. In normal colonic tissue, the signal intensity, ie, the number of procollagen $\alpha 1(\text{I})$ -specific silver grains attributable to individual cells, did not differ significantly between sSEMF, pcSEMF, and other lamina propria cells (Figures 2A and 3). In contrast, CC specimens displayed a significantly enhanced signal intensity ($P < 0.001$) in sSEMF cells in an almost linear, uninterrupted distribution (Figures 2B, 2C, and 3). The relative number of cells expressing procollagen $\alpha 1(\text{I})$, however, was similar in normal mucosa and CC specimens. Pericryptal subepithelial cells and some of the superficial subepithelial cells stained for smooth-muscle α -actin, identifying these cells as myofibroblasts. By combining immunohistology and *in situ* hybridization, we found that at least a proportion of approximately 30% of the procollagen $\alpha 1(\text{I})$ -expressing cells of the sSEMF sheet were α -actin-positive myofibroblastic cells (Figure 1, G and H).

In normal mucosa, the cellular $\alpha 1(\text{IV})$ procollagen transcript levels were near the threshold of detection and required extended autoradiographic exposure of up to 36 days to be clearly labeled. These cells were found underneath the surface and crypt epithelium and along small blood vessels (Figure 2E). In CC, procollagen $\alpha 1(\text{IV})$ RNA expression showed a significantly elevated number of labeled cells in all compartments of the LP and an increased signal intensity in the sSEMF ($P < 0.01$) (Figures 2F and 4). The procollagen $\alpha 1(\text{IV})$ -expressing cells displayed a procollagen $\alpha 1(\text{I})$ -like linear distribution pattern with increased transcript levels in sSEMF cells.

MMP-1 RNA transcript levels were close to the threshold of detection in normal mucosa. In CC, MMP-1 transcript levels were elevated in 8 of 12 cases. The number of MMP-1 expressing cells was increased in both the sSEMF and pcSEMF cells, but cells with highly elevated signal intensity were exclusive to the sSEMF sheet and showed a discontinuous and patchy distribution (Figures 5, A–C, and 6). sSEMF cell clusters with elevated MMP-1

Figure 1. Immunostaining specific for procollagen type III (A, B), tenascin (C, D), and undulin (E, F). Procollagen type III specific staining is present in the lamina propria of normal mucosa, particularly underneath the superficial and crypt epithelium (A), and accumulates in the superficial collagenous layer of CC (B). Tenascin-specific staining is largely localized to a thin subepithelial layer in beneath the surface epithelium of normal colon mucosa (C) and stains the subepithelial matrix deposits of CC in their full thickness (D). Tenascin staining is virtually absent from the pericryptal area in CC and normal mucosa. In normal mucosa, undulin is found in small amounts along vessels in the lamina propria and decorating dense fibrillar bundles of the submucosal stroma (E). At variance with tenascin, undulin is absent from the subepithelial collagenous layer of CC (F). Combined immunohistology and *in situ* hybridization for smooth-muscle α -actin and procollagen $\alpha 1(\text{I})$ RNA in CC (cryostat sections; G, H). A proportion of the procollagen $\alpha 1(\text{I})$ RNA-positive cells underneath the superficial collagenous band is decorated by α -actin (arrows, CC case no. 8, Table 1). APAAP technique, original magnification, $\times 100$.

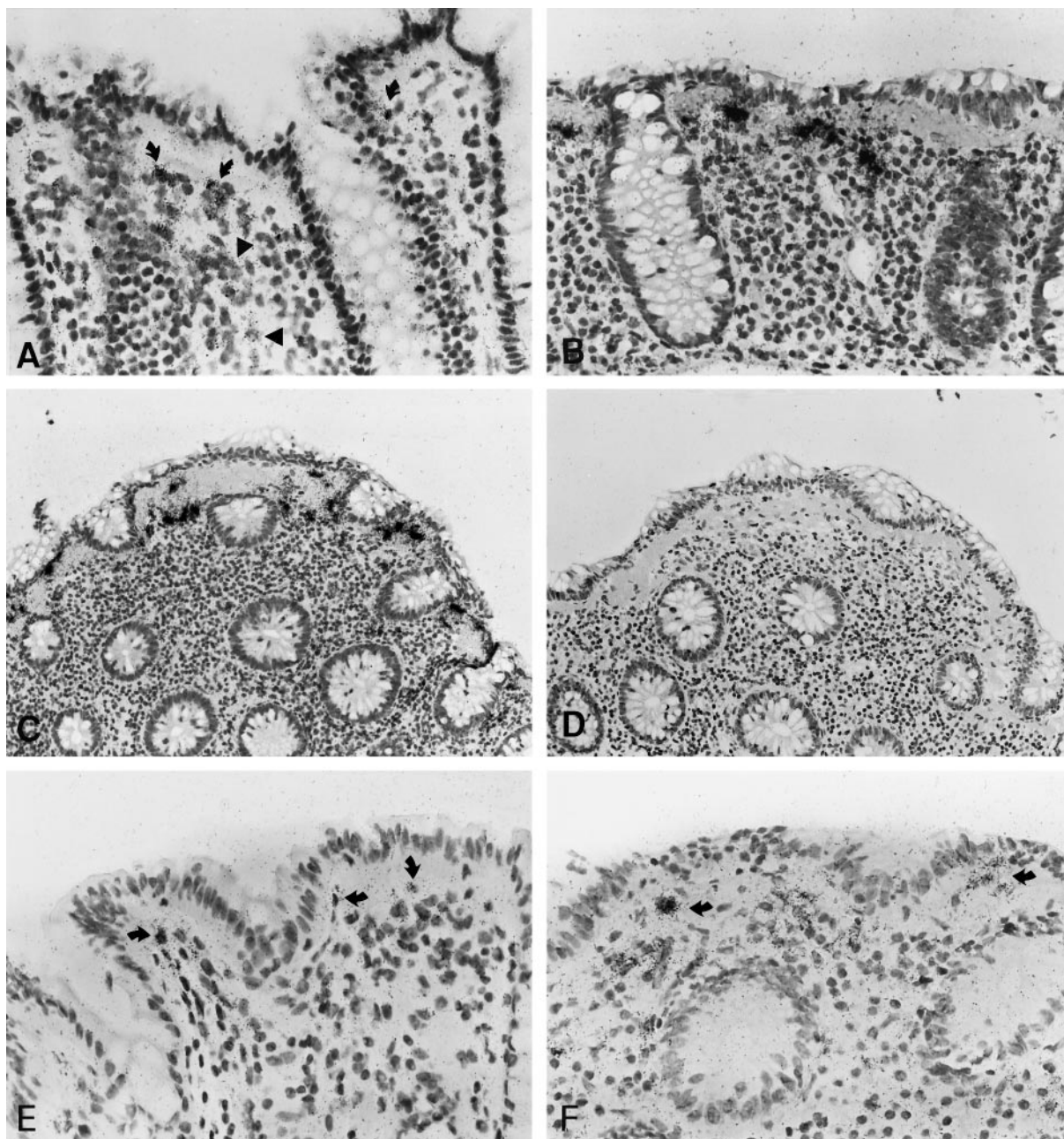


Figure 2. Patterns of procollagen $\alpha 1(I)$ and $\alpha 1(IV)$ expression in normal mucosa and CC as revealed by *in situ* hybridization with [^{35}S]-labeled RNA probes. B, C, and D represent adjacent serial sections of the same biopsy (case no. 8). Procollagen $\alpha 1(I)$ -specific labeling is displayed by cells of the subepithelial myofibroblast sheet (curved arrows) and the lamina propria (arrowheads) of normal colonic mucosa (A). In CC, the same probe reveals cells with high transcripts levels in an almost linear distribution underneath the surface epithelium (B and C). A weak background signal is seen after hybridization with the sense (control) probe (D). As compared to normal mucosa (E), elevated levels of procollagen $\alpha 1(IV)$ transcripts are found in myofibroblasts in a linear distribution underneath the surface epithelium in CC (arrows, case no. 6, F). Autoradiographic exposure time 14 days (A–D) and 36 days (E, F). Original magnification, $\times 100$ (A, B, E, F) and $\times 50$ (C, D).

transcript levels alternated with extended regions devoid of any specific signal. The peculiarity of this labeling pattern was evident when compared to procollagen $\alpha 1(I)$ -specific hybridizations on neighboring serial sections (Figure 5D).

TIMP-1 RNA expression was found in moderate levels underneath the surface epithelium of normal mucosa (Figure 5E). In the sSEMF sheet of CC biopsies, TIMP-1 expression was increased as well ($P < 0.025$), both with regard to the number of cells and the signal intensity per

cell (Figures 5F and 7). TIMP-1-positive cells showed an almost linear distribution in the sSEMF sheet including clusters of strongly labeled sSEMF cells. In serial sections, these clusters of strongly TIMP-1 RNA-expressing cells colocalized with clusters of cells displaying excess MMP-1 transcripts. Thus, the cellular distribution of TIMP-1 RNA appeared as a superimposition of both the procollagen $\alpha 1(I)$ and the MMP-1 characteristic patterns.

Labeling patterns for the above listed probes in ulcerative colitis and Crohn's disease specimens were as

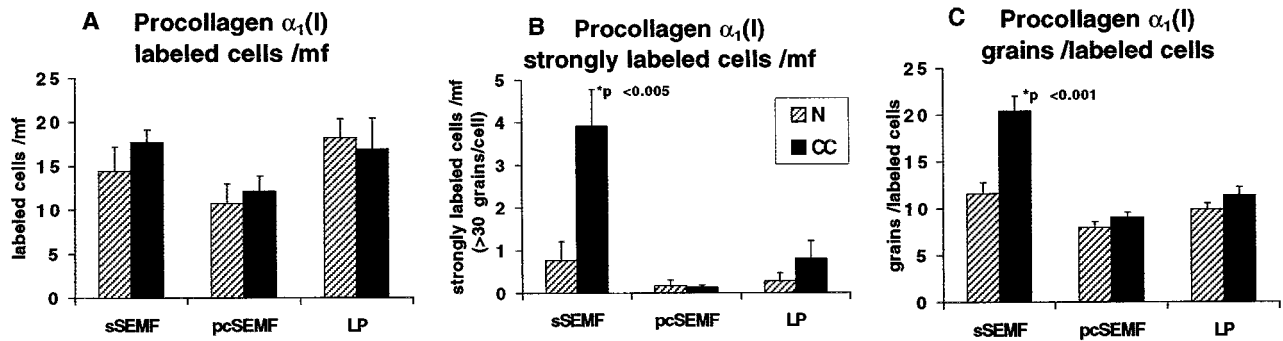


Figure 3. Semiquantitative evaluation of autoradiographic signals (silver grains) obtained with the procollagen $\alpha_1(I)$ probe in CC cases (closed bars) and in normal mucosa (hatched bars). The histograms represent the average number of all specifically labeled cells (A) and of the strongly labeled cells (>30 grains/cells; B) per microscopic field (mf) in the subepithelial myofibroblast (sSEMF) and pericryptal myofibroblast (pcSEMF) layer and the remainder of the lamina propria (LP). A third panel (C) displays the evaluation of the average number of silver grains per labeled cell. In comparison to normal control mucosa, the number of strongly labeled cells (B) and the average number of grains per labeled cell (C) were significantly increased in the sSEMF sheet of CC biopsies, whereas the relative number of labeled cells did not differ significantly. The evaluation is based on sections of all cases of collagenous colitis ($n = 12$) and normal controls ($n = 7$) which had been carried out in parallel through the entire *in situ* hybridization and autoradiographic procedures.

described previously^{27,31} and differed from those in CC by strong labeling of cells throughout the entire LP or even other layers of the bowel wall.

MMP-13 RNA transcripts were not detectable in CC specimens and in three cases of ulcerative colitis. The MMP-13 probe produced, however, a clear signal in few submucosal mesenchymal cells of 1 of 3 cases of Crohn's disease (Figure 6A), and a very intense autoradiographic signal in peritumoral mesenchymal cells in 4 of 9 squamous cell carcinomas of the head and neck region (Figure 6B). Hybridizations with sense (control) probes revealed a weak, nonspecific background labeling (Figure 2D).

Discussion

Collagenous colitis is a rare, only recently recognized intestinal disease entity of unknown etiology. At variance with other forms of inflammatory bowel diseases, the macroscopic and radiological findings are inconspicuous. Unlike Crohn's disease and ulcerative colitis, which may involve the entire bowel wall or the full thickness of the mucosa, respectively, CC displays deposits of interstitial ECM confined to a layer beneath the surface epi-

thelium. More recent evidence suggests that, upon fecal stream diversion, these deposits can be removed within periods as short as a few months,¹² pointing to a peculiar composition of the subepithelial ECM layer favoring a high turnover.

The enhanced immunostaining for tenascin in the absence of any undulin/collagen XIV-specific labeling within the ECM layer may provide an explanation for these findings. Tenascin is considered a marker of mesenchymal cell proliferation and migration, and undulin is usually found in dense and differentiated collagen matrices. The tenascin/undulin dichotomy has previously been described in the context of matrix deposits that are subject to rapid remodeling, such as the ECM of tumors or in hepatic fibrosis.^{22,32} Thus, an ECM that is subject to rapid turnover, regardless of the collagen composition, is made up of loosely connected interstitial collagen fibers with a high tenascin content. In contrast, matrices of long-standing scar tissue such as those present in Crohn's disease or cirrhotic liver display a high undulin but a low tenascin content.^{22,33-36} Apparently, undulin condenses interstitial ECM by binding and interconnecting collagen type I, III, V, and VI fibers and decorin.^{37,38} These staining patterns can explain some aspects of the above-mentioned clinical observations.

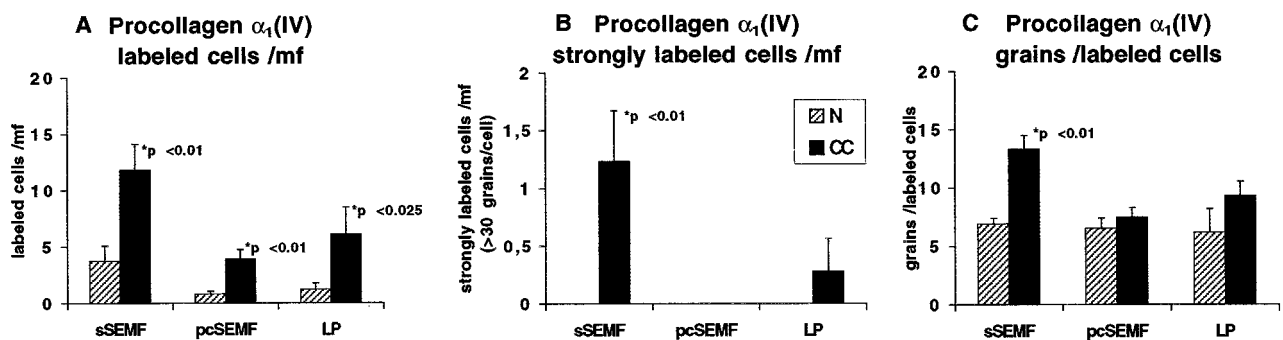


Figure 4. Semiquantitative evaluation of autoradiographic signals (silver grains) obtained with the procollagen $\alpha_1(IV)$ probe in CC cases (closed bars) and in normal mucosa (hatched bars) as outlined in Figure 3. In CC, the number of cells expressing procollagen $\alpha_1(IV)$ transcripts was significantly increased in all compartments of the lamina propria (A). In the sSEMF sheet, the number of strongly labeled cells (B) and the average signal intensity (C) were also significantly higher in CC as compared to normal mucosa.

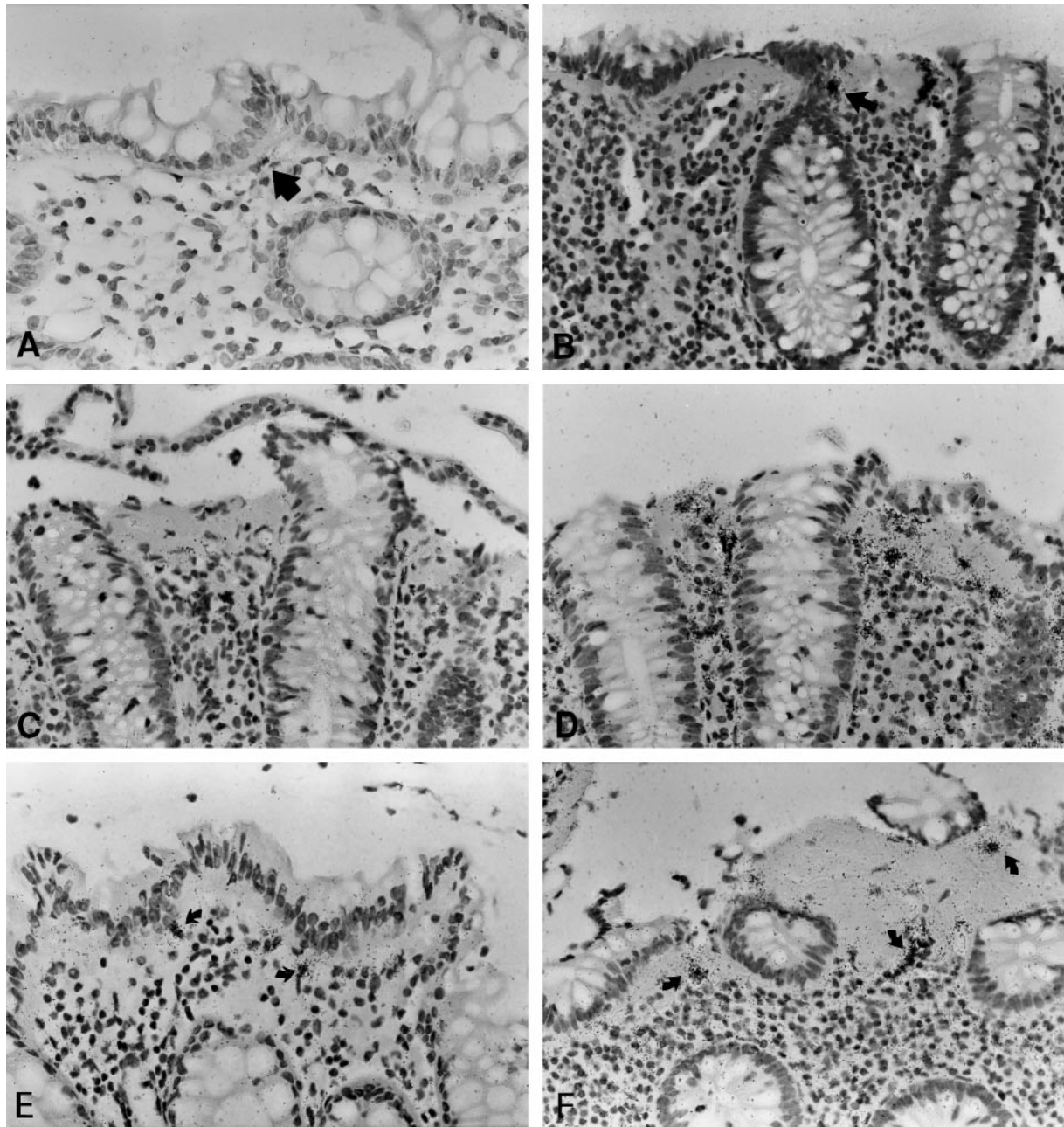


Figure 5. *In situ* hybridization with MMP-1, procollagen $\alpha 1(I)$ and TIMP-1 specific [^{35}S]-labeled RNA probes in normal mucosa and CC. In normal mucosa, MMP-1 expression is restricted to few weakly labeled cells (arrow, A). In CC, MMP-1 labeling is found in few cells clustered within the subepithelial myofibroblast sheet (arrow, B). These foci of expression are interrupted by long stretches of myofibroblasts without detectable MMP-1 RNA transcript levels (B, C). The paucity of these cells is particularly evident when compared to the procollagen $\alpha 1(I)$ specific signal on an adjacent serial section (D). TIMP-1 expression is found in a few cells of normal colon (arrows, E), but upregulated in sSEMF, pcSEMF and LP cells of CC specimens (arrows, F). CC cases no. 8 (B, F) and no. 1 (C, D). Autoradiographic exposure time 28 days (A–C), 14 days (D), and 25 days (E, F). Original magnification, $\times 100$ (A–F).

In situ hybridization demonstrated synthesis of procollagen $\alpha 1(I)$ and $\alpha 1(IV)$ at significantly increased transcript levels by resident sSEMF in all of our CC specimens. In case of procollagen $\alpha 1(I)$, an almost linear distribution of sSEMF cells with high transcript levels became evident. For procollagen $\alpha 1(IV)$, an essentially identical distribution of labeled cells was found, albeit with lower autoradiographic signal intensity requiring extended autoradiographic exposure. These observations indicate a continuously up-regulated synthesis of both

prototypic ECM molecules during the course of CC. Our findings are well in keeping with the recently published immunostaining patterns for collagens type I, III, IV, and VI.³⁹ The nonradioactive *in situ* hybridization patterns for the corresponding procollagens reported in the latter study are largely congruent with the distribution of (myo-) fibroblasts displaying >30 silver grains in our hybridizations.

In CC, up-regulated MMP-1 expression was restricted to few sSEMF cells in clustered, patchy distribution, con-

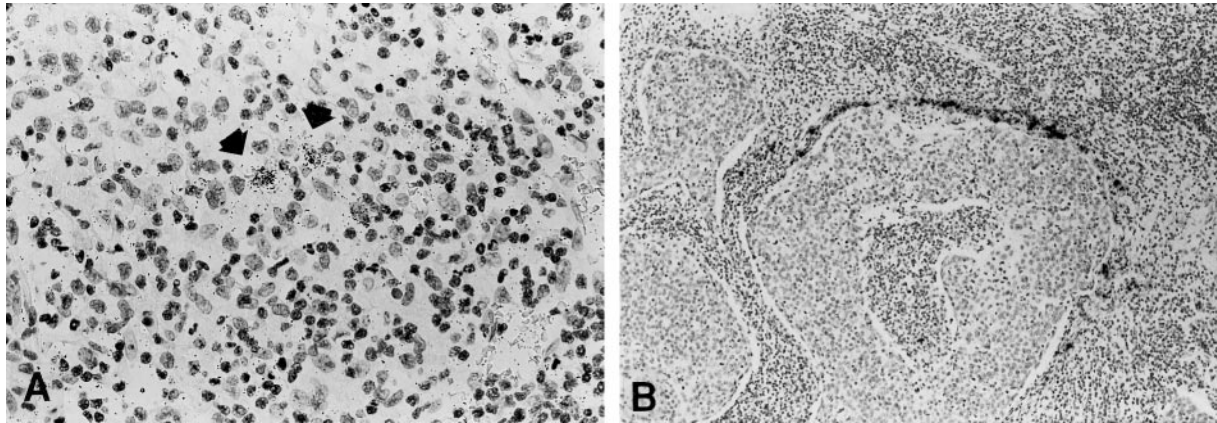


Figure 6. *In situ* hybridization with the MMP-13 specific [³⁵S]-labeled RNA probe in Crohn's disease and a tonsillar squamous cell carcinoma. A clear autoradiographic signal is seen in few submucosal mesenchymal cells in Crohn's disease (A), whereas a very intense signal is found in the peritumoral mesenchymal cells in the squamous cell carcinoma (B). Autoradiographic exposure time 36 days. Original magnifications, ×100 (A) and ×40 (B).

trasting with ECM molecules, whereas its physiological inhibitor, TIMP-1, displayed a partially superimposable expression pattern. Degradation of the various ECM components is a function of an array of enzymes with diverse, partially overlapping substrate specificities.¹⁴⁻¹⁶ The observation of MMP-1 RNA suggests local enzyme synthesis, but does not prove the presence of active enzyme. Similarly, the immunostaining pattern for MMP-1 was inconclusive as it does not distinguish among zymogen, active, and inhibited forms of the enzyme. TIMP-1 expres-

sion seems to occur independently of MMP-1, as demonstrated by the weak, linear distribution of TIMP-1 RNA along the subepithelial collagenous layer. In areas with expression of MMP-1, MMP-1 protein, if activated, may be inhibited by TIMP-1 and other inhibitors unless, at balance, MMP-1 molecules prevail. Local and temporary changes of this balance in favor of active MMP-1 may then result in a rapid increase of the fibrolytic potential.

The expression of MMP-13 has previously been documented in ulcerative colitis and Crohn's disease at the

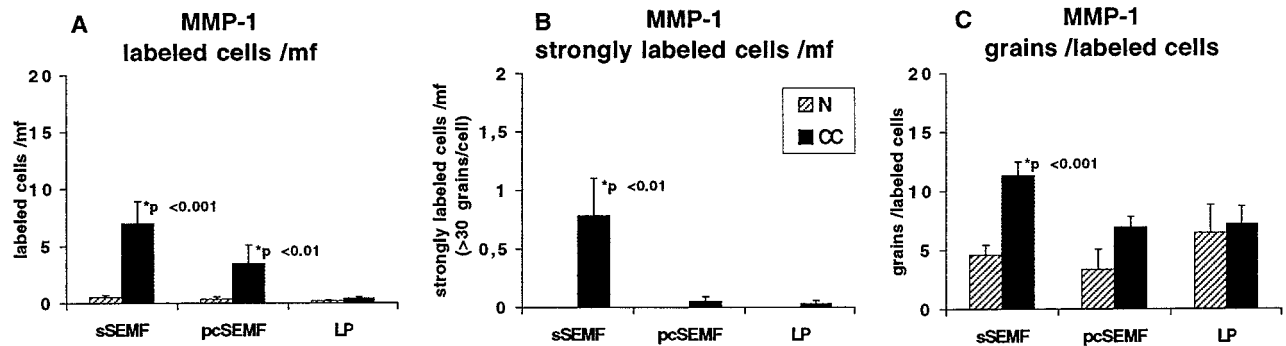


Figure 7. Semiquantitative evaluation of autoradiographic signals (silver grains) obtained with the MMP-1 probe in CC cases (closed bars) and in normal mucosa (hatched bars) as outlined in Figure 3. In comparison to normal control mucosa, the number of MMP-1-expressing cells in general (A) and of cells with high specific transcript levels (B), and the average number of grains per labeled cell is (C) increased in CC. Note that these cells display a clustered distribution (Figure 5).

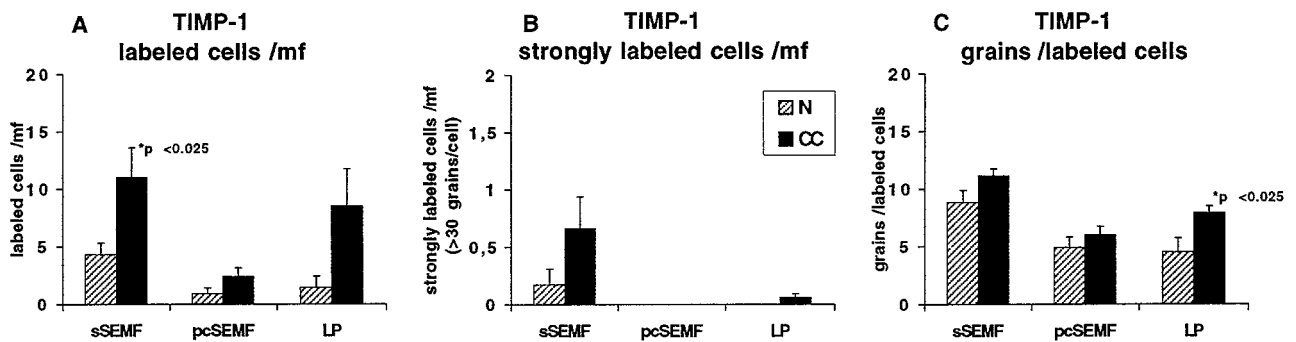


Figure 8. Semiquantitative evaluation of autoradiographic signals (silver grains) obtained with the TIMP-1 probe in CC cases (closed bars) and in normal mucosa (hatched bars) as outlined in Figure 3. In comparison to normal control mucosa, the number of TIMP-1-expressing cells in general (A) and of cells with high specific transcript levels (B), and the average number of grains per labeled cell (C) is increased in CC. The relative frequency of sSEMF cells with strong MMP-1- and TIMP-1-specific labeling is similar. Note that these cells display a clustered distribution (Figure 5).

base of ulcerations and proved to be absent within intact mucosa.⁴⁰ MMP-13 was also found in squamous cell carcinomas of the head and neck region.⁴¹ Our *in situ* hybridization results confirm these patterns. In our CC biopsies, however, MMP-13 transcript levels were below the threshold of detection, suggesting that this collagenase may not play a major role in nonulcerating intestinal lesions such as CC. This finding and the observation of only few and scattered granulocytes as potential sources of MMP-8 (neutrophil collagenase) in the LP underline the suggested central role of MMP-1 expression for interstitial collagen type I degradation in CC. A future approach to the study of collagen degradation in CC should analyze the MMP-1 activation cascade including expression patterns of plasmin-generating molecules and stromelysins (MMP-3 and -10), which serve as co-activators of MMP-1.^{15,16}

MMP expression in inflammatory bowel diseases has recently been studied by several groups reporting enhanced expression of MMP-1, -3, -7, -9, -10, -12, and -13 in ulcerative lesions of Crohn's disease, ulcerative and ischaemic colitis, and in wound healing.^{40,42-44} However, analysis of such lesions does not warrant conclusions about CC, because CC differs from the former lesions and entities by its peculiar clinical presentation and histology as a non-ulcerating, potentially reversible intestinal disorder.

In summary, the patterns of procollagen $\alpha 1(I)$ -specific labeling in CC, which is accompanied by an enhanced deposition of tenascin in the absence of undulin, indicate synthesis of an immature, loosely connected interstitial collagenous matrix that may be rapidly degraded. The restricted occurrence of MMP-1 transcripts in the presence of increased TIMP-1 and interstitial procollagen expression, however, suggests a local imbalance of fibrogenesis and fibrolysis in CC, resulting in the characteristic, linear ECM deposits. Thus, future therapeutic strategies may be directed at influencing the levels of TIMP-1 produced by sSEMF cells.

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