

Increased Expression of the 72-kd Type IV Collagenase in Prostatic Adenocarcinoma

Demonstration by Immunohistochemistry and *in Situ* Hybridization

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The expression of the 72-kd type IV collagenase has been implicated as an important factor in determining the invasive potential of malignant tumors. Using immunohistochemistry and nonisotopic *in situ* hybridization, type IV collagenase expression was assessed in benign and malignant prostatic tissue obtained from 117 surgical and autopsy specimens. Diffuse strong staining for type IV collagenase mRNA and protein was identified in the malignant cells of more than 85% of prostatic adenocarcinomas and the dysplastic cells of high grade prostatic intraepithelial neoplasia. Benign hyperplastic epithelium showed moderate expression in basal cells and mild expression in secretory cells. The qualitative patterns of type IV collagenase expression in prostatic epithelium at the protein and mRNA levels in individual cases were identical. There was no correlation between the level of type IV collagenase expression and either tumor grade or stage. In 10% of adenocarcinomas, focal mild to moderate stromal cell immunoreactivity was present but mRNA was not detectable in the stromal compartment in any case. The enhanced expression of type IV collagenase in dysplastic epithelium and prostatic adenocarcinoma suggests it contributes to the development of the invasive phenotype. The vast majority of the enzyme present in these tumors is synthesized by malignant cells and its production by stromal cells is negligible. (Am J Pathol 1994, 144:585-591)

(MMPs) comprise a group of related enzymes that appear to have an important role in this process.^{1,2} The 72-kd type IV collagenase (cIVase) is an MMP that cleaves type IV collagen, the main structural component of the basement membrane. Because the basement membrane is the first matrix compartment to be breached during tissue invasion by carcinomas, increased cIVase activity may represent a critical early event in the development of the invasive phenotype. The cIVase activity of tumor cells has been shown to correlate with their ability to cross basement membranes *in vitro* and produce metastases in animal models.³ In several human tumors including breast,⁴ colonic,⁵ ovarian,⁶ and hepatocellular carcinomas,⁷ cIVase expression has positively correlated with tumor invasiveness.

The relative contribution of neoplastic, stromal, and inflammatory cells to cIVase activity in human tumors remains unclear. By immunohistochemistry, cIVase expression has been shown to be largely restricted to malignant cells in breast, colonic, and hepatocellular carcinomas.^{4,5,7} However, *in situ* hybridization (ISH) studies of skin, lung, and colorectal carcinomas have localized cIVase transcripts predominantly in stromal cells,⁸⁻¹⁰ suggesting the mesenchymal compartment is the primary site of cIVase synthesis in these tumors.

The factors that determine the heterogeneous metastatic potential of prostatic adenocarcinoma are poorly understood, although it has been shown that progressive basement membrane loss occurs with increasing Gleason grade,¹¹ which in turn correlates with invasive potential. cIVase has previously been shown to be present in prostatic carcinomas and absent in normal prostates by Northern analysis.¹² We

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Degradation of the extracellular matrix is required for tumor invasion. The matrix metalloproteinases

have recently demonstrated enhanced expression of *clVase* in prostatic adenocarcinoma relative to benign prostatic epithelium using immunohistochemistry.¹³ In this study we have used a combination of nonisotopic ISH and immunohistochemistry to extend the analysis of *clVase* in prostatic adenocarcinomas of varying grades and stages as well as normal and hyperplastic prostates. The results demonstrate that *clVase* expression is up-regulated at the mRNA and protein level in dysplastic and malignant epithelial cells but is not synthesized in significant amounts by tumor stroma.

Materials and Methods

Tissue

A retrospective immunohistochemical analysis of *clVase* expression was performed on 95 archival cases (including the 50 cases previously reported¹³) comprising 10 normal prostates (from autopsies on men under the age of 45 years), 29 cases of prostatic hyperplasia, and 56 cases of prostatic adenocarcinoma of varying grades and stages. All cases had been fixed in 10% neutral-buffered formalin and tissue sections were cut onto poly-L-lysine-coated slides.

In addition, immunohistochemistry and mRNA ISH were conducted on benign and malignant prostatic tissue derived from 22 prospectively gathered fresh radical prostatectomy specimens. Tissue blocks were fixed in 4% paraformaldehyde in phosphate-buffered saline, pH 7.0, for 4 hours at room temperature, processed using standard techniques and embedded in paraffin. Sections were cut onto 3-aminopropyltriethoxysilane (Aldrich Chemical, Milwaukee, WI)-coated slides.

Antibodies

Two polyclonal affinity-purified antitype IV collagenase antibodies were a generous gift of Dr. W. G. Stetler-Stevenson of the National Cancer Institute, NIH. Antibodies KLH 31 and KLH 48 were raised against synthetic peptides corresponding to the 17 NH₂-terminal amino acids of type IV procollagenase and the putative metal binding domain of *clVase*, respectively.⁴

Immunohistochemistry

Tissue sections were deparaffinized then rehydrated and blocked with 0.45% H₂O₂ in methanol for 30 minutes followed by normal goat serum (Dako Labs, Car-

pentaria, CA) diluted 1/20 for 20 minutes. The primary antibodies KLH 31 and KLH 48, diluted to 1 and 15 mg/ml, respectively, were applied and incubated at 4 C overnight. A 15-minute preincubation at 37 C was performed for KLH 48. A secondary biotinylated antibody (goat anti-rabbit, Zymed, San Francisco, CA), diluted 1/200, and peroxidase-conjugated streptavidin (Dakopatts, Glostrup, Denmark), diluted 1/300, were applied sequentially for 30 minutes, followed by 0.33 mg/ml 3-3' diaminobenzidine tetrahydrochloride (Sigma, St. Louis, MO) in 50 nM Tris, pH 7.6, containing 0.045% H₂O₂ for 8 minutes. The sections were counterstained lightly with hematoxylin. Phosphate-buffered saline was used to dilute the antibodies and to rinse sections between steps. The primary antibodies were replaced by normal rabbit serum for negative control slides. Immunoreactivity was assessed using routine light microscopy and intensity graded from 0 to 4+.

Synthesis of Digoxigenin-Labeled cRNA Probes

Sense and antisense cRNA probes were transcribed from a polymerase chain reaction (PCR)-amplified segment of the *clVase* gene using a previously described technique.¹⁴ Briefly, a 175 base sequence of the gene including nucleotides 2250-2425^{15,16} was amplified from crude genomic DNA using 20-mer oligonucleotide sense and antisense primers. This sequence resides within a larger cRNA probe that was used previously for ISH.⁹ The SP6 and T7 RNA polymerase promoters were incorporated into the amplified DNA by including these sequences in the 5' termini of the sense and antisense primers, respectively. The PCR reaction mixture included: 40 μ l of diethylpyrocarbonate (DEPC)-treated water, 10 μ l of PCR buffer (650 mM Tris, pH 8.0, 20 mM MgCl₂, 160 mM ammonium sulfate, and 0.7% β -mercaptoethanol), 16 μ l of 800 mM deoxyribonucleotide triphosphate mixture, 1 μ l (20 pmol) of each primer, 30 μ l of crude DNA extract, and 0.5 μ l (2.5 U) Taq polymerase (Perkin-Elmer Cetus, Norwalk, CT). The DNA extract was obtained by boiling human blood in an equal volume of water for 10 minutes. After centrifugation, the supernatant was used as the template for the PCR. The reaction was performed for 30 cycles using a DNA thermal cycler (Perkin-Elmer Cetus) using the following profile: 94 C for 2 minutes, 50 C for 2 minutes, and 72 C for 1 minute lengthened by 5 seconds in each successive cycle. Analysis of the PCR product by polyacrylamide gel electrophoresis

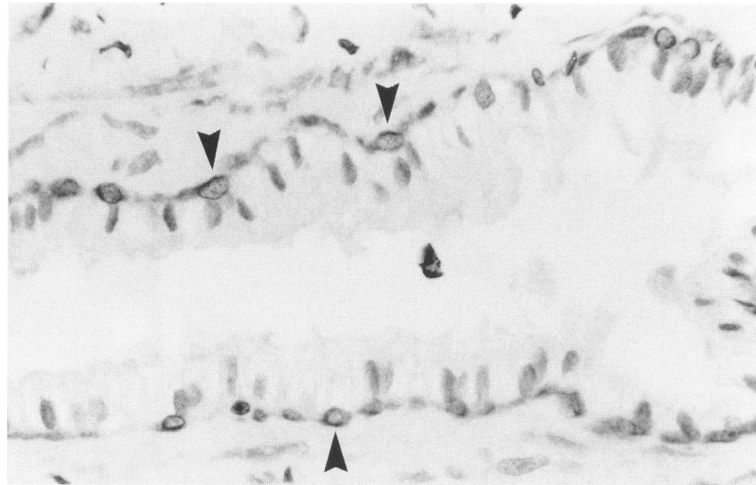


Figure 1. Normal prostate gland showing moderate immunoreactivity for cIVase in basal cells (arrowheads) and minimal immunoreactivity in secretory epithelium ($\times 400$, hematoxylin counterstain).

demonstrated a single, discrete DNA band of predicted size (data not shown).

Digoxigenin-labeled cRNA probes were transcribed from the amplified template by incubation of the following reaction mixture at 37 C for 2 hours: 1 μ l PCR product, 2 μ l transcription buffer (400 mM Tris-HCl, pH 8.0, 60 mM MgCl₂, 20 mM spermidine, 100 mM NaCl), 2 μ l ribonucleotide triphosphate mixture

(10 mM ATP, 10 mM GTP, 10 mM CTP, 3.5 mM digoxigenin-11-UTP, Boehringer Mannheim, Mannheim, Germany, 6.5 mM UTP), 40 U of SP6 or T7 RNA polymerase (Bio/Can Scientific, Inc., Mississauga, Ontario, Canada), and DEPC-treated water to a final volume of 20 μ l. DNA was digested with 20 U of RNase-free DNase (Pharmacia, Uppsala, Sweden) for 20 minutes at 37 C and the DNase inactivated by

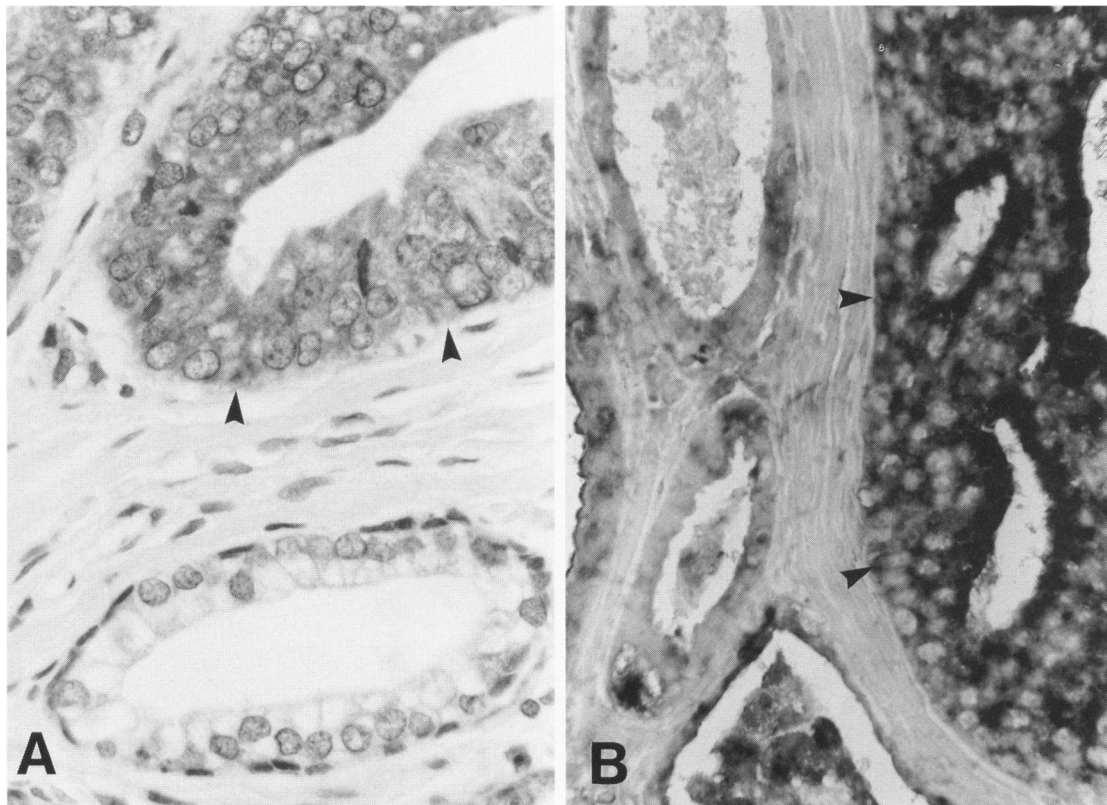
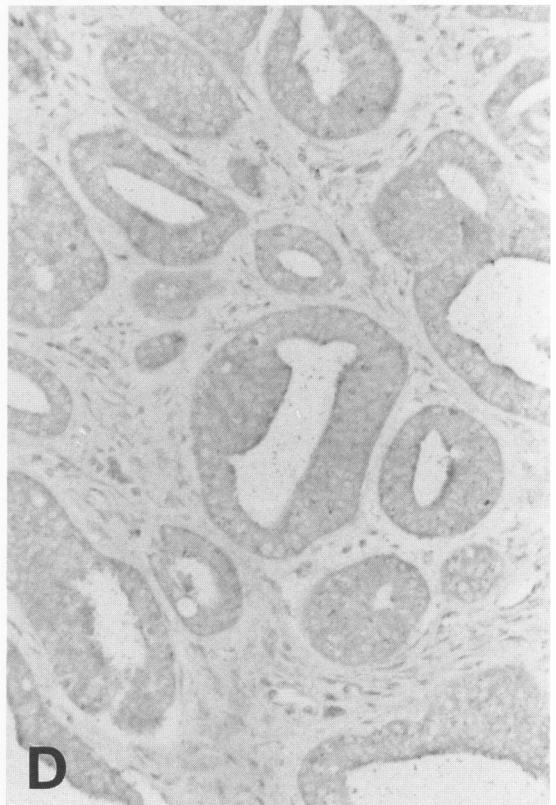
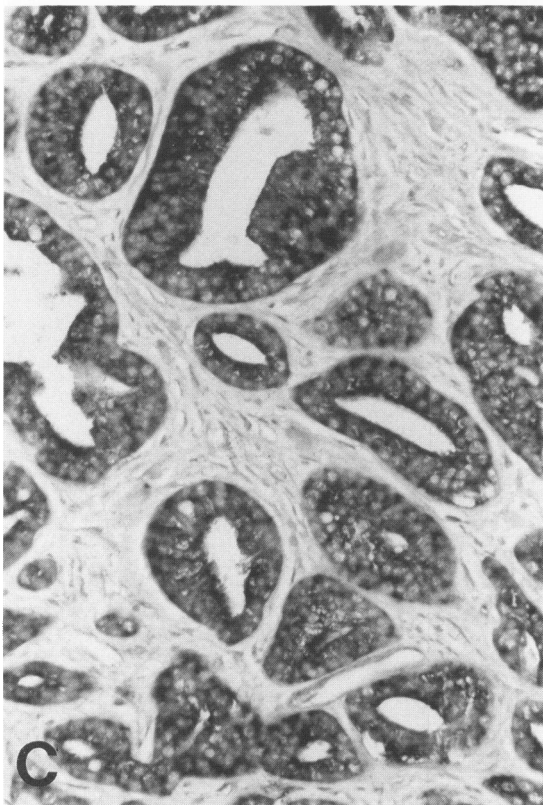
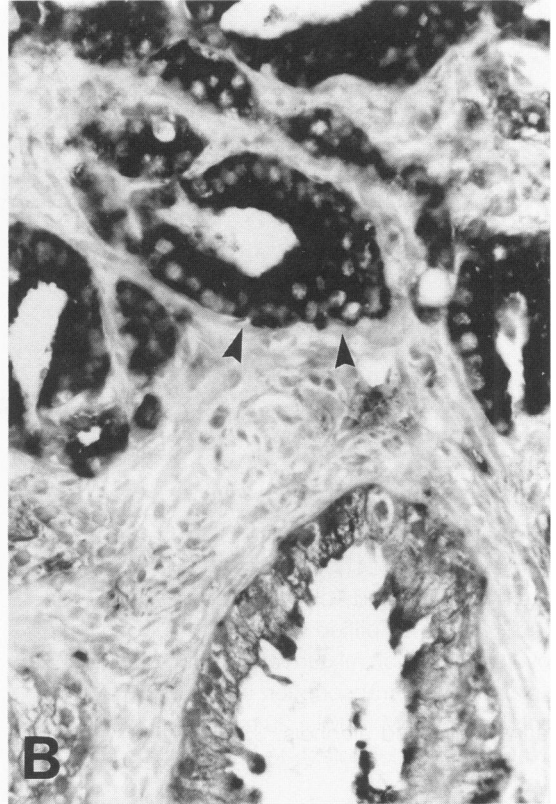
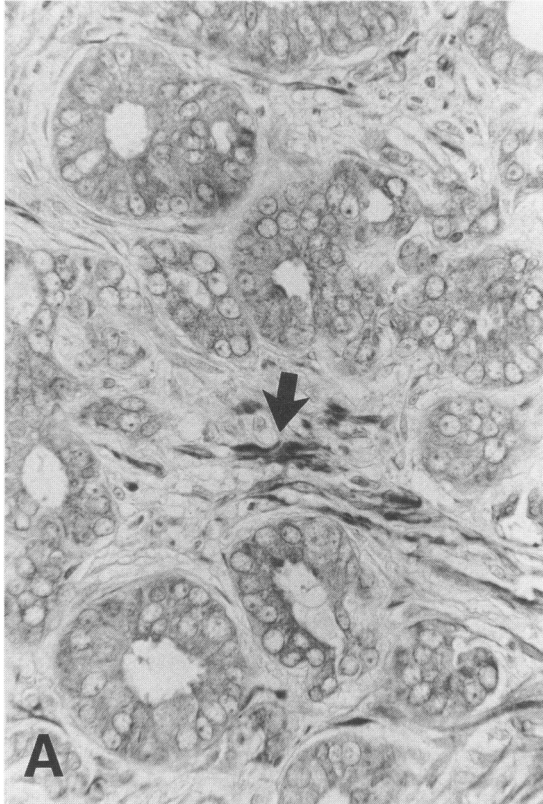


Figure 2. High grade prostatic intraepithelial neoplasia (arrowheads) shows high levels of cIVase expression by both immunohistochemistry (A) and ISH (B). Adjacent stroma and benign glands show minimal expression (A, $\times 250$, hematoxylin counterstain; B, $\times 200$, methyl green counterstain).



heating to 65 C for 5 minutes. The probes were precipitated at -20°C overnight after addition of 1/8 volume of 4 M LiCl and three volumes of cold absolute ethanol. The precipitate was washed three times in 75% ethanol, dissolved in DEPC-treated water, and the probe concentration determined by measuring optical density at 260 nm. To verify the identity of antisense probe. Northern blots of the obtained transcripts were hybridized under high stringency conditions with a [δ - 32] PATP end-labeled 18-mer oligonucleotide complementary to an internal sequence of the antisense transcript. This internal oligonucleotide showed specific hybridization to the antisense transcript (data not shown).

ISH

Tissue sections were deparaffinized, rehydrated, washed in 0.2 M HCl for 10 minutes, and rinsed in nanopure water. They were then washed with 0.25% acetic anhydride in buffer 1 (150 mM NaCl, 100 mM Tris-HCl, pH 7.5) for 10 minutes then rinsed in buffer 1 and washed in 0.2% glycine in buffer 1. After dehydration in graded alcohols, 30 μl of hybridization buffer (50 mM Tris, pH 7.6, 1 mM ethylenediamine tetraacetic acid, 0.02% Ficoll, 0.02% polyvinylpyrrolidone, 500 $\mu\text{g}/\text{ml}$ yeast tRNA [Boehringer Mannheim], 10 mM dithiothreitol, 0.3 M NaCl, 10% dextran sulfate, and 50% formamide) containing 500 $\mu\text{g}/\text{ml}$ boiled sheared salmon sperm DNA (Sigma) was applied per cm^2 of tissue section. The slides were placed in a metal pan that was floated in a 95 C water bath for 10 minutes. Excess prehybridization solution was drawn off the slides and 30 μl of hybridization buffer containing 20 $\mu\text{g}/\text{ml}$ denatured probe was applied. The hybridization was performed at 50 C for 2 hours in a humidified box. In addition to using sense probes, negative controls included elimination of the antisense probe and pretreatment of sections with RNase A (Sigma) before hybridization.

After hybridization, sections were washed sequentially with 2 \times standard saline citrate (SSC) (1 \times SSC is 0.15 M NaCl, 0.015 M sodium citrate) for 5 minutes and 0.2 \times SSC for 30 minutes at room temperature. Sections were blocked with 1% casein (U.S. Biochemical Corp., Cleveland, OH) in buffer 1 for 30 minutes and then 100 μl of alkaline phosphatase-conjugated antidigoxigenin F(ab) fragments (Boehringer Mannheim) diluted 1:1000 in

buffer 1 was applied for 2 hours. The slides were washed twice in buffer 1 for 15 minutes and rinsed in buffer 2 (100 nM NaCl, 20 nM MgCl_2 , 100 nM Tris-HCl, pH 9.5). The sections were covered with 100 μl chromogen substrate solution (45 μl of 90 mM nitroblue tetrazolium salt in 70% [volume/volume] dimethylformamide and 35 μl of 120 mM 5-bromo-4-chloro-3-indolylphosphate toluidinium salt in 100% dimethylformamide in 10 ml buffer 2) containing 0.5 mM levamisole and incubated overnight at room temperature. The sections were rinsed well with nanopure water, counterstained with 2% chloroform-extracted methyl green, and coverslipped with an aqueous mounting medium.

Results

Immunohistochemistry

Most (over 75%) basal epithelial cells in all cases of normal and hyperplastic prostatic tissue showed moderate (3+) cytoplasmic immunoreactivity for cI-Vase (Figure 1). Acinar and ductal secretory epithelial cells demonstrated immunostaining of lower intensity (1 to 2+), although these cells did show moderate immunoreactivity in some cases. Atrophic epithelium showed absent to focal slight (0 to 1+) immunoreactivity. Endothelium and transitional epithelium lining the urethra and distal prostatic ducts showed moderate immunostaining. The stroma in normal prostates and most cases of hyperplasia showed no immunoreactivity. However, in 10% of cases of hyperplasia, focal areas of moderate cytoplasmic immunoreactivity in stromal cells were identified, a finding that was more pronounced with antibody KLH 31 than KLH 48.

In 96% of the archival cases of prostatic adenocarcinoma and 85% of paraformaldehyde-fixed cases, tumor cells showed uniformly moderate to strong (3 to 4+) cytoplasmic immunoreactivity. Tissue fixation in the negative cases was suboptimal and may have contributed to the lack of staining. The overall staining intensity was less in paraformaldehyde-fixed than formalin-fixed tissue. High grade prostatic intraepithelial neoplasia (PIN) showed strong immunoreactivity similar to that seen in invasive cancers and significantly more intense than adjacent nondysplastic ductal epithelium (Figure 2A). There was no

Figure 3. A: Prostatic adenocarcinoma demonstrates diffuse immunoreactivity for cI-Vase in malignant epithelial cells and focal immunoreactivity in stromal cells (arrow). B: ISH localizes very abundant cI-Vase mRNA in the epithelial compartment of prostatic adenocarcinoma (arrowheads) compared with minimal expression in adjacent benign epithelium. C: ISH in the case depicted in (A) shows cI-Vase mRNA is restricted to neoplastic epithelium with no stromal expression apparent. D: ISH with sense cRNA shows no staining (A, $\times 200$, hematoxylin counterstain; B to D, $\times 200$, methyl green counterstain).

correlation between cIVase immunoreactivity and tumor grade or stage, although a trend to decreased expression was seen in some very poorly differentiated carcinomas. Tumor infiltrating periprostatic tissue, and foci of metastatic carcinoma in lymph nodes and soft tissue, all showed immunoreactivity of similar intensity to tumor within the prostate.

Prostatic stromal cells in malignant prostates were generally negative. However, in 10% of carcinomas there were focal areas of slight to moderate cytoplasmic immunostaining of stromal cells with KLH 31 but not KLH 48 (Figure 3A). Mesenchymal cells associated with foci of metastatic tumor showed no immunoreactivity.

ISH

There was a strong positive correlation between the ISH and immunohistochemistry results in both carcinomas and benign epithelium. High levels of cIVase mRNA, as indicated by dense cytoplasmic staining of malignant cells, were apparent in 85% of prostatic carcinomas. There was no hybridization in two highly anaplastic carcinomas and in two cases in which tissue fixation was suboptimal. Apart from a trend to decreased expression in very poorly differentiated tumors, no correlation of mRNA level with grade was found. Metastatic tumor in the single pelvic lymph node studied and foci of locally invasive tumor in the urinary bladder wall in a second case showed similar intensities of staining to the corresponding primary tumors within the prostate. There was no correlation between the ISH results and pathological stage. Foci of PIN also expressed abundant cIVase mRNA, whereas hyperplastic epithelium showed mild to moderate positivity and atrophic glands were generally negative (Figure 2B). The ISH signal intensity was clearly lower in benign prostatic epithelium when compared with prostatic adenocarcinoma.

ISH detected no cIVase mRNA in any stromal cells in either benign or malignant prostates, including those cases in which focal stromal cell immunoreactivity for cIVase was identified (Figure 3).

ISH performed using the sense probe as a negative control showed minimal background staining (Figure 3D). Treatment of tissue sections with RNase A before hybridization completely eliminated the signal (data not shown). The specificity of the ISH is also supported by the excellent correlation with the immunohistochemistry performed on the same tissue samples. The two anaplastic carcinomas in which ISH was negative showed weak stain-

ing by immunohistochemistry and two cases in which fixation appeared poor show minimal staining by both methods. Otherwise, all cases were positive by both techniques.

Discussion

cIVase is thought to play a role in the proteolysis of basement membrane by malignant cells that occurs early in the development of invasive tumors. The results of the current study demonstrate that cIVase expression is enhanced at both the protein and mRNA levels in prostatic adenocarcinoma, consistent with such a role for cIVase in this cancer. High grade PIN was also found to display markedly elevated levels of cIVase. This probable precursor of prostatic adenocarcinoma¹⁷ is analogous to atypical epithelial hyperplasia and *in situ* carcinoma of the breast in which increased levels of cIVase have also been found.⁴ These data indicate that overexpression of cIVase is an early event that precedes the invasion of carcinomas in general and prostatic adenocarcinomas in particular.

Similar levels of expression of cIVase were found in well differentiated tumors restricted to the prostate and poorly differentiated tumors with extensive local invasion and distant metastases. This contrasts with the findings of studies on hepatocellular carcinoma⁷ and colonic carcinomas^{5,9} in which increased cIVase expression correlated with higher tumor grades and stages. Other variables are presumably involved in determining the full invasive phenotype of prostatic adenocarcinomas. A tipping of the balance between cIVase activation and inhibition could in part explain the heterogeneous invasive potential in prostatic carcinomas that show similar levels of cIVase protein and mRNA.^{18,19} Other possible factors could include differential expression of other matrix metalloproteases or molecules involved in cell adhesion and migration.

Further evidence that elevated cIVase levels alone do not confer invasive behavior is provided by the finding of moderate levels of cIVase expression in basal cells. This probably reflects a greater role in normal ongoing basement membrane metabolism for these cells compared with the secretory cells that generally showed lower levels of cIVase expression. An analogous observation has been made in normal breast tissue in which the myoepithelial cells show higher levels of cIVase expression when compared with the secretory epithelium.⁴

The ISH data from this study suggest that cIVase production in prostatic adenocarcinoma occurs ex-

clusively in the neoplastic cells. It is possible that *clVase* transcripts are present in very low abundance in prostatic stromal cells and that our ISH technique was inadequately sensitive to detect them. However, a recent study using Northern blot analysis has also suggested that *clVase* mRNA is largely restricted to the neoplastic cells in prostatic adenocarcinoma.²⁰ In previous studies of skin, lung, and colorectal carcinomas,⁸⁻¹⁰ ISH localized *clVase* mRNA principally in tumor stromal cells. Immunoreactive *clVase* was identified primarily in neoplastic cells, suggesting that *clVase* synthesized and secreted by tumor stroma was secondarily taken up by malignant cells. In prostatic adenocarcinoma, however, malignant prostatic epithelial cells express both abundant *clVase* mRNA and protein, demonstrating they are producing the enzyme. Stromal cells in prostatic tumors are not a significant source of *clVase*.

In summary, prostatic adenocarcinomas showed increased expression of *clVase*, which is produced almost exclusively by neoplastic cells. Similar levels of *clVase* are present in high grade PIN and invasive carcinomas of varying grades and stages, indicating that although *clVase* may play a role in prostate cancer invasion, other factors must be involved in determining the full invasive potential of this tumor.

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