Prostate-specific Membrane Antigen Expression in Normal and Malignant Human Tissues

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ABSTRACT

Prostate-specific membrane antigen is a type II membrane protein with folate hydrolase activity produced by prostatic epithelium. The expression of this molecule has also been documented in extraprostatic tissues, including small bowel and brain. In the present study, an extensive immunohistochemical analysis was performed on a panel of well-characterized normal and malignant human tissues to further define the pattern of prostate-specific membrane antigen (PSMA) expression.

Detectable PSMA levels were identified in prostatic epithelium, duodenal mucosa, and a subset of proximal renal tubules. A subpopulation of neuroendocrine cells in the colonic crypts also exhibited PSMA immunoreactivity. All other normal tissues, including cerebral cortex and cerebellum, had undetectable levels of PSMA. Thirty-three of 35 primary prostate adenocarcinomas and 7 of 8 lymph node metastases displayed tumor cell PSMA immunostaining. Eight of 18 prostate tumors metastatic to bone expressed PSMA. All of the other nonprostatic primary tumors studied had undetectable PSMA levels. However, intense staining was observed in endothelial cells of capillary vessels in peritumoral and endotumoral areas of certain malignancies, including 8 of 17 renal cell carcinomas, 7 of 13 transitional cell carcinomas, and 3 of 19 colon carcinomas.

Extraprostatic PSMA expression appears to be highly restricted. Nevertheless, its diverse anatomical distribution implies a broader functional significance than previously suspected. The decrease in PSMA immunoreactivity noted in advanced prostate cancer suggests that expression of this molecule may be linked to the degree of tumor differentialization. The neoexpression of PSMA in endothelial cells of capillary beds in certain tumors may be related to tumor angiogenesis and suggests a potential mechanism for specific targeting of tumor neovascularature.

INTRODUCTION

PSMA1 is a Mr 100,000 type II membrane protein consisting of 750 amino acids (1, 2). Although PSMA exhibits in vitro neuropeptidase activity (3), its function in vivo has not been fully elucidated. The PSMA gene is located on chromosome 11 (4, 5). To date, two molecular forms of the protein, designated PSMA and PSMA', have been identified (6). Monoclonal antibody 7E11-C5 (designated CYT 351) recognizes an intracellular epitope of PSMA (7) and has been previously utilized for its immunohistochemical detection (6, 8). In addition, a radioimmunoconjugate of the 7E11-C5 antibody (designated CYT 356) is in use as an imaging agent for prostate tissues (7, 9–12). The clinical utility of PSMA as a marker for prostate disease depends on the pattern of its expression in vivo. An extensive immunohistochemical evaluation was performed to further characterize the pattern of PSMA expression in normal human tissues. A selected panel of primary and metastatic prostate carcinomas was also evaluated as well as a representative cohort of other epithelial malignancies, including renal cell carcinoma, bladder transitional cell carcinoma, and colon carcinoma.

MATERIALS AND METHODS

Tissues. Normal and neoplastic formalin-fixed, paraffin-embedded tissue samples were obtained from the Department of Pathology at the Memorial Sloan-Kettering Cancer Center. Thirty-five primary prostate adenocarcinoma specimens were evaluated as well as 8 metastases to lymph nodes and 18 metastases to bone. Table 1 summarizes the clinical characteristics of these tumors in terms of pathological stage and prior treatment. Additional primary tumors included 17 renal cell carcinomas, 13 bladder transitional cell carcinomas, and 19 colon adenocarcinomas.

Antibodies. Mouse monoclonal antibody CYT-351 (clone 7E11-C5) (Cytogen, Princeton, NJ) was used as the primary antibody. This clone is derived from the original hybridoma reported by Horoszewicz et al. (7). Secondary antibodies consisted of biotinylated horse anti-mouse polyclonal IgG (Vector Laboratories, Inc., Burlingame, CA). The proper concentration of each reagent was determined by titration experiments prior to staining.

Immunohistochemistry. An avidin-biotin peroxidase method was used. Sections were deparaffinized, and endogenous peroxidase activity was blocked in 1.0% hydrogen perox-

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The abbreviation used is: PSMA, prostate-specific membrane antigen.
Table 1: Primary prostate tumors: stage and prior treatment

<table>
<thead>
<tr>
<th>Stage</th>
<th>ADT*</th>
<th>RT</th>
<th>Chemo</th>
<th>ADT/RT</th>
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<tbody>
<tr>
<td>P,N0</td>
<td>1</td>
<td>14</td>
<td></td>
<td></td>
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<td>M+</td>
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<td>1</td>
<td>5</td>
<td>3</td>
<td>2</td>
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</tbody>
</table>

*ADT, androgen deprivation therapy; RT, radiation therapy; Chemo, chemotherapy.

RESULTS

Table 2 summarizes immunoreactivities identified in the normal tissues studied. In normal and hyperplastic prostate glands, staining was either weak and luminal or absent. In several tissues, the immunohistochemical procedure routinely utilized was modified to include blocking of endogenous biotin to avoid false-positive reactions. Renal tubules, initially noted to display intense cytoplasmic staining, exhibited identical reaction patterns with class-matched primary antibody substitution and negative controls (Fig. 1a). Blocking of endogenous biotin abolished the background cytoplasmic staining and revealed immunoreactivity that was reproducibly restricted to a subset of proximal tubules (Fig. 1b). A similar situation was encountered in the gastrointestinal tract, with intense staining of the duodenal and colonic mucosa. Blocking of endogenous biotin revealed persistent immunoreactivity limited to the duodenal brush border (Fig. 1c). Rare cells in the deepest portions of the colonic crypts were immunoreactive (Fig. 1d); these had a morphology and distribution similar to those of chromogranin-positive cells in serial sections (data not shown), implying a possible neuroendocrine origin.

Table 3 summarizes immunoreactivities identified in the tumors studied. Significant PSMA expression was detectable in 33 of 35 primary prostate tumors. The pattern of staining varied with the degree of differentiation, with the most intense and homogeneous reactivity located at the luminal site of the glands in well-differentiated tumors (Fig. 2a). Immunoreactivity was more heterogeneous in less well-differentiated lesions (Fig. 2b). Considerable heterogeneity of expression within the same tumor was noted in most cases. No immunoreactivity was present in prostatic stromal elements, including blood vessels.

Similarly, seven of eight prostate carcinomas metastatic to lymph nodes expressed detectable PSMA levels (Fig. 2c). In the majority of cases, the staining pattern was reminiscent of that observed in poorly differentiated primary tumors, without any noticeable subcellular orientation. In one case, pseudogland formation was present with intense reactivity at the luminal site. Staining within a metastatic deposit was less heterogeneous than that in the primary tumors, with cells virtually all positive or all negative. lymphoid elements did not exhibit immunoreactivities. The 18 osseous metastases of prostate carcinoma were divided between cases with and without detectable PSMA ex-

The 18 osseous metastases of prostate carcinoma were divided between cases with and without detectable PSMA expression.
expression of PSMA by a subset of renal tubules cannot be controlled sections utilizing a class-matched primary antibody. PSMA expression in normal tissues, with several distinctions.

**DISCUSSION**

None of the 17 renal cell carcinomas, 13 bladder transitional cell carcinomas, and 19 colon adenocarcinomas evaluated showed detectable PSMA levels in the tumor cells. Stromal components were similarly negative, except for some blood vessels. Capillary endothelial cell immunoreactivity restricted to the region of the tumor was noted in 8 of 17 renal cell carcinomas (Fig. 3a), in 7 of 13 transitional cell carcinomas (Fig. 3b), and in 3 of 19 colon tumors (Fig. 3, c and d). Capillaries located in normal tissue adjacent to the tumors were not immunoreactive. Considerable heterogeneity of expression was evident, with virtually all peritumor capillaries positive in some cases and only a few capillaries positive in others. Blocking of endogenous biotin did not change this result, and it was not seen in control sections utilizing a class-matched primary antibody.

**Table 3** PSMA expression in tumor tissues

<table>
<thead>
<tr>
<th>Carcinoma</th>
<th>No. studied</th>
<th>Tumor cells</th>
<th>Neovascularity</th>
</tr>
</thead>
<tbody>
<tr>
<td>Prostate, primary</td>
<td>35</td>
<td>33</td>
<td>0</td>
</tr>
<tr>
<td>Prostate, metastatic</td>
<td>33</td>
<td>18</td>
<td>8</td>
</tr>
<tr>
<td>Lymph node</td>
<td>8</td>
<td>7</td>
<td>0</td>
</tr>
<tr>
<td>Bone</td>
<td>18</td>
<td>8</td>
<td>0</td>
</tr>
<tr>
<td>Renal cell</td>
<td>17</td>
<td>0</td>
<td>8</td>
</tr>
<tr>
<td>Bladder transitional cell</td>
<td>13</td>
<td>0</td>
<td>7</td>
</tr>
<tr>
<td>Colon</td>
<td>19</td>
<td>0</td>
<td>3</td>
</tr>
</tbody>
</table>

expression. The eight cases with immunostaining demonstrated considerable heterogeneity within each specimen (Fig. 2d). No staining of osseous or hematopoietic elements was observed.

None of the 17 renal cell carcinomas, 13 bladder transitional cell carcinomas, and 19 colon adenocarcinomas evaluated showed detectable PSMA levels in the tumor cells. Stromal components were similarly negative, except for some blood vessels. Capillary endothelial cell immunoreactivity restricted to the region of the tumor was noted in 8 of 17 renal cell carcinomas (Fig. 3a), in 7 of 13 transitional cell carcinomas (Fig. 3b), and in 3 of 19 colon tumors (Fig. 3, c and d). Capillaries located in normal tissue adjacent to the tumors were not immunoreactive. Considerable heterogeneity of expression was evident, with virtually all peritumor capillaries positive in some cases and only a few capillaries positive in others. Blocking of endogenous biotin did not change this result, and it was not seen in control sections utilizing a class-matched primary antibody.

**DISCUSSION**

The present study is supportive of previous evaluations of PSMA expression in normal tissues, with several distinctions. Expression of PSMA by a subset of renal tubules cannot be regarded as artifactual, since controls in biotin-blocked sections confirm this finding. The identification of rare PSMA-expressing cells in the colonic crypts represents a new finding. The morphology and immunohistochemical characteristics of these cells indicate a neuroendocrine origin. The significance of this finding is not clear, but it parallels the recent report by Carter et al. (3) of a carboxypeptidase involved in central nervous system glutamate metabolism with remarkable homology to PSMA. The finding of PSMA expression in the duodenum is consistent with the previous detection of PSMA mRNA transcripts in small bowel (13) and of PSMA in small bowel protein extracts (14). Additionally, monoclonal antibody 7E11-C5 has recently been shown to precipitate a molecule with folate hydrolase activity from prostate carcinoma cell line extracts (15), a finding which parallels the known high level of folate hydrolase activity in duodenal mucosa. Folate hydrolase is a carboxypeptidase and, like the brain enzyme, liberates glutamate as a reaction product. The possible function of PSMA as a folate hydrolase in the duodenum and in the prostate is currently under investigation. PSMA mRNA transcripts were also identified in central nervous system (13). However, immunohistochemically detectable PSMA expression was not seen in either cerebral cortex or cerebellum in the present study. This may represent expression of the alternatively spliced molecular form of PSMA (PSMA') lacking the epitope recognized by 7E11-C5 or expression at a specific brainstem or gangionic locus not analyzed.

The present study confirms results from previous analyses with respect to the immunohistochemical detection of PSMA expression in primary and metastatic prostate cancer. Horoszewicz et al. (7) described immunoreactivity in frozen prostate tissues, including nine of nine normal prostates, nine of nine primary prostatic carcinomas, and two of two lymph node metastases. Lopes et al. (8) compared staining patterns of 7E11-C5 and the radionuclide-labeled immunon conjugate CYT-356 in frozen prostate tissues. They noted immunohistochemical
Fig. 2 PSMA expression in prostatic carcinoma. Intense PSMA immunoreactivity in the glandular epithelium located mainly at the luminal site of a well-differentiated primary tumor (a). More homogeneous cytoplasm and membrane immunostaining of a poorly differentiated primary tumor (b). PSMA expression by tumor cells of lymph node metastasis. Note the absence of staining in lymphoid elements (c). Osseous metastasis showing a heterogeneous pattern of PSMA immunoreactivity (d). a–c, X200; d, X400.

Detection of PSMA by 7E11-C5 in an unspecified number of normal prostates and in 10 primary prostatic carcinomas. Wright et al. (9) found PSMA immunoreactivity in all normal prostates analyzed, 157 of 165 primary tumors, and 72 of 79 lymph node metastases.

With respect to prostate cancer metastatic to bone, Wright et al. (9) noted that all of the seven cases examined expressed PSMA. This is at variance with the current study, in which PSMA expression could be detected in only 8 of 18 osseous metastases (Table 2). This difference may be due to sample size or it may be related to the degree of differentiation and extensive prior treatment (androgen deprivation, radiation, chemotherapy) of the lesions analyzed. It is also possible that some bone metastases express the alternatively spliced form of PSMA (PSMA') lacking the epitope recognized by 7E11-C5. Additionally, although down-regulation of PSMA mRNA expression in response to androgen has been demonstrated in vitro, with the greatest expression noted at castrate levels of androgen (13), PSMA detection in the present study was lowest in the group of patients failing androgen deprivation. These patients represent those with tumor progression to osseous metastases despite hormonal manipulation. These findings support the hypothesis

Fig. 3 PSMA expression by neovascular capillary endothelial cells in peritumoral areas of selected primary epithelial malignancies. Renal cell carcinoma (a), transitional cell carcinoma of the urinary bladder (b), and colonic adenocarcinoma (c and d). Note the intense immunostaining of endothelial cells, whereas tumor cells had undetectable PSMA levels. a and d, X400; b and c, X200.
that the interaction of tumor with the metastatic site has an effect on tumor phenotype (16).

PSMA expression was not detected in a variety of primary epithelial tumors. The lack of PSMA in renal cell carcinomas is of interest, in view of its expression in a subset of proximal tubules. It is known that renal cell carcinomas, specifically clear cell and granular cell carcinomas, are derived from proximal epithelial cells. The undetectable PSMA levels in the renal cell carcinomas analyzed may be due to the loss of PSMA during malignant transformation. Alternatively, the lack of PSMA in the renal tumors studied may indicate that they are derived from cells not displaying the PSMA-positive phenotype. Similarly, the cells which express PSMA in colonic crypts are of neuroendocrine derivation. Since these cells are not the precursors of colonic adenocarcinomas, the lack of PSMA staining in tumor cells from these neoplasms is not unexpected.

An important finding of the present study is the novel demonstration of PSMA expression by neovascular capillary endothelium in the peritumoral areas of a variety of epithelial malignancies. The significance of this finding in terms of the function of PSMA is presently unclear; however, it may have therapeutic implications. Humanized anti-PSMA antibodies could be used to deliver a variety of agents aimed at destroying neovascularature, ranging from conventional cellular toxins to peptide-based produg activators. Additionally, analysis with RNase protection techniques has demonstrated the presence of PSMA mRNA in both healthy and diseased prostate tissue (13).

Further understanding of the PSMA gene’s control mechanisms may be useful in the development of promoter-driven gene therapy for both benign and malignant prostate diseases.

In summary, PSMA appears to be highly expressed in normal prostate tissue as well as primary and nodally metastatic prostate cancer. In the present study, 40% of prostate cancers metastatic to bone expressed PSMA. Examination of normal tissues revealed PSMA expression in prostate epithelium, duodenal mucosa, a subset of renal tubules, and certain neuroendocrine cells in colonic crypts. Carcinomas arising in the bladder, kidney, and colon do not appear to express PSMA. PSMA expression by peritumor capillaries must be examined in other malignancies to establish the range of this phenomenon.

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