Analysis of p53 expression and proliferative assessment using PCNA in localized prostate carcinoma

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Abstract

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Received July 3, 1998 Accepted November 17, 1998 The surgical specimens from 51 men submitted to radical prostatectomy for localized prostate cancer were examined by immunohistochemistry using proliferation cell nuclear antigen (PCNA) monoclonal antibody to evaluate the proliferative index (PI). The relationship between PI, biological variables and p53 protein expression was evaluated by immunohistochemistry. PI was low in invasive localized prostate carcinoma (mean, 12.4%) and the incidence of PCNA-positive cells was significantly higher in tumors with p53 expression (P = 0.0226). There was no statistical difference in PCNA values when biological parameters such as Gleason score, tumor volume, extraprostatic involvement, seminal vesicle infiltration or lymph node metastasis were considered. We conclude that proliferative activity is usually low in prostate carcinoma but is correlated with p53 immune staining, indicating that p53 is important in cell cycle control in this neoplasm.

Key words

- · Prostate carcinoma
- p53
- Proliferating cell nuclear antigen
- PCNA

Introduction

The assessment of proliferation is important to characterize the malignant phenotype and biological behavior of neoplasms. Several techniques such as tritium [³H]-thymidine radioautography, 5-bromo- and 5-iododeoxyuridine (Brdu) immunohistochemistry, mitotic figure counts, S-phase evaluation by flow cytometry or the observation of living cells in tissue culture by phase contrast microscopy are some of the tools to determine the proliferative activity. Proliferation cell nuclear antigen (PCNA) is an auxiliary enzyme protein for DNA polymerase-delta in-

volved in DNA synthesis during the S-phase of the cell cycle (1,2). The PCNA antibody can be used in formalin-fixed, paraffin-embedded tissue to evaluate cell proliferation. Cells in the late G1 and S phase show strong nuclear staining, which is not detected in mitotic or arrested cells. In prostate carcinoma the proliferative activity has been related to disease progression (3) and risk of death (4) and has been used to demonstrate highgrade prostatic intraepithelial neoplasia (HG-PIN) as a precursor of the invasive carcinoma (5). p53 is a tumor suppressor gene important in cell cycle control, DNA repair and apoptosis (6). Its role in prostate cancer has been

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evaluated by many investigators since Isaacs (7) described a suppression of tumorigenesis in prostate cancer cell lines by transfecting wild type p53. The most common alteration in the p53 gene, located in the 17p13.1 chromosome, is the missense mutation, that produces a protein with extended half-life that can be detected by immunohistochemistry using monoclonal antibodies (8). p53 immunostaining has been related to high histological grade, DNA aneuploidy and proliferation (9). Others consider the p53 mutation to be a late event related to androgen-independent growth (10). The aim of the present study was to evaluate proliferation, its relation to p53 expression and other biological variables. A retrospective study using paraffinembedded tissue from 51 men submitted to radical prostatectomy for localized prostate cancer was performed using anti-PCNA antibody (PC10) for the evaluation of proliferative activity. The PI, expressed as percent of positive cells in 500 cells sampled, was compared with p53 protein expression, Gleason score, tumor volume, extraprostatic involvement, seminal vesicle infiltration and lymph node metastasis.

Material and Methods

Fifty-one patients with prostate cancer clinically staged as T1 and T2N0M0 (11) were submitted to radical prostatectomy between March and June 1997 at the Sírio Libanês Hospital. The specimens were fixed in buffered formalin for 4 to 16 h. The entire surgical margin was stained with India ink. The left and right lobes were separated, 3mm transversal serial sections were obtained from each lobe and the whole tissue was submitted to histological examination. Sections of bladder neck, prostate apex, seminal vesicles and pelvic lymph nodes were also examined. An average of 10 to 16 slides from each gland were examined. The Gleason score was used for grading. Tumor volume was evaluated using a grid below the slides on which the tumor areas were previously sketched out. Tumor percentage was calculated by counting the number of squares on the grid involved by tumor and its relation to the number of squares of the entire area of the whole section on each slide. The mean percentage of tumor affecting the gland was considered and will be referred to as tumor volume in the text. The extraprostatic involvement was defined as tumor infiltrating the adipose tissue or neurovascular plexus. Only the parenchymatous infiltration of the seminal vesicle was considered positive. The tumors were divided into two groups for statistical comparison: favorable group which included tumors without extraprostatic involvement, seminal vesicle infiltration or lymph node metastasis, and the unfavorable group which included tumors with extraprostatic tissue involvement with seminal vesicle infiltration and/or lymph node metastasis.

Immunohistochemistry

Three-micrometer sections from paraffin-embedded specimens on adhesive-coated slides were submitted to heat antigen retrieval using a domestic microwave oven (high power) and citrate buffer, pH 6.0, 3 times, for 8 min (12). The endogenous peroxidase activity was quenched by exposing the sections to a solution of H_2O_2 for 5 min. The sections were incubated overnight with anti-PCNA (PC10) and anti-p53 (DO-7) monoclonal antibodies (Dako, Carpinteria, CA) at 1:100 dilution in PBS at 4°C. Biotinylated antimouse immunoglobulin G was applied at 1: 400 dilution for 60 min at room temperature (RT). After rinsing with PBS for 30 min, slides were incubated with peroxidase-conjugated streptavidin at 1:400 dilution for 45 min at RT and then rinsed with PBS for 30 min. Color was developed by incubating the slides with 0.06% diaminobenzidine in PBS for 15 min; slides were then rinsed in tap water, counterstained with Harris hematoxylin, dehydrated and coverslipped (13). One slide that best represented the tumor was selected and at least 500 cells in a "hot spot" area were counted with a high-power field light microscope. Only the dark nuclear staining was considered positive for PCNA and p53. The percentage of anti-PCNA-stained cells and 3 groups of p53 expression were used for statistical analysis. p53 expression was considered negative when less than 20% of cells were stained, low when p53 was between 20 to 50% and high when the percentage was higher than 50%.

Statistics

The patients were divided into favorable and unfavorable groups considering invasion of periprostatic tissue, seminal vesicle infiltration and lymph node metastasis. Mean PCNA values were compared between these groups by the chi-square test. Differences among the three groups in p53 immune expression in relation to PCNA were evaluated by analysis of variance.

Results

Patient age ranged from 50 to 74 years (mean, 65 years; median, 67 years). The Gleason score was 3-6 in 32 patients (62.7%) and 7-10 in 19 (37.3%). The mean tumor volume was 23.8% (median, 20%) ranging from 0.6 to 79%. Twenty-six (51%) patients had less than a 20% tumor volume and 25 (49%) more than a 20% tumor volume. Thirty-three (64.7%) patients were staged as T1-T2N0M0, 17 (33.3%) as T3N0M0 and 1 (2%) as T3N1M0. Extraprostatic tissue extension was observed in 13 patients (25.5%), seminal vesicle involvement in 11 (21.6%) and lymph node metastasis in only 1 case (2%). The percent of anti-PCNA-stained cells ranged from 0.5 to 46% (mean, 12.4%; median, 10%) (Figure 1). Eleven patients (21.6%) were in the unfavorable group and the PCNA value ranged from 3 to 29% (mean, 12.85%; median, 12%). The remaining cases (78.4%) were in the favorable group, with PCNA ranging from 0.5 to 46% (mean, 12.28%; median, 10%). There was no significant difference between groups (P = 0.8460) (Table 1). In 8 tumors p53 was considered negative, in 21 there was low immune reactivity and in 22 high immune reactivity (Figure 2). Mean p53 levels were 45.9%, ranging from 0 to 99% of positive cells. There was a statistically significant difference among the mean PCNA values of the three groups of p53 immune expression. Mean PCNA was 8.3% in the p53-negative

Table 1 - Proliferative cell nuclear antigen (PCNA) values of favorable and unfavorable groups considering extraprostatic involvement, seminal vesicle infiltration and lymph node metastasis.

*P = 0.8460. PCNA data are reported as percent of PCNA-stained cells.

Group		PCNA (%)						
	N	Mean (± SD)	Median	Range				
Favorable Unfavorable	40 11	12.28 ± 8.89* 12.85 ± 7.08*	10 12	0.5-46 3.0-29				

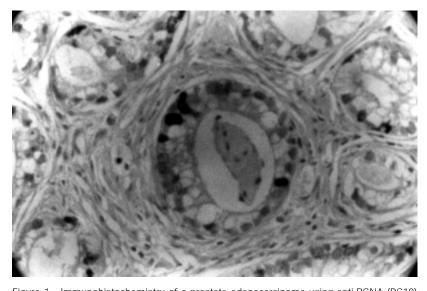


Figure 1 - Immunohistochemistry of a prostate adenocarcinoma using anti-PCNA (PC10) antibody. The dark nuclei are positive.

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group, 10.2% in the low expression group and 16% in the high expression group (P = 0.0226) (Table 2). No significant correlation was detectable between PCNA and any of the biological variables considered separately.

Discussion

The index of tumor cell proliferation can be used to define the behavior and prognosis in a series of human tumors, including breast cancer (14) and sarcomas (15). Proliferation activity in prostate can-

Table 2 - Proliferative cell nuclear antigen (PCNA) values and p53 immune expression.

PCNA data are reported as percent of PCNAstained cells. *P = 0.0226.

p53 expression	PCNA (%)					
	N	Mean (± SD)	Median	Range		
Negative	8	8.31 ± 4.89*	10.5	0.5-13		
Low	21	10.19 ± 6.20*	9.0	1.0-29		
High	22	16.02 ± 10.06*	15.5	4.0-46		

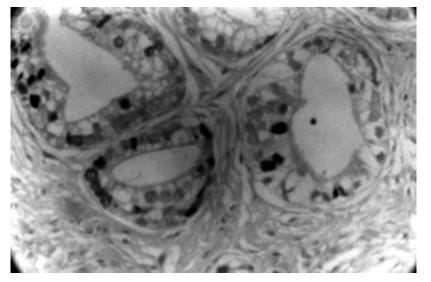


Figure 2 - Immunohistochemistry of a prostate adenocarcinoma using anti-p53 (DO7) antibody. Only the dark nuclei are considered to be positive.

cer has been assessed by immunohistochemistry using Ki-67 monoclonal antibody (16,17), PCNA (18) and tritium[3 H]-thymidine or bromodeoxyuridine labeling (19,20). Some studies demonstrated progressively rising levels of proliferative index (PI) values from PIN to invasive and metastatic carcinoma, illustrating the cancer progression (19,20). The mean PI values have been shown to range from 1.7 to 17.6% indicating low cell turnover consistent with a natural slow disease progression. Visakorpi et al. (9), studying a heterogeneous group of patients including advanced metastatic disease using the same antibody (PCNA-PC10), found mean PI values ranging from 0.3 to 29.1% and established a strong association between proliferation and p53 immunoreactivity. We could also demonstrate a strong association between PI value and p53 expression in our cases and since only localized prostate cancers were selected a lower PI value was determined, with a mean PCNA value of 12.4% and a median of 10%. Montironi et al. (18), also using PCNA antibody, found the same mean PI value in their cases and were able to correlate PI to histological grading. In our study we were unable to show a relationship between PI and Gleason grade, tumor volume or stage. Bubendorf et al. (21) obtained the same results using the MIB-1 antibody directed at recombinant parts of Ki-67 to evaluate proliferation only in localized prostate adenocarcinoma. They found a mean PI value of 7.7% and were also unable to correlate proliferation with tumor stage or histological grade. The tumor growth results from high cellular proliferation and a low rate of cell death by apoptosis. The p53 protein coded by the p53 gene, localized on the short arm of chromosome 17, is important for the cell-cycle control. This protein acts as a DNA guardian allowing expression of DNA repair genes as well as promoting apoptosis (6). In more than 50% of human tumors the p53 protein is inactivated. The

most common cause of p53 inactivation is missense mutation that promotes protein stabilization, allowing detection by immunohistochemistry (8). There are also alternative ways for inactivation: a) associations of p53 protein with other tumor suppressor genes such as WT1 and IRF1; b) inactivation by binding with oncoproteinlike adenovirus E1B and Mdm2 (murine double minute 2) protein (22), and c) destruction via a ubiquitin-dependent proteolytic pathway after binding to E6 protein, the product of human papillomavirus (HPV) 16 and 18 (23). We found p53 expression in 84% of our patients, with a mean value of 45.9%. Bauer et al. (24) found almost the same proportion of p53 expression (70.5%) using the same methodology, and the low or high expression was considered important to predict the relative risk of recurrence. The high level of nuclear p53 protein accumulation detected in our cases may be related to methodology since we used heat antigen retrieval with a microwave oven and citrate buffer, which considerably enhances antigen detection, unmasking antigenic epitopes (25,26). This method was not mentioned in other studies (9,10). We could not demonstrate a correlation between proliferation and biological parameters in localized prostate carcinoma, but we were able to find a strong relationship between PI and p53 protein expression (P = 0.0226), suggesting that P53 mutation gives a proliferative advantage to a subgroup of these neoplasms.

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