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## Implication of Cell Kinetic Changes during the Progression of Human Prostatic Cancer<sup>1</sup>

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### Abstract

The daily percentage of cells proliferating and dying were determined for normal, premalignant, and cancerous prostatic cells within the prostate as well as for prostatic cancer cells in lymph node, sort tissue, and bone metastases from untreated and hormonally failing patients. These data demonstrate that normal prostatic glandular cells have an extremely low but balanced rate of cell proliferation and death (*i.e.*, both  $<0.20\%/day$ ). This results in a steady-state, self-renewing condition in which there is no net growth, although the glandular cells are continuously being replaced (*i.e.*, turnover) every  $500 \pm 79$  days. Transformation of these cells into high-grade prostatic intraepithelial neoplastic cells initially involves an unbalanced increase in the daily percentage of cells proliferating *versus* dying, such that net continuous growth occurs (*i.e.*, mean doubling time,  $154 \pm 22$  days). As these early proliferation lesions continue to grow into late stage high-grade prostatic intraepithelial neo-plastic cells, the daily percentage of cells dying increases further to a point equaling the daily percentage of proliferation. This results in cessation of net growth while inducing a 6-fold increase in the turnover time of these cells (*i.e.*,  $56 \pm 12$  days), increasing their risk of further genetic changes. The transition of late stage high-grade prostatic intraepithelial neoplastic cells into localized prostatic cancer cells involves no further increase in proliferation but a decrease in death resulting in net continuous growth of localized prostatic cancers with a mean doubling time of 475 days. As compared to localized prostatic cancer cells, metastatic prostatic cancer cells within lymph nodes or bones of untreated patients have an increase in daily rate of proliferation coupled with a reduction in their daily percentage of cell death, producing net growth rates with a mean doubling time of  $33 \pm 4$  days and  $54 \pm 5$  days, respectively. Remarkably, there is no further increase in proliferation in hormonally failing patients, but instead an increase in the daily percentage of androgen-independent prostatic cancer cells dying within sort tissue or bone metastases. These changes result in doubling times which are two to three times longer (*i.e.*,  $126 \pm 21$  and  $94 \pm 15$  days) in these lymph node and bone metastatic sites, respectively, compared to similar sites in hormonally untreated patients. These data

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demonstrate that the daily percentage of proliferation for either androgen-dependent or -independent metastatic prostatic cancer cells is remarkably low (*i.e.*,  $<3.0\%/day$ ), consistent with why antiproliferative chemotherapy has been of such limited value against such metastatic cells. These results also suggest that prostatic carcinogenesis starts in the second to third decade of life and may require over 50 years for progression to pathologically detectable metastatic disease.

## Introduction

Metastatic prostatic cancers, like the normal prostates from which they arise, are sensitive to androgenic stimulation of their growth. This is due to the presence of androgen-dependent prostatic cancer cells within such metastatic patients. These cells are androgen dependent since androgen stimulates their daily percentage of proliferation (*i.e.*,  $K_p$ ) while inhibiting their daily percentage of death (*i.e.*,  $K_d$ ; Ref. 1 ). In the presence of adequate androgen, continuous net growth of these dependent cells occurs since the daily percentage of cells proliferating exceeds the daily percentage of cells dying. In contrast, following androgen ablation, androgen-dependent prostatic cancer cells stop proliferating and activate a cellular suicide pathway, termed programmed cell death or apoptosis (2). This activation results in the elimination of these androgen-dependent prostatic cancer cells since their daily percentage of cell death now exceeds their daily rate of cell of proliferation. Due to this elimination, 80–90% of all men with metastatic prostatic cancer treated with androgen ablation therapy have an initial positive response (3). All of these patients eventually relapse to a state unresponsive to further antiandrogen therapy, no matter how completely given (3). This is due to the heterogeneous presence of androgen-independent prostatic cancer cells within such metastatic patients (4). These latter cells are androgen-independent since their rate of proliferation exceeds their rate of cell death even after complete androgen blockage is performed (5).

Attempts to use nonandrogen ablative chemotherapeutic agents to adjust the kinetic parameters of these androgen-independent prostatic cancer cells so that their daily percentage of death exceeds their daily percentage of proliferation have been remarkable in their lack of success (6). The agents tested in androgen ablation-failing patients have been targeted at inducing DNA damage directly or indirectly via inhibition of DNA metabolism or repair (6). These agents are thus critically dependent upon an adequate rate of proliferation to be cytotoxic (7). *In vitro* cell culture studies have demonstrated that when androgen-independent, metastatic, prostatic cancer cells have a high daily percentage of cell proliferation (*i.e.*,  $K_p \approx 50\%$ ), these cells are highly sensitive to the induction of programmed cell death via exposure to the same antiproliferative chemotherapeutic agents which are of limited value when used *in vivo* in prostatic cancer patients (8). The apparent paradox between the *in vitro* and *in vivo* responsiveness to the same chemotherapeutic agents by androgen-independent prostatic cancer cells may reflect major differences in the daily percentage of proliferation occurring in the two states. Likewise, for chemotherapeutic agents to be effective, not only must the cancer cells have a critical daily percentage of proliferation but also a critical sensitivity to induction of cell death (9). This sensitivity to induction of cell death is reflected in the magnitude of the daily percentage of cell death in the untreated condition (9).

Thus, knowledge concerning the magnitude of the daily percentage of cell proliferation (*i.e.*,  $K_p$ ) and death (*i.e.*,  $K_d$ ) for androgen-independent metastatic prostatic cancer cells could provide a logical explanation of why the previously tested chemotherapeutic agents have been of limited success against these devastating cells. PSA<sup>4</sup> is a unique prostatic marker whose serum level is related to prostatic cancer burden (10, 11). Several studies have attempted to use serial measurements of PSA levels to estimate the growth rate of prostatic cancer cells in patients with both localized or metastatic disease. Using serial serum PSA values, doubling times for localized *versus* metastatic disease were reported to be  $2.4 \pm 0.6$  years *versus*  $1.8 \pm 0.2$  years in the Carteret *et al.* (10) study and 5.8 *versus* 3.6 years in the Schmid *et al.* (11) study. Since it requires at least 30 population doublings for a tumor to reach 1 cm<sup>3</sup> (12), these PSA-based growth rates predict that it should take minimally 72 years, based on the Carteret *et al.* (10) data, or 174 years, based on the Schmid *et al.* (11) data, for a patient to develop a 1 cm<sup>3</sup> primary prostatic cancer. Since the average time of diagnosis for clinical prostatic cancer is 72 years of age (13), these PSA-based calculations clearly are underestimates of the true growth rates for prostatic cancer cells. To obtain more accurate estimates of the growth rates of prostatic cancers and to clarify how these cell kinetic parameters are affected during prostatic cancer progression, the present study was performed. In this study, the *in vivo* daily percentage of cell proliferation and death were determined for normal, premalignant, and cancerous prostatic cells within the prostate as well as for prostatic cancer cells in lymph node, soft tissue, and bone metastases from untreated and hormonally failing patients.

## Materials and Methods

### Tissue

For determination of cell kinetic parameters of normal prostate glandular cells, prostatic intraepithelial neoplastic cells, and primary prostatic cancer cells, radical prostatectomy specimens were used. Prostates from 27 patients undergoing radical prostatectomy for pathological stage B localized prostatic cancer were processed for routine formalin fixation, paraffin embedding, and histological sectioning. Consecutive 4- $\mu$ m-thick step sections were obtained for each prostate with representative sections being hematoxylin and eosin stained to determine the total cancer volume (*i.e.*,  $2.9 \pm 0.3$  cm<sup>3</sup>; range, 0.3–7.5 cm<sup>3</sup>) and Gleason sum (*i.e.*, range, 4–10) and to define the area of prostatic cancer and the areas of peripheral prostatic tissue composed of only normal prostatic glandular cells. In this way, localized prostatic cancer cells could be age and patient matched to normal prostatic glandular cells for the kinetic analysis. An additional group of 20 prostates, also obtained from patients undergoing radical prostatectomy, was used to obtain histological sections containing high-grade PIN not associated with cancer (*i.e.*, early stage PIN) from 10 patients and sections containing high-grade PIN adjacent to primary prostatic cancer (*i.e.*, late stage PIN) from 10 patients. Classification of these lesions as high-grade PIN was as described previously (14). Pelvic lymph nodes containing metastatic prostatic cancer cells from a series of 30 patients with pathological stage D<sub>1</sub> metastatic prostatic cancer were also used. Also used were 23

<sup>4</sup>The abbreviations used are: PSA, prostatic specific antigen; PIN, prostatic intraepithelial neoplasia; GF, growth fraction; Tc, intermitotic cell cycle time; BBRF, Biological Research Faculty and Facilities; TTF, terminal transferase fraction; TT, terminal transferase.

bone metastases and 5 soft tissue metastases (*i.e.*, from epidural, bronchus, breast, laminectomy, and mesenteric/omental sites) obtained from a series of patients with pathological D<sub>2</sub> metastatic prostatic cancer whose treatment status and response data were known.

### Determination of the Daily Rate of Cell Proliferation ( $K_p$ )

The  $K_p$  value, expressed as the percentage of a particular cell type proliferating per day, was calculated by dividing the GF for the particular cell type by the intermitotic  $T_c$ , expressed in days, and then multiplying this number by 100 (15). The GF for a particular cell type was determined by immunocytochemical staining of histological sections using the commercially available mouse mAb MIB1 (AMAC, Westbrook, Maine) to detect cells in the cell cycle. This mAb has been demonstrated by Cattoretti *et al.* (16) to recognize epitopes in the Ki67 antigen in microwave-processed formalin-fixed, decalcified, paraffin sections. The Ki67 antigen is a nuclear nonhistone protein of 395 and 345 kDa present in all parts of the proliferative cell cycle (*i.e.*, G<sub>1</sub>, S, G<sub>2</sub>, and mitosis) but is absent when cells are out of cycle (*i.e.*, in G<sub>0</sub>; Ref. 16). For these immunocytochemical stains, the microwave protocol described by Cattoretti *et al.* (16) was used, except that the secondary antibody used was a biotinylated rabbit anti-mouse IgG obtained from Vector (Burlington, CA) and detection was via a Vectostain ABC peroxidase kit (Vector) using diaminobenzamide/nickel as the peroxidase substrate. After immunocytochemical staining, sections were counterstained with ethyl green. Two thousand cells per cell type per patient were chosen using the random sampling technique as we have described previously (17), and the fraction of cells whose nuclei were positively stained with the MIB1 antibody determined to calculate the GF.

The  $T_c$  was determined using primary cultures of both normal prostatic glandular cells and prostatic cancer cells. To do this normal prostatic glandular and cancerous tissues were obtained from radical prostatectomy specimens from patients with clinically localized prostatic cancers. These tissues were enzymatically dissociated into epithelial organoids by 8-12-h treatment with a 0.05% collagenase (GIBCO), 0.75% BSA (GIBCO), and antibiotics in RPMI 1640 media. The resultant epithelial organoids were plated in tissue culture dishes coated with a commercially available attachment solution containing bovine fibronectin, collagen, and BSA [*i.e.*, FNC coating mix (cat. AF-10) from BRFF, Ijamsville, MD]. The cells were maintained using a commercially available serum-free tissue culture media containing 10<sup>-8</sup> M dihydrotestosterone (*i.e.*, HPC-1) from BRFF supplemented with 25 ng/ml cholera toxin, 5 ng/ml epidermal growth factor, 5 µg/ml insulin, and 100 µg/ml bovine pituitary extract (additive commercially available from BRFF). In this media, prostatic epithelial cells, but not the fibroblasts, attach and spread over a 4-day period. These primary epithelial cultures were time-lapse video recorded over the next week using an inverted microscope as described previously (18). Using these video recordings, the time between successive mitosis (*i.e.*,  $T_c$ ) for both normal or malignant prostatic cells from five distinct patients were determined to be 48 ± 5 h (*i.e.*, 2 days). To determine whether the  $T_c$  changes when prostatic cancer progresses from a localized to a more advanced malignant phenotype, the DU-145, PC-3, and LNCaP human prostatic cell lines (*i.e.*, each established from a metastatic site) were video recorded as described previously (18). The mean  $T_c$  value was determined to be 48 ± 6 h for these three cell lines. These results demonstrated that there is

no change in the  $T_c$  value during the progression from localized to more advanced prostatic cancer. Thus, a value of 2 days was used as the  $T_c$  value for normal and malignant prostatic cells.

### Determination of the Daily Rate of Cell Death ( $K_d$ )

The  $K_d$  value, expressed as the percentage of a particular cell type dying per day, was calculated by dividing the fraction of the cells whose DNA is end labeled with exogenous *in situ* treatment with purified terminal transferase (*i.e.*, termed TTF) by the half-life of the labeled cells (*i.e.*, termed  $TT_{1/2}$ ) and then multiplying this number by 100. This calculation is based upon the fact that within the prostate, cell death normally occurs via programmed (*i.e.*, apoptotic) cell death and that during this programmed death, genomic DNA is double-strand fragmented, producing large numbers of free 3'-hydroxyl deoxynucleotide ends (19, 20). These newly produced 3'-hydroxyl groups can be end labeled *in situ* in formalin-fixed, paraffin-embedded histological sections as described by Gavrieli *et al.* (21) by incubation of the deparaffinated sections with 300 units/ml purified terminal deoxynucleotide transferase (*i.e.*, TT; Boehringer Mannheim) and 6.25  $\mu$ M biotinylated dUTP (Boehringer Mannheim) with subsequent detection of the end-labeled DNA performed with Vectostain ABC peroxidase kit using diaminobenzamidine/nickel as substrate. Sections were counterstained with ethyl green. Two thousand cells per cell type per patient were chosen using the random sampling technique described previously (17), and the fraction of cells whose nuclear DNA was positively end labeled but whose nuclei had not yet undergone apoptotic fragmentation was determined to calculate the TTF. Cells whose nuclear DNA was positively end labeled, but whose nuclei had already undergone apoptotic fragmentation, were excluded from the TTF determination. This exclusion is based on the observation that there can be substantial variation between the length of time such apoptotic cell fragments remain before becoming undetectable due to the phagocytosis and degradation by neighboring cells in normal and malignant tissue. In contrast to the variation in the time apoptotic cell fragments remain detectable, 12 h is consistently the average half-life (*i.e.*,  $TT_{1/2}$ ) for cell nuclei to be terminal transferase end labelable before undergoing apoptotic fragmentation. This mean  $TT_{1/2}$  value was determined by treating primary cultures of both normal and prostatic cancer cells from five separate patients and three established cell lines (*i.e.*, DU-145, PC-3, and LNCaP) with the 10  $\mu$ M of the calcium ionophore, ionomycin (Calbiochemical, CA), for 4 days *in vitro*. As described previously, such chronic treatment with ionomycin elevates the intracellular free  $Ca^{2+}$  ( $Ca_i$ ) and within 4 days induces programmed death of prostatic cells (18). After 4 days, triplicate cultures were fixed with 10% formalin, terminal transferase end labeled as described, and the TTF determined. Additional cultures had their ionomycin-containing media replaced with media lacking ionomycin, and then at 6-h intervals triplicate cultures were fixed and TTF determined. Previously we have demonstrated that once ionomycin is removed from the media, the  $Ca_i$  returns to baseline and no additional cells subsequently initiate their programmed death pathway (18). Thus, by plotting the log of TTF *versus* time of removal of ionomycin, the half-life of terminal transferase-labeled nuclei before they undergo apoptosis (*i.e.*,  $TT_{1/2}$ ), was determined to be  $12 \pm 2$  h (*i.e.*, 0.5 days) for normal prostatic cells,  $12 \pm 3$  h for the localized prostatic cancer cells, and  $12 \pm 3$  h for the cell lines. Thus, a value of 0.5 days was used as the  $TT_{1/2}$  for normal and malignant prostatic cells.

## Determination of Net Growth Rate, Doubling Time, and Turnover Time

The net growth rate, expressed as the percentage of cells accumulating per day, was determined by subtracting the  $K_d$  value from the  $K_p$  value for the particular cell type (15). When  $K_p > K_d$ , the doubling time was calculated according to the formula:  $\ln 2 / [K_p - K_d]$  (15). When  $K_p = K_d$ , the turnover time (TT) was determined using the formula:  $T_T = 1 / K_p$  (15).

## Statistical Analysis

Numerical values are expressed as the mean  $\pm$  SE. Statistical analyses of significance were made by a one-way ANOVA with the Kruskal-Wallis test.

## Results

### Kinetic Parameters for Normal and Premalignant Prostatic Glandular Cells

The percentage of normal glandular cells proliferating per day (*i.e.*,  $K_p$ ) is remarkably low in the prostate, Table 1 and Fig. 1A. However, this low rate of glandular cell proliferation is large enough to balance the equally low percentage of these cells spontaneously dying per day (*i.e.*,  $K_d$ , Table 1). This demonstrates that these normal cells are in a steady-state, self-renewing condition in which neither over-growth nor regression of these cells continuously occurs. During this maintenance condition, the turnover time (*i.e.*, the time required to renew these cells) is  $500 \pm 79$  days.

When such normal prostatic glandular cells undergo transformation into early stage high-grade PIN, there is a 6.9-fold increase ( $P < 0.05$ ) in the  $K_p$  value for these cells (Fig. 1C), with only a 4-fold increase ( $P < 0.05$ ) in the  $K_d$  value (Table 1 and Fig. 18). This leads to a predicted net growth rate (*i.e.*,  $K_p - K_d$ ) of  $0.45 \pm 0.11\%$  of cells accumulating per day. This translates into a doubling time of  $154 \pm 22$  days for these early high-grade PINs. For late stage high-grade PIN, there is a 9-10-fold increase ( $P < 0.05$ ) in both the  $K_p$  and  $K_d$  values as compared to the normal prostatic glandular cells. Since the  $K_p = K_d$ , however, late stage high-grade PINs do not continue to undergo net growth but instead are in a steady state of self-renewal. These late stage high-grade PINs, however, do have a 9-fold higher ( $P > 0.05$ ) turnover time (*i.e.*,  $56 \pm 12$  days for late stage high-grade PIN *versus*  $500 \pm 79$  days for normal prostatic glandular cells), thus increasing their risk of further genetic changes needed to acquire the ability for net continuous growth.

### Kinetic Parameters for Localized Prostatic Cancer Cells

There is an  $\approx 6$ -fold increase ( $P < 0.05$ ) in  $K_p$  values for localized prostatic cancer cells within the prostate of pathological stage B patients compared to normal prostatic glandular cells (Table 1 and Fig. 1E). However, there is no difference between the  $K_p$  values for high-grade PIN and localized prostatic cancer (Table 1). Likewise, there is no correlation between  $K_p$  values and patient age or primary tumor volume and the histological tumor grade (*i.e.*, Gleason sum; Note: there were no primary cancers with a Gleason sum below 4 in this series). As compared to the high  $K_d$  value in late stage high-grade PIN, there is a 40% decrease ( $P < 0.05$ ) in the  $K_d$  values for the localized prostatic cancer cells which is still, however, 6–7-fold higher than for the normal prostatic glandular cells (Table 1 and Fig. 1D).



There is no correlation between the  $K_d$  values and patient age or primary tumor volume histological tumor grade. These kinetic changes predict a net growth for these prostatic cancer cells. The mean net growth rate for low *versus* high Gleason sum localized prostatic cancers is  $0.12 \pm 0.03$  *versus*  $0.14 \pm 0.04\%$  of cells accumulating/day, respectively. This translates into a mean doubling time of  $577 \pm 68$  and  $495 \pm 56$  days, respectively, for low *versus* high Gleason sum localized cancers.

### Kinetic Parameters for Metastatic Prostatic Cancer Cells in Metastatic Sites

The mean  $K_p$  value of metastatic prostatic cancer cells in pelvic lymph nodes of hormonally untreated pathological stage D<sub>1</sub> patients was increased by 15-fold ( $P > 0.05$ ) compared to the normal prostate glandular cells and 1.5–2-fold ( $P > 0.05$ ) compared to either late stage prostatic intraepithelial neoplastic cells or localized prostatic cancer cells. There was no correlation between the  $K_p$  values and patient age or Gleason sum. (Note: there were no metastases with a Gleason sum below 6.) Likewise, there is no correlation between the  $K_d$  values and patient age or Gleason sum in the metastatic prostatic cancer cells growing in pelvic lymph nodes of untreated patients (Table 1). The  $K_d$  values for these metastatic cells in lymph nodes were reduced  $\approx 40\%$  ( $P < 0.05$ ) and  $\approx 60\%$  ( $P < 0.05$ ) compared to localized prostatic cancer cells and late stage high-grade prostatic intraepithelial neoplastic cells, respectively. These kinetic changes result in a predicted net growth rate of  $2.1 \pm 0.30\%$  of prostatic cancer cells accumulating in lymph nodes/day. This translates into a doubling time of  $33 \pm 4$  days.

The mean  $K_p$  value for metastatic prostatic cancer cells within the bone of untreated patients was 36% higher ( $P < 0.05$ ) than that of localized prostatic cancer cells and 10.7-fold increased ( $P > 0.05$ ) compared to normal prostatic glandular cells (Table 1). There is also an approximately 50% reduction ( $P < 0.05$ ) in the  $K_d$  value in these bone metastases as compared to localized prostatic cancer cells and a 60% reduction ( $P < 0.05$ ) compared to late stage high-grade prostatic intraepithelial neoplastic cells (Table 1 and Fig. 1F). The  $K_d$  value for bone, however, is still 3.8-fold ( $P > 0.05$ ) higher than for normal prostatic glandular cells. Thus, the predicted net growth rate for these bone metastatic prostate cancer cells is  $1.28 \pm 0.23\%$  of prostatic cancer cells accumulating in the bone/day, which translates into a doubling time of  $54 \pm 5$  days. This value is nearly 8–9-fold faster than that of localized prostatic cancer cells but approximately 40-fold lower ( $P < 0.05$ ) than those for the net growth rate of metastatic cells in lymph nodes.

In hormonally untreated patients, it is unknown what proportion of the metastatic prostatic cancer cells analyzed for the  $K_p$  and  $K_d$  determinations are androgen dependent *versus* independent. In contrast, in patients failing androgen ablation therapy, these metastatic cancer cells are androgen independent. Androgen-independent metastatic prostatic cancer cells in either bone or soft tissue sites in patients failing hormonal treatment have no significant difference in their  $K_p$  values compared to metastatic cells in the respective sites in hormonally untreated patients. There is, however, a 2-fold increase in the  $K_d$  values ( $P < 0.05$ ) for androgen-independent metastatic prostatic cancer cells in soft tissue and bone sites in failing *versus* hormonally untreated patients (Table 1). These kinetic changes result in predicted net growth rates of  $0.55 \pm 0.09\%$  and  $0.74 \pm 0.12\%$  of the androgen-independent

metastatic prostatic cancer cells accumulating/day respectively in soft tissue *versus* bone sites. This translates into a doubling time of  $126 \pm 21$  days and  $94 \pm 15$  days respectively for soft tissue *versus* bone sites in these hormonally failing patients.

## Discussion

The daily percentage of cells proliferating (*i.e.*,  $K_p$ ) can be calculated from the formula:  $K_p = \text{growth fraction/intermitotic cell cycle time (days)} \times 100$  (15). The GF can be determined using immunocytochemical staining with the M1B1 mAb to detect the fraction of a particular cell type positively expressing the Ki67 antigen in formalin-fixed, paraffin-embedded, histo-logical sections (16). The Ki67 antigen is a nuclear nonhistone protein expressed in all parts of the proliferative cell cycle, but absent where cells are out of cycle (16, 22). Baisch and Gerdes (23) demonstrated that identical GF values were obtained using Ki67 immunostaining or other methods, including stathmokinetic measurements using colchicine blockade or flow cytometry measurement with bromodeoxyuridine labeling, except when cells are nutritionally deprived. When cells are nutritionally deprived, the GF determined on the basis of Ki67 immunostaining is higher than the GF determined by S-phase progression (23). This suggests that under such nutritional deprivation, cells spend longer times in  $G_1$  than normal (*i.e.*,  $T_c$  increase). *In vivo*, the hallmark of such nutritional deprivation is the appearance of morphological signs of necrosis (*i.e.*, areas within tissues in which multiple adjacent cells have undergone cellular edema, vacuolization, and/or cellular lysis). Based on such standard morphological criteria, necrosis is not a commonly identified morphological features in any of the tissue samples examined in this study. Thus, within these samples, GF estimated on the basis of positive staining for the Ki67 antigen should be reasonably accurate and allow valid estimation of the  $K_p$  values, if appropriate  $T_c$  values are used in the calculation.

Presently, it is impossible to determine the  $T_c$  value within a particular cancer specimen based on immunocytochemical staining of a formalin-fixed, paraffin-embedded histological section without injecting the patient with nucleotide precursors and harvesting and analyzing repeated biopsies after varying times (9, 15). Thus, in order to allow retrospective analysis of archival pathology specimens from untreated patients a “standardized” value for  $T_c$  must be estimated and used in the  $K_p$  calculation. In the present studies, we used time-lapse videomicroscopy to determine the  $T_c$  directly for: (a) normal prostatic glandular cells, (b) primary cultures of pathological stage B (*i.e.*, localized) prostate cancer cells, and (c) continuously passageable cell lines established using prostatic cancer tissue obtained from metastatic sites from patients failing hormonal therapy. Thus, within this series of samples, the full range of progression (*i.e.*, normal prostate  $\rightarrow$  localized prostatic cancer  $\rightarrow$  metastatic androgen-independent prostatic cancer) is represented. Regardless of the progressional state, the mean  $T_c$  was determined to be 2 days within this series of samples. A similar mean  $T_c$  value of 2 days has been reported by Tubiana and Malaise (9) using the percentage of [ $^3\text{H}$ ]thymidine-labeled mitosis method on a series of more than 40 human solid cancers (9). Thus,  $T_c$  value of 2 days was used in all of our calculations. The accuracy of these calculated  $K_p$  values, therefore, is related to the validity of the estimated 2 days for the  $T_c$  value.



For determination of the daily percentage of prostatic cells dying (*i.e.*,  $K_d$ ), cell death was assumed to occur exclusively via apoptosis and not via necrosis in the various prostatic tissue samples analyzed. This assumption is based on histological evaluation using standard morphological criteria which demonstrated that necrosis is not a commonly identified morphological feature in any of the tissue samples examined. Since DNA fragmentation is a characteristic of apoptotic death, terminal transferase end labeling can be used to identify cells undergoing apoptosis in histological sections (21). Thus, the daily percentage of cells dying was determined using the formula:  $K_d = \text{fraction of cells terminal transferase labelable but not yet undergoing fragmentation into apoptotic bodies (TTF)} \div \text{the half-life (in days) of terminal transferase end-labeled nuclei before undergoing fragmentation into apoptotic bodies (TT}_{1/2}) \times 100$ . Presently, it is impossible to determine the  $TT_{1/2}$  value for each individual cell type based on immunocytochemical staining alone. Therefore, the average half-life for terminal transferase end-labeled cell nuclei before undergoing fragmentation into apoptotic bodies ( $TT_{1/2}$ ) was determined for: (a) normal prostatic glandular cells, (b) primary culture of pathological stage B (*i.e.*, localized) prostatic cancer cells, and (c) continuous cell lines established using prostatic cancer tissue obtained from metastatic sites from patients failing hormonal therapy. Thus, within this series of samples, the full range of the progressional state is presented. Regardless of the progressional state, the mean half-life for terminal transferase-labeled nuclei before undergoing fragmentation into apoptotic bodies ( $TT_{1/2}$ ) was 0.5 days. Thus, in the present study, 0.5 days was used as the standardized  $TT_{1/2}$  value for all subsequent calculations of  $K_d$ . The accuracy of these calculated  $K_d$  values is related to the validity of the estimated 0.5 days for the  $TT_{1/2}$  value.

Using the described formulas, the  $K_p$  and  $K_d$  values were estimated for normal prostatic glandular cells, which are the cells of origin for the majority of prostatic adenocarcinomas (24). These calculations demonstrate that there is an extremely low, but balanced, daily percentage of proliferation and death of glandular cells within the prostate (*i.e.*,  $\approx 0.2\%/day$ ). These results are consistent with previous reports that the rates of proliferation of normal prostatic glandular cells are very low (24–27). Under such maintenance conditions, the turnover time for these normal glandular cells is  $500 \pm 79$  days. This estimate is consistent with a previous estimate that prostatic glandular cells have an average life span of longer than 2 years using a more indirect method of calculation (28).

High-grade PIN is believed to be the major premalignant lesion for prostatic cancer (29). Early high-grade PINs are characterized by a hyperplastic increase in their  $K_p$  value with a smaller increase in their  $K_d$  values. Using these cell kinetic parameters (*i.e.*,  $K_p - K_d$ ), it is estimated that these early high-grade PINs are growing with a mean doubling time of  $154 \pm 22$  days. Since one cell must undergo 26 population doublings to reach a reasonable size for detection (*i.e.*,  $60 \text{ mm}^3$  or  $5 \text{ mm} \times 5 \text{ mm} \times 5 \text{ mm}$ ; Ref. 12),  $\approx 11$  years are required for the clonal outgrowth of a high-grade prostatic intraepithelial neo-plastic cell to reach such a detectable size. It is likely, however, that this is a substantial underestimate of the actual time required for two reasons. First, the average volume of PIN in a series of 54 consecutive patients undergoing radical prostatectomy was determined to be  $\approx 2000 \text{ mm}^3$  (30). Second, as high-grade PINs grow (*i.e.*, becoming late stage high-grade PINs), their  $K_d$  values increase to a point equaling their  $K_p$  values. This increase in the  $K_d$  value could be

explainable due to the observation that the growth of these cells is centripetal into the luminal (*i.e.*, intraepithelial) space (29). Such centripetal growth is away from the capillary blood supply in the stroma. Recent studies have demonstrated that tumor angiogenesis is a late event in prostatic carcinogenesis occurring after a hyperplastic response (31). Thus, an increasing degree of cellular hypoxia as the high-grade prostatic intraepithelial neoplastic cells continues to expand centripetally could be the mechanism for the increase in  $K_d$  in these late stage high-grade prostatic intraepithelial neoplastic lesions.

The transition of these premalignant lesions into localized prostatic cancer cells within the prostate involves no further hyperplastic increase in the  $K_p$  value. Instead, the major change during this transition is a decrease in the  $K_d$  value. This may be due to the change from centripetal to centrifugal growth of these cancer cells, which is associated with loss of the basal epithelial layer and breakdown of the basement membrane (32). Such changes allow the centrifugally expanding cancer cells to invade the stroma, becoming closer to the capillary blood supply and thus less hypoxic. Because of the decrease in  $K_d$ , localized prostatic cancer cells within the prostate grow with an estimated mean doubling time of  $479 \pm 56$  days. This remarkably slow rate of growth of prostatic cancer cells within the prostate has been previously demonstrated by a series of earlier studies (24-27, 33-36). At this slow growth rate,  $\approx 39.4$  years would be required for the clonal outgrowth of such a localized prostatic cancer cell to reach  $1 \text{ cm}^3$  [*i.e.*, it takes 30 population doublings to reach  $1 \text{ cm}^3$  (12), thus  $30 \times 479 \text{ days} = 39.4 \text{ years}$ ]. Since the mean age of prostatic cancer diagnosis is 72 years (13), this suggests that prostatic carcinogenesis starts in the third to fourth decade of life. This estimate is consistent with histological data demonstrating that the frequency of high-grade PIN is highly age related, increasing rapidly between the third and fourth decade of life (37).

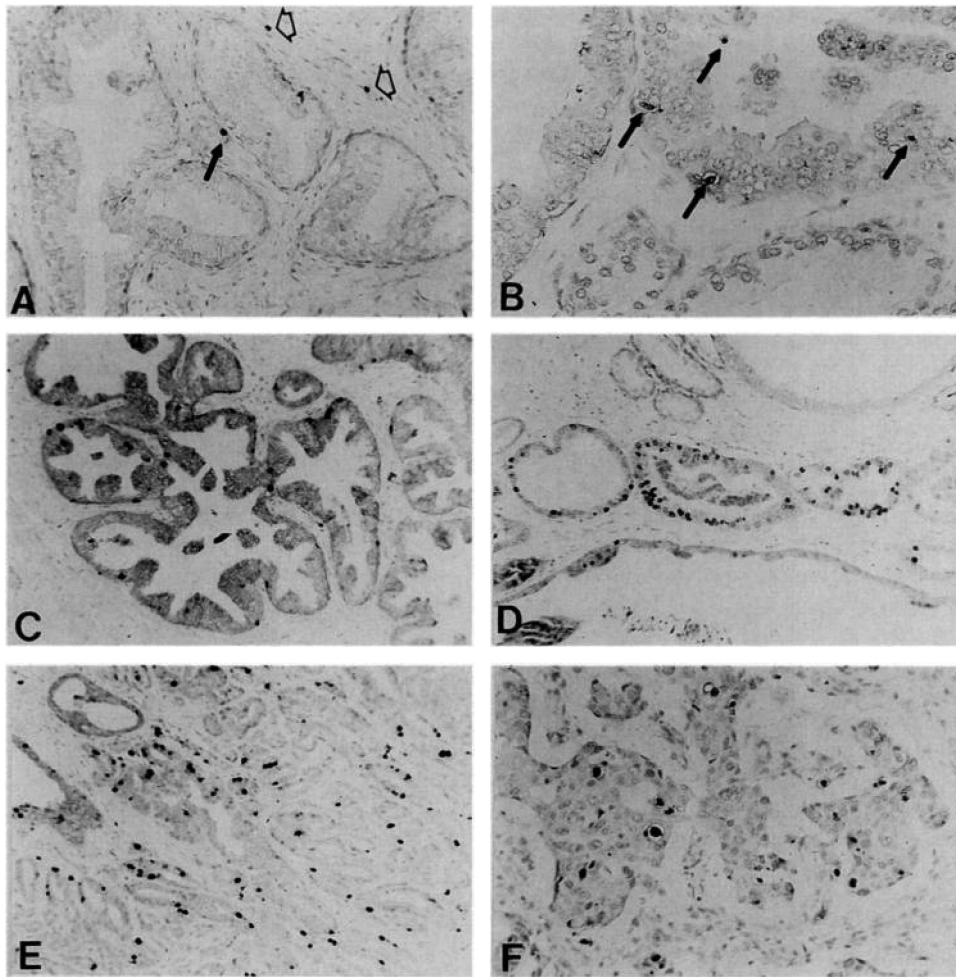
Due to the multistep nature and the remarkably long latent time period required to produce clinically detectable disease, prostatic carcinogenesis may be one of the most sensitive and appropriate disease processes for chemoprevention attempts. Chemoprevention is particularly appealing since the majority of available cytotoxic chemotherapies require a critical daily percentage of cell proliferation to be effective (7). Thus, the observation that the  $K_p$  values are very low (*i.e.*,  $<3\%/ \text{day}$ ) for metastatic, as well as localized, prostatic cancer cells is entirely consistent with the clinical observations that treatment of the metastatic patients with standard antiproliferation chemotherapeutic agents have been of only limited success. These data emphasize the need for the use of chemotherapeutic agents which can activate programmed death of androgen-independent metastatic prostatic cancer cells efficiently in a proliferation-independent manner. Indeed, such agents have already been identified (8, 38).

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**Fig. 1.** Immunocytochemical detection of prostatic cells in the proliferative cell cycle (A, C, and E) and undergoing cell death (B, D, and F). M1B1 antibody staining of proliferating cells in: A, normal prostatic glandular tissue; C, high-grade early PIN, primary prostatic cancer. Terminal transferase *in situ* end labeling of cells dying in: B, high-grade early PIN; D, primary prostatic cancer; F, bone metastases from hormonally untreated patients. All areas are representative of random fields except D, which represents a selected area of unusually high cell death, and E, which represents a selected area of unusually high cell proliferation. *Open arrows*, tissue in the stroma; *closed arrows*, tissue in the glandular epithelium.

**Table 1**

Kinetic parameters of normal and neoplastic prostatic epithelial cells

Cell type	Histological grade (Gleason sum)	% Cells		Doubling time (days)
		Proliferating/day	Dying/day (*d)	
Normal prostatic glandular epithelial cells (N = 27)		0.19 ± 0.03 <sup>a</sup> (0.05-0.30)*	0.20 ± 0.03 <sup>a</sup> (0.05-0.35)	
High-grade prostatic intraepithelial neoplastic cells				
Early (N = 10)		1.25 ± 0.30 <sup>c</sup> (0.25-2.60)	0.80 ± 0.24 <sup>a,c</sup> (0.20-0.80)	154 ± 22
Late (N = 10)		1.80 ± 0.28 <sup>c</sup> (0.20-2.50)	1.80 ± 0.16 <sup>a,c</sup> (0.20-4.96)	
Localized prostatic cancer cells within the prostate				
Low Gleason sum (>6) (N = 15)	5.3 ± 0.3 (4-6)	1.54 ± 0.17 <sup>c</sup> (0.40-2.60)	1.42 ± 0.20 <sup>c</sup> (0.25-5.25)	577 ± 68
High Gleason sum (>6) (N = 12)	7.9 ± 0.4 (7-10)	1.46 ± 0.20 <sup>c</sup> (0.80-2.85)	1.32 ± 0.18 <sup>c</sup> (0.10-3.10)	495 ± 56
Metastatic prostatic cancer cells from hormonally untreated patients				
Within Lymph node (N = 30)	8.1 ± 0.5 (6-10)	2.90 ± 0.30 <sup>a,c</sup> {0.45-3.10}	0.85 ± 0.21 <sup>c</sup> (0.15-1.30)	33 ± 4
Within the Bone (N = 13)	8.2 ± 0.20 (6-10)	2.04 ± 0.29 <sup>a,c</sup> (0.1-2.85)	0.76 ± 0.16 <sup>a,c</sup> (0.10-3.60)	54 ± 5
From hormonally failing patients				
Within distinct nonbone sites (N = 5)	10 (10)	2.77 ± 0.31 <sup>a,c</sup> (0.65-3.90)	2.22 ± 0.40 <sup>a,c</sup> (1.50-3.65)	126 ± 21
Within the bone (N = 10)	9.2 ± 0.3 (7-10)	2.42 ± 0.40 <sup>a,c</sup> (0.10-4.60)	1.68 ± 0.38 <sup>c</sup> (2.10-7.75)	94 ± 15

<sup>a</sup>  $P < 0.05$  compared to primary prostatic cancer cells.<sup>b</sup> Values in parentheses, range for particular cell type.<sup>c</sup>  $P < 0.05$  compared to normal prostatic glandular epithelial cells.