Cannabinoid Receptor as a Novel Target for the Treatment of Prostate Cancer

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Abstract

Cannabinoids, the active components of Cannabis sativa Linnaeus (marijuana) and their derivatives have received renewed interest in recent years due to their diverse pharmacologic activities such as cell growth inhibition, anti-inflammatory effects and tumor regression. Here we show that expression levels of both cannabinoid receptors, CB1 and CB2, are significantly higher in CA-human papillomavirus-10 (vireally transformed cells derived from adenocarcinoma of human prostate tissue), and other human prostate cells LNCaP, DU145, PC3, and CWR22Rv1 than in human prostate epithelial and PZ-HPV-7 (vireally transformed cells derived from normal human prostate tissue) cells. WIN-55,212-2 (mixed CB1/CB2 agonist) treatment with androgen-responsive LNCaP cells resulted in a dose- (1-10 μmol/L) and time-dependent (24-48 hours) inhibition of cell growth, blocking of CB1 and CB2 receptors by their antagonists SR141716 (CB1) and SR144528 (CB2) significantly prevented this effect. Extending this observation, we found that WIN-55,212-2 treatment with LNCaP resulted in a dose- (1-10 μmol/L) and time-dependent (24-72 hours) induction of apoptosis (a), decrease in protein and mRNA expression of androgen receptor (b), decrease in intracellular protein and mRNA expression of prostate-specific antigen (c), decrease in secreted prostate-specific antigen levels (d), and decrease in protein expression of proliferation cell nuclear antigen and vascular endothelial growth factor (e). Our results suggest that WIN-55,212-2 or other non–habit-forming cannabinoid receptor agonists could be developed as novel therapeutic agents for the treatment of prostate cancer. (Cancer Res 2005; 65(5): 1635-41)

Introduction

Because prostate cancer has become the most common cancer diagnosed in men, developing novel targets and mechanism-based agents for its treatment has become a challenging issue. In recent years cannabinoids, the active components of Cannabis sativa Linnaeus (marijuana) and their derivatives have drawn renewed attention because of their diverse pharmacologic activities such as cell growth inhibition, anti-inflammatory effects, and tumor regression (1-5). Cannabinoids have been shown to induce apoptosis in gliomas (6), PC-12 pheochromocytoma (7), CHP 100 neuroblastoma (8), and hippocampal neurons (9) in vitro, and most interestingly, regression of C6-cell gliomas in vivo (10). Further interest in cannabinoid research came from the discovery of specific cannabinoid systems and the cloning of specific cannabinoid receptors (10). These diversified effects of cannabinoids are now known to be mediated by the activation of specific G protein-coupled receptors that are normally bound by a family of endogenous ligands, the endocannabinoids (11, 12). Two different cannabinoid receptors have been characterized and cloned from mammalian tissues: the “central” CB1 receptor (13), and the “peripheral” CB2 receptor (14).

In the present study, we show for the first time that expression levels of both cannabinoid receptors, CB1 and CB2, are higher in human prostate cancer cells than in normal cells. Importantly, we also show that WIN-55,212-2 (CB1/CB2 agonist) treatment with androgen-responsive LNCaP cells results in a dose- and time-dependent inhibition of cell growth with a concomitant induction of apoptosis, decrease in protein and mRNA expression of androgen receptor and prostate-specific antigen (PSA), decrease in secreted PSA levels, protein expression of proliferating cell nuclear antigen (PCNA), and vascular endothelial growth factor (VEGF). We suggest that cannabinoid receptor agonists may be useful in the treatment of human prostate cancer.

Materials and Methods

Materials. R-(+)-WIN-55,212-2 (2,3 dihydro-5-methyl-3 [(morpholinyl)methyl]pyrrollo (1,2,3 de)-1,4-benzoxazinyl]-[1-naphthaleny]methanone, C27H26N2O3CH3SOH was purchased from Sigma (St. Louis, MO), CB1 receptor antagonist SR141716 (SR1) and CB2 receptor antagonist SR144528 (SR2) were procured from Dr. Herbert H. Seltzman (National Institute on Drug Abuse, Division of Neuroscience and Behavioral Research, through RTI International, Research Triangle Park, NC). DMEM and fetal bovine serum were procured from Life Technologies, Invitrogen Corporation (Grand Island, NY). Human PSA ELISA kit from yes Biotech Laboratories (Ontario, Canada) and annexin-V-FLUOS staining kit were from Roche Diagnostic Corporation (Indianapolis, IN). Antibiotics (penicillin and streptomycin) were used obtained from Cellgro Mediatech, Inc. (Herndon, VA). Apo-Direct kit for measuring apoptosis by flow cytometry was procured from ApoDirect (San Diego, CA). RNA isolation kit was from Qiagen, Inc. (Valencia, CA). Monoclonal antibodies for PSA, androgen receptor, PCNA, and VEGF, were purchased from Santa Cruz Biotechnology, Inc. (Santa Cruz, CA). Anti-mouse secondary horseradish peroxidase conjugate was obtained from Amersham Biosciences Limited (Buckinghamshire, United Kingdom). Protein was estimated using bicinchoninic acid protein assay kit obtained from Pierce (Rockford, IL).

Cell Culture. The LNCaP, DU145, PC-3, CWR22Rv1, CA-HPV-10, and PZ-HPV-7 cells were obtained from American Type Culture Collection (Manassas, VA). LNCaP and DU145 cells were cultured in DMEM supplemented with 5% heat-inactivated fetal bovine serum and 1% antibiotic penicillin and streptomycin. PC-3 and CWR22Rv1 cells were cultured in RPMI 1640 medium supplemented with 5% heat-inactivated fetal bovine serum and 1% antibiotic penicillin and streptomycin. CA-HPV-10 and PZ-HPV-7 were grown in keratinocyte serum-free medium (17005-042, Life Technologies) supplemented with 5 ng/mL human recombinant EGF and 50 μg/mL bovine pituitary extract. Human prostate epithelial cells (PrEC) were obtained from Cambrex Bioscience (Walkersville, MD) and grown in prostate epithelial basal cell medium (Cambrex Bioscience)
according to the manufacturer's instructions. The cells were maintained under standard cell culture conditions at 37°C and 5% CO₂ in a humid environment.

**Treatment of Cells.** WIN-55,212-2 (dissolved in DMSO), was used for the treatment of cells. The final concentration of DMSO used was 0.1% (v/v) for each treatment. For dose-dependent studies, cells were treated with WIN-55,212-2 at final concentrations of 1.0, 2.5, 5.0, 7.5, and 10.0 μmol/L for 24 hours in complete cell medium, whereas for time-dependent studies, the cells (50-60% confluent) were treated with 5 μmol/L dose of WIN-55,212-2 for 24, 48, and 72 hours. For time-dependent study, we included a control treated with DMSO for 72 hours because it was the longest time point post-WIN-55,212-2 treatment in our experimental protocol. To establish the role of CB1 and CB2 receptors in the effect of WIN-55,212-2, cells were treated with 5 μmol/L dose of WIN-55,212-2 for 24, 48, and 72 hours. For time-dependent studies, the cells were treated with 2 μmol/L of SR141716 or SR144528 alone for 24 hours. In the second experiment, cells were pretreated with each of these antagonists (2 μmol/L) for 3 hours following incubation with 7.5 μmol/L WIN-55,212-2 for 24 hours. In pilot experiments, it was established that DMSO (0.1% v/v) had no effects when measured at 24, 48, or 72 hours.

**Cell Viability.** The effect of WIN-55,212-2 on the viability of cells was determined by [3-(4,5-dimethylthiazol-2-yl)-2,5-diphenyl tetrazolium bromide] (MTT) assay. The cells were plated at 1 × 10⁵ cells per well in 200 μL of complete cell culture medium containing 1.0, 2.5, 5.0, 7.5, and 10.0 μmol/L concentrations of WIN-55,212-2 in 96-well microtiter plates for 24 and 48 hours at 37°C in a humidified chamber. Each concentration of WIN-55,212-2 was repeated in 10 wells. After incubation for specified times at 37°C in a humidified incubator, MTT reagent (4 μL, 5 mg/mL in PBS) was added to each well and incubated for 2 hours. The microtiter plate containing the cells was centrifuged at 1,800 rpm for 5 minutes at 4°C. The MTT solution was removed from the wells by aspiration and the formazan crystals were dissolved in DMSO (150 μL). Absorbance was recorded on a microplate reader at 540 nm wavelength. The effect of WIN-55,212-2 on growth inhibition was assessed as the percentage of inhibition in cell growth where vehicle-treated cells were taken as 100% viable.

**Preparation of Cell Lysates and Western Blot Analysis.** Following treatment of cells with WIN-55,212-2, the medium was aspirated and the cells were washed with cold PBS [10 mmol/L (pH 7.45)]. The cells were then incubated in ice-cold lysis buffer (50 mmol/L Tris-HCl, 150 mmol/L NaCl, 1 mmol/L EDTA, 1 mmol/L NaF, 100 mmol/L Na₃VO₄, 0.5 mmol/L PMSF, 1% Triton X-100, 1 mmol/L phenylmethylsulfonyl fluoride, pH 7.4), with freshly added protease inhibitor cocktail (protease inhibitor cocktail set III, Calbiochem, La Jolla, CA) over ice for 20 minutes. The cells were scraped and the lysate was collected in a microfuge tube and passed through a 21.5-gauge needle to break up the cell aggregates. The lysate was cleared by centrifugation at 14,000 × g for 15 minutes at 4°C, and the supernatant (total cell lysate) was collected, aliquoted, and used on the day of preparation or immediately stored at −80°C for use at a later time. For Western blotting, 25 to 50 μg of protein were resolved over 12% polyacrylamide gels and transferred onto a nitrocellulose membrane. The nonspecific sites on blots were blocked by incubating in blocking buffer [5% nonfat dry milk/1% Tween 20 in 20 mmol/L TBS (pH 7.6)] for 1 hour at room temperature, incubated with appropriate monoclonal primary antibody in blocking buffer for 90 minutes to overnight at 4°C, followed by incubation with anti-mouse or anti-rabbit secondary antibody horseradish peroxidase conjugate and detected by chemiluminescence and autoradiography using Hyperfilm obtained from Amersham Biosciences. Densitometric measurements of the bands in Western blot analysis were done using digitalized scientific software program UN-SCAN-IT purchased from Silk Scientific Corporation (Orem, UT).

**ELISA for PSA.** The human PSA ELISA kit was used for the quantitative determination of PSA levels in culture medium according to the vendor's protocol. This kit uses a technique of quantitative sandwich immunoassay for determination of PSA with an estimated sensitivity of 1 ng/mL PSA antigen.

**Detection of Apoptosis and Necrosis by Confocal Microscopy.** The annexin-V-FLUOS staining kit was used for the detection of apoptotic and necrotic cells according to vendor's protocol. This kit uses a dual-staining protocol in which the apoptotic cells are stained with annexin-V (green fluorescence), and the necrotic cells are stained with propidium iodide (PI; red fluorescence). LNCaP cells were grown to about 70% confluence and then treated with WIN-55,212-2 (1.0, 2.5, 5.0, 7.5, and 10.0 μmol/L) for 24 hours. The fluorescence was measured by a Zeiss 410 confocal microscope (Thornwood, NY). Confocal images of green annexin-FTC fluorescence were collected using 488 nm excitation light from an argon/krypton laser, a 560 nm dichroic mirror, and a 514 to 540 nm bandpass barrier filter. Images of red PI fluorescence were collected using a 568 nm excitation light from the argon/krypton laser, a 560 nm dichroic mirror, and a 590-nm-long pass filter. In a selected field, the cells stained with annexin-V and PI as well as unstained cells were counted to ascertain the extent of apoptosis and necrosis.

**Quantification of Apoptosis by Flow Cytometry.** The cells were grown at density of 1 × 10⁶ cells in 100 mm culture dishes and were treated with WIN-55,212-2 (1.0, 2.5, 5.0, 7.5, and 10.0 μmol/L doses) for 24 hours. The cells were trypsinized, washed with PBS, and processed for labeling with fluorescein-tagged dUTP nucleotide and PI by use of an Apo-Direct apoptosis kit obtained from Phoenix Flow Systems (San Diego, CA) and was used according to the manufacturer's protocol. The labeled cells were analyzed by flow cytometry.

**Quantitative Real-Time PCR for mRNA Expression of Androgen Receptor and PSA.** Total RNA was isolated from LNCaP cells using RNeasy kit according to the vendor's protocol. The ratio of optical densities of RNA samples at 260 and 280 nm was consistently >1.8, cDNA was synthesized by reverse transcription of 1 μg of extracted RNA with 200 units of Moloney murine leukemia virus reverse transcriptase (Promega, Madison, WI) in the presence of oligo dT and deoxynucleotide triphosphate (Promega). Androgen receptor and PSA were amplified using a FailSafe real-time PCR system obtained from Epicentre (Madison, WI). The thermal cycle used for amplification was an ABI-PRISM 7000 sequence detection system (Applied Biosystems, Foster City, CA). Primers were designed as follows: androgen receptor forward, 5′-AAGACGCTTCTACAGGCTCAACCA; reverse, 5′-TCCACAGAAAGGATCTTGGGCCACTT; PSA forward, 5′-ACTCCAGCAGAGATTGGGACTGAA; reverse, 5′-TGAGGGTGTCTCGGAGGACTTCA. The cycler was programmed with the following conditions (a) initial denaturation at 94°C for 2 minutes, followed by 35 cycles of (b) 94°C for 40 seconds, (c) annealing of the primer template at 58°C for 40 seconds, and (d) extension at 72°C for 40 seconds.

**Results**

**Expression of Cannabinoid Receptor in Normal and Prostate Cancer Cells.** We first compared the expression levels of both cannabinoid receptors CB₁ and CB₂ in PrEC and a series of human prostate cancer cells. We also included a pair of cells, PZ-HPV-7 (virally transformed cells, derived from normal human prostate tissue) and CA-HPV-10 (virally transformed cells, derived from the adenocarcinoma of human prostate tissue) derived from the same individual. Immunoblot data shown in Fig. 1 revealed that expression of both CB₁ and CB₂ receptors was significantly higher in prostate cancer cells LNCaP, DU145, PC3, CWR22Rv1, and CA-HPV-10 as compared with normal prostate cells PZ-HPV-7 and PrEC cells. To establish the specificity of the cannabinoid receptor antibodies used in the blotting experiments, antigen preabsorption experiments were carried out. The peptides blocked anti-CB₁ and anti-CB₂ antibody binding in all cells (data not shown).

**Effect of WIN-55,212-2 on Cell Viability of PrEC and LNCaP Cells.** To evaluate the cell viability response of WIN-55,212-2 on PrEC and LNCaP cells, we used a MTT assay. The fluorescence was measured by a Zeiss 410 confocal microscope (Thornwood, NY). Confocal images of green annexin-FTC fluorescence were collected using 488 nm excitation light from an argon/krypton laser, a 560 nm dichroic mirror, and a 590-nm-long pass filter. In a selected field, the cells stained with annexin-V and PI as well as unstained cells were counted to ascertain the extent of apoptosis and necrosis.
treatment of LNCaP cells with similar doses of WIN-55,212-2 in a dose-dependent manner significantly decreased the viability of LNCaP cells at 24 and 48 hours (Fig. 2A). The IC_{50} for inhibition of cell viability at 24 and 48 hours was 6.0 and 5.0 \mu M, respectively. CB_{1} and CB_{2} receptor activation signals growth inhibition in LNCaP cells. To study the possible implication of CB_{1} and CB_{2} receptors in WIN-55,212-2-induced cell death, the effect of their antagonists were evaluated using MTT assay. Cells pretreated with 2 \mu M of SR141716 (CB_{1} antagonist) or SR144528 (CB_{2} antagonist) had no effect on cell viability but exhibited significant protective effect when coadministered with WIN-55,212-2 (7.5 \mu M) at a molar ratio of 1:3.75 (Fig. 2B). These data suggest that both CB_{1} and CB_{2} receptors may be involved in WIN-55,212-2-mediated growth inhibition and apoptosis.

Effect of WIN-55,212-2 on Apoptosis and Necrosis of LNCaP Cells. We next assessed whether the cell growth inhibitory effect of WIN-55,212-2 was associated with induction of apoptosis. The induction of apoptosis by WIN-55,212-2 was evident from the analysis of the data obtained by confocal microscopy after labeling the cells with annexin-V (Fig. 2B). This method was used because it identifies the apoptotic (green fluorescence) as well as necrotic (red fluorescence) cells. As shown by the data, WIN-55,212-2 treatment resulted in a dose-dependent apoptosis in LNCaP cells. In a time-dependent study with a 5 \mu M dose of WIN-55,212-2, there was an increasing trend of apoptotic cells at 72 compared with 48 hours after treatment (Fig. 2C). We next quantified the extent of apoptosis by flow cytometric analysis of the cells labeled with dUTP and PI. LNCaP cells were treated with WIN-55,212-2 (1-10 \mu M) for 24 hours. As shown by the data in Fig. 2D, WIN-55,212-2 treatment resulted in 18.3% and 25.6% of apoptotic cells at a dose of 7.5 and 10 \mu M, respectively. Whereas the induction of apoptosis was almost negligible at the lowest dose (1.0 \mu M) used, the highest dose employed (10 \mu M) resulted in a massive induction of apoptosis as determined by flow cytometry. A similar trend was evident when apoptosis was measured by ladder formation on agarose gel electrophoresis (data not shown).

Effect of WIN-55,212-2 on Androgen Receptor and PSA Protein and mRNA Expression in LNCaP Cells. Androgens are involved in the development and progression of prostate cancer where androgen receptor is assumed to be the essential mediator for androgen action (15, 16). In the next series of experiments, we determined the effect of WIN-55,212-2 on protein and mRNA expression of androgen receptor. In a dose-dependent study, we found that treatment of LNCaP cells with WIN-55,212-2 resulted in a marked decrease in androgen receptor protein expression (Fig. 3A). Relative density data of these immunoblots revealed that the decrease in androgen receptor protein expression was 50% and 90% at 5.0 and 7.5 \mu M of WIN-55,212-2, respectively. In a time-dependent study with 5 \mu M dose of WIN-55,212-2, there was a marked decrease in androgen receptor protein expression and this corresponded with the relative density data showing a decrease of 61% and 69% at 48 and 72 hours, respectively (Fig. 3B). Studies have also shown that modulation in androgen receptor leads to alteration in androgen-responsive genes (17). PSA is an androgen-responsive gene and is regarded as the most sensitive biomarker and screening tool for prostate cancer in humans (18). The dose-dependent effect of WIN-55,212-2 on LNCaP cells showed a significant decrease in PSA protein expression at 5.0, 7.5, and 10 \mu M concentrations when assessing at 24 hours post-treatment (Fig. 3A). Densitometric analysis data revealed that the decrease was 48%, 75%, and 90% at 5.0, 7.5, and 10.0 \mu M concentrations (Fig. 3A). For time-dependent studies, cells were treated with 5 \mu M of WIN-55,212-2 for 24, 48, and 72 hours. Employing Western blot analysis, we found a significant decrease in a time-dependent manner in PSA protein expression. Densitometric analysis revealed a decrease in PSA protein expression by 48% and 60% at 48 and 72 hours, respectively (Fig. 3B). We also evaluated the effect of WIN-55,212-2 on mRNA levels of androgen receptor and PSA. As shown by the real time-PCR analysis data, there was an inhibition in mRNA levels of androgen receptor (Fig. 3C) and PSA (Fig. 3D) at 7.5 and 10.0 \mu M concentrations.

We next examined the effect of WIN-55,212-2 on secreted PSA in LNCaP cells. Employing ELISA technique, we found that treatment of LNCaP cells with WIN-55,212-2 resulted in a dose-dependent decrease in the secreted levels of PSA by 30%, 53%, and 62% at 5.0, 7.5, and 10 \mu M, respectively. At similar doses of WIN-55,212-2, but varying the time point by 48 hours, PSA levels decreased by 53%, 77%, and 80% (Fig. 3E). Furthermore, at 72 hours post-treatment of WIN-55,212-2, secreted PSA levels decreased by 58%, 82%, and 88% (Fig. 3E). From these data, it seems that the decrease in LNCaP cell growth was concomitant with a decrease in androgen receptor protein expression as well as a decrease in both intracellular and secreted PSA levels.

Effect of WIN-55,212-2 on Cell Proliferation Marker, PCNA. We next determined the effect of WIN-55,212-2 on PCNA which serves as a requisite auxiliary protein for DNA polymerase δ-driven DNA synthesis and is cell-regulated (19, 20). The dose-dependent study treatment of LNCaP cells with WIN-55,212-2 (1-10 \mu M) resulted in a significant decrease in protein expression of PCNA. Western blot analysis and relative density of these bands showed that the decrease in protein expression of PCNA was 71% at 7.5 \mu M of WIN-55,212-2 (Fig. 4A). In a time-dependent study, treatment of LNCaP cells with 5 \mu M of WIN-55,212-2 resulted in >50% inhibition in PCNA protein expression at 48 and 72 hours of treatment (Fig. 4B).
Effect of WIN-55,212-2 on VEGF. Because VEGF is a marker for angiogenesis, blocking the angiogenic process may represent a promising way of treating the tumor. Studies have shown that androgen regulates VEGF content in prostate cancer (21). As WIN-55,212-2 treatment resulted in a decrease in androgen receptor expression, the effects on VEGF were also determined. It was observed that WIN-55,212-2 treatment also resulted in a decrease in VEGF protein expression (Fig. 4A). Densitometric analysis data showed a decrease of 40% at 7.5 μmol/L concentration of WIN-55,212-2. In a time-dependent study at 5 μmol/L WIN-55,212-2 treatment, VEGF protein expression decreased in a time-dependent manner (Fig. 4B).

Discussion

It is now well accepted that uncontrolled cellular growth, which may be a result of defects in cell cycle and apoptotic machinery, is responsible for the development of most of the cancers including prostate cancer. Thus, the agents which can modulate apoptosis in cancer cells may be able to affect the
steady-state cell population and may be useful in the management and therapy of cancer. Consistent with this notion, there is a need to develop novel targets and mechanism-based agents for the management of prostate cancer. One of the most exciting and promising areas of current cannabinoid research is the ability of these compounds to control the cell survival/death decision (1). In this study, we found that compared with PrEC and PZ-HPV-7 cells, the expression levels of both cannabinoid receptors CB1 and CB2 were significantly higher in CA-HPV-10 and other human prostate cells LNCaP, DU145, PC-3, and CWR22Rv1. These data suggest that CB1 and CB2 receptors could be a target for novel treatment options for prostate cancer. We also found that mixed CB1/CB2 agonist WIN-55,212-2 treatment of LNCaP cells resulted in a decrease of cell viability as determined by MTT assay at varying doses and time points (Fig. 2A), suggesting the involvement of both CB1 and CB2 in the antiproliferative action of cannabinoids (Fig. 2B). It is widely recognized that apoptosis is an ideal way of elimination of cancer cells and that selective apoptotic events could provide suitable targets for cancer treatment and prevention. In this study, we also observed an increase in apoptosis of LNCaP cells by treatment with WIN-55,212-2. This observation was confirmed by employing confocal microscopy (Fig. 2C and D) and flow cytometry (Fig. 2E). This could be an important observation which might be useful for devising strategies for the management of human prostate cancer because apoptosis is a physiological and discrete way of cell death different from necrotic cell death and is regarded to be an ideal way of cell elimination.

Androgens are essential for the growth, differentiation, and functioning of the prostate as well as in increasing prostate cancer development (22, 23). Many molecular mechanisms have been suggested for the development of recurrent hormone refractory...
tumors. Most of these mechanisms postulate an alteration in the function of the androgen receptor and its signaling pathways (23). The overexpression of androgen receptor in prostate cancer may promote cell growth. Hence, elimination or reducing the androgen receptor in prostate cancer should help in treating this neoplastic disease. As most of the molecular mechanism for the development of prostate cancer involves modulation in the function of androgen receptor and its signaling pathways, we further studied the effect of WIN-55,212-2 on androgen receptor protein and mRNA expression and its subsequent effect on PSA production. Our results indicate that WIN-55,212-2 treatment significantly decreases androgen receptor protein (Fig. 3A) and mRNA expression (Fig. 3C) in LNCaP cells.

PSA belongs to the kallikrein family (17), is a serine protease with highly prostate-specific expression, and is the most widely employed marker in the detection of early prostate cancer. For these reasons, it is considered that agents which could reduce PSA levels may have important clinical implications for prostate cancer. Earlier studies reported that PSA is primarily regulated by androgens (24). This observation was based on the fact that the antiandrogen, cyproterone acetate, had the ability to induce PSA, and that hydroxyflutamide could block androgen and progesterone induction of PSA glycoprotein, thus suggesting that PSA glycoprotein expression is influenced predominantly by androgens via its receptor, and the mutation of the receptor can affect the expression of this gene by steroids other than androgens (24). Recent studies have established that androgen receptor functions as a transcriptional regulator via its binding to androgen response elements within promoter and enhancer regions of PSA. PSA is currently the most accepted marker for assessment of prostate cancer progression in humans and is being detected in the serum of patients with prostate diseases including prostatitis, benign prostatic hypertrophy, and prostate cancer (18). It is reported that in LNCaP cells, androgens regulate PSA glycoprotein expression and mRNA via androgen receptor (25, 26). Our studies show a significant decrease in intracellular, mRNA (Fig. 3D), as well as secreted levels of PSA by WIN-55,212-2 treatment of cells, suggesting that cannabinoid receptor agonists may be exploited to prevent prostate cancer progression.

PCNA recognizes nuclear antigens and its overexpression is associated with increase in PSA serum levels (27). PCNA expression has significant prognostic value and it seems to be a significant biomarker in prognosis and treatment of prostate cancer (27). Our results also suggest that concomitant with the decrease in PCNA protein expression (Fig. 4A), there was a decrease in PSA serum levels following WIN-55,212-2 treatment (Fig. 3E).

VEGF is a ubiquitous cytokine that regulates embryonic vasculogenesis and angiogenesis. Normal prostate epithelium expresses low levels of VEGF, whereas premalignant lesions have increased VEGF expression, which is additionally increased in prostate carcinoma (28). Studies have shown that cannabinoid treatment markedly reduced the expression of VEGF in gliomas, the most potent proangiogenic factor and also of angiopoietin 2, which contributes to the angiogenic process by preventing vessel maturation (29). Our results showed that treatment of LNCaP with WIN-55,212-2 inhibits growth and VEGF protein expression (Fig. 4A and B).

Recently, cannabinoids have received considerable attention due to their diverse pharmacologic activities such as cell growth inhibition, anti-inflammatory effects, and tumor regression. Our results suggest that treatment of androgen-responsive human prostate carcinoma LNCaP cells resulted in a decrease in intracellular and secreted levels of PSA, with concomitant inhibition of androgen receptor, cell growth, and induction of apoptosis. We conclude that cannabinoids should be considered as agents for the management of prostate cancer. If our hypothesis is supported by in vivo experiments, then the long-term implications of our work could be to develop non–habit-forming cannabinoid agonist(s) for the management of prostate cancer.