The Role of CD44 in Glucose Metabolism in Prostatic Small Cell Neuroendocrine Carcinoma

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Abstract

While prostatic adenocarcinomas are relatively indolent, some patients with advanced adenocarcinomas recur with small cell neuroendocrine carcinoma which is highly aggressive and lethal. Because glycolysis is a feature of malignancy and the degree of glycolysis generally correlates with tumor aggressiveness, we wanted to compare the metabolic differences and the molecular mechanisms involved between the two tumor types. In this study, and based on previous characterization, LNCaP and PC-3 prostate cancer cell lines were selected as models of prostatic adenocarcinoma and small cell neuroendocrine carcinoma, respectively. In addition to measuring glucose consumption, lactate secretion, and reactive oxygen species (ROS) levels, we performed metabolic profiling in these two model systems. The role of CD44 was studied by RNAi and lentivirus-mediated overexpression. Expression of key enzymes in glycolysis was studied using human tissue microarrays containing benign prostate, adenocarcinoma, and small cell neuroendocrine carcinoma. Results showed that glycolytic features of PC-3 cells were higher than that of LNCaP cells. PFKFB4 was overexpressed in human small cell carcinoma tissue versus adenocarcinoma tissue. CD44 regulated glucose metabolism, intracellular ROS, and cell proliferation in PC-3 cells. Inhibition of CD44 also sensitized PC-3 cells to carboplatin. In conclusion, this study suggests different pathways of glucose metabolism contribute to the disparate biologic behaviors of these two tumor types.

Implications: CD44 is an important regulator of glucose metabolism in small cell neuroendocrine carcinoma and may be an important therapeutic target. Mol Cancer Res; 14(4): 344–353. ©2016 AACR.

Introduction

Prostate cancer is the most common cancer in men in the United States with 233,000 new cases and approximately 30,000 deaths estimated in 2014 (1). Hormonal therapy, to lower androgen levels and/or block androgen receptor (AR) function, is currently used to treat advanced and metastatic prostate cancer, which provides temporary symptomatic relief. However, after an average of 18 months, the cancer invariably recurs as castration-resistant prostate cancer (CRPC). To treat CRPC, newer agents have been approved, including enzalutamide and abiraterone, which better block AR signaling and inhibit intratumoral androgen synthesis, respectively. Unfortunately, their effects are generally short-lived, and many patients will quickly develop resistance. Importantly, although well over 90% of primary prostate cancers are classified as adenocarcinoma (AdenoCa) with glandular features of carcinoma known as small cell neuroendocrine carcinoma (SCNC; ref. 2). In an ongoing large-scale study of metastatic prostate cancer in 300 men after treatment with enzalutamide and abiraterone, we have observed that about 20% of the biopsied tumors are SCNC (Targeting Resistance Pathways in Metastatic Castration Resistant Prostate Cancer; AACR). In contrast with the relatively slow-growing and indolent AdenoCa, SCNC is highly aggressive and rapidly lethal. Histologically, SCNC is composed of neuroendocrine cells that do not form glands, are negative for AR and PSA, and are refractory to currently available therapies (3). Understanding the fundamental molecular mechanism of SCNC and discovering novel therapeutics are urgent unmet needs.

In contrast with benign cells, cancer cells generally follow the Warburg effect, displaying increased glycolysis for energy production (4). This feature is accompanied by suppression of mitochondrial respiration and increased flux through the pentose phosphate pathway (PPP). The Warburg effect, initially described in 1924, has received renewed attention recently in the field of cancer metabolism due to the widespread clinical application of fluorodeoxyglucose (FDG)-PET imaging which is based on increased glucose uptake by cancer cells. Unlike most tumors, however, FDG-PET imaging cannot detect localized, untreated...
Glucose Metabolism in Prostatic SCNC

prostate cancer (5), but can image SCNC (6), suggesting that metabolic differences may be an underlying mechanism for the vastly different biologic behaviors of AdenoCa and SCNC. In addition, literature has shown that prostate AdenoCa utilizes mitochondrial respiration which is unique to the metabolism displayed in other tumor cells (7).

CD44 is a cell-surface protein with functions in many biologic processes, including cell adhesion and proliferation. In cancer cells, CD44 plays a role in tumor growth and metastasis through multiple means: angiogenesis, cell survival, cell migration, etc. (8, 9). It has also been suggested that CD44 is a cell-surface marker for cancer stem cells. A recent report showed that CD44 was involved in regulating the glycolytic pathway in colorectal cancer and lung carcinoma cell lines (10). This report also found that expression of CD44 in human cancer cell lines induces a more invasive and drug-resistant phenotype (11). Our lab has reported that CD44 expression is a feature of prostatic SCNC, whereas the bulk tumor cells of AdenoCa are negative for CD44 (12). With reports that p53, a commonly mutated tumor suppressor in SCNC, regulates CD44 (8) and evidence that CD44 interacts with PKM2 (10), the glycolytic isozyme of pyruvate kinase, we propose that CD44 may be modulating metabolism in SCNC. The goal of this study is to investigate the differences in glucose metabolism between the two forms of prostate cancer and understand the underlying molecular mechanism. We hypothesize that prostatic SCNC is more glycolytically active than AdenoCa and CD44 is a key regulator of glucose metabolism that maybe a potential therapeutic target for SCNC.

Materials and Methods

Cell lines
Prostate cancer cell lines LNCaP, VCaP, PC-3, CWRR1, DU145, and 22RV1 cells were obtained from the American Type Culture Collection. The cells were cultured in RPMI-1640 medium supplemented with 10% FBS (Hyclone) and penicillin–streptomycin. The cells were incubated in a humidified incubator at 37°C and 5% CO2.

LNCaP and PC-3 cell lines were authenticated by DDC Medical.

Preparation of low glucose RPMI-1640 medium
Ten percent glucose RPMI-1640 medium was prepared with glucose-free RPMI-1640 (Gibco BRL) and dextrose (EMD chemicals Inc.), and supplemented with 10% FBS (Hyclone) and penicillin–streptomycin.

Measurement of glucose consumption and lactate secretion
LNCaP and PC-3 cells were seeded in a 96-well plate at 8,000 cells/well and incubated in low glucose RPMI-1640 medium with 10% FBS and penicillin–streptomycin for 18 hours. Glucose concentrations in the media were determined by a glucose assay kit (Sigma) following the manufacturer’s instructions.

Lactate concentration was determined by using an enzymatic method. Briefly, the media were collected, and hydrazine buffer, nicotinamide adenine dinucleotide (NAD), and lactate dehydrogenase (LDH) (Sigma) were added. Distilled water was used as blanks. Absorbance at 340 nm was measured to determine the lactate levels.

Measurement of reactive oxygen species
Intracellular reactive oxygen species (ROS) levels were detected using H2DCF-DA (Invitrogen). Cells were cultured in a 96-well black assay plate (Coming) and washed once with PBS before incubation with serum-free media containing 10 μmol/L H2DCF-DA for 30 minutes at 37°C. ROS levels were determined at excitation and emission wavelengths of 485 and 520 nm, respectively, by using a Synergy H1 Hybrid multi-mode microplate reader (Bio-Tek Instruments). Control media without dye were used to determine background fluorescence. Cell numbers were normalized before measurement.

Metabolic profiling
LNCaP and PC-3 cells were plated in 6-well plates (500,000 cells/well; 3 wells per condition) and incubated overnight. Cells were incubated in medium with dialyzed FBS containing labeled glucose (4.5 g/L of U-13C6 D-glucose; Cambridge Isotope Laboratories, Inc.) for 24 hours. For spent media samples, 20 μL was collected from each well (in triplicate). For cell samples, cells were rinsed with cold 150 mmol/L ammonium acetate. Then cells were scraped off with 1 mL of 80% cold methanol. An internal standard of 10 nmol norvaline was added, and the cell suspension was spun down at top speed for 5 minutes at 4°C. The supernatant was transferred into a glass vial, and the pellet was resuspended in 200 μL 80% methanol, spun again, and the supernatant was combined into the glass vial. The metabolites were dried in EZ-ZElite evaporator at 30°C using aqueous program and stored at −80°C until mass spectrometry was performed.

Samples were resuspended in 70% acetonitrile, and 5 μL were injected onto a LunaNH2 (150 mm × 2 mm; Phenomenex) column. Analysis was performed with an UltiMate 3000RSLC (Thermo Scientific) couple to a Q Exactive mass spectrometer (Thermo Scientific). The spectrometer ran with polarity switching (+4.00 kV/−4.00 kV) in full scan mode containing an m/z range of 70 to 1,050. Metabolites were separated with (1) 5 mmol/L ammonium acetate (pH 9.9) and (2) acetonitrile. The start of the gradient was 15% to 90% (1) for 18 minutes, then an isocratic step for 9 minutes, and back to 15% (1) for 7 minutes. The metabolites and their isotopomers were quantified using TraceFinder 3.1 with accurate mass measurements (≤3 ppm) and retention times. Correction for naturally occurring 13C was considered for isotopologue distribution measurements as described (13). R programming language was used for data analysis.

Immunohistochemistry
A total of 146 cases were studied, including 73 cases of normal prostate, 60 cases of prostatic adenocarcinoma, and 13 cases of SCNC. All samples were built into tissue microarrays (TMA), and each sample was represented by three cores. Sections were deparaffinized with xylene and rehydrated through graded ethanol, cleared with xylene, and cover-slips.

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Evaluation of immunohistochemical staining

All sections were blindly analyzed by two experienced investigators (W. Li and Z. Li). The staining intensity was scored as negative (0), weakly positive (1), moderately positive (2), and strongly positive (3). Results of the cores from the same case were averaged to arrive at a score for the case, which were analyzed by a team of statisticians (G. Li and L. Du).

RNA interference

CD44 and PFKFB4 were knocked down with siRNA using siTRAN 1.0 Transfection Reagent (OriGene). PC-3 cells were transfected with 10 nmol/L of CD44 siRNA (SR300683; OriGene), PFKFB4 siRNA (SR303465; OriGene), or Universal scrambled negative control siRNA duplex (SR30004; OriGene) for 48 hours. siRNA-mediated knocking down was confirmed by Western blot analyses for CD44 (ABGENT) and PFKFB4 (Abcam).

Infection with retrovirus vector expressing CD44

As previously described (14), lentiviral vector expressing CD44 was introduced into LNCaP cells through spin infection at 1,500 rpm for 45 minutes with 8 µg/mL polybrene. After 48 hours, cells were selected in puromycin and expanded. The expression of CD44 was analyzed using Western blot analysis.

Western blot

Western blot was performed as described (14). Briefly, cultured cells were lysed with cell lysis buffer. The protein concentration of each sample was measured with a Bio-Rad protein assay kit. Equal amounts of protein were separated on 8% SDS-polyacrylamide gels followed by transfer to polyvinylidene fluoride membrane. The membrane is blocked with nonfat milk, incubated with primary antibody at 4°C overnight, followed by secondary antibody incubation, and developed with the West Femto Kit (Pierce).

qPCR analysis

Total RNA was extracted from cultured cells with animal RNA miniprep plus kit (Bioland Scientific LLC) according to the manufacturer’s protocol, normalized based on the amount of β-actin mRNA. The following specific forward and reverse primers were used: PFKFB4: 5’-CAGAACAGCTGCCCTACCTC-3’ and 5’-GCCTCCTGAGGACTGCTGA-3’.

Cell proliferation assay

Cell proliferation and viability were quantified using the Quick Cell Proliferation Colorimetric Assay Kit Plus (Biovision). Briefly, cells were cultured at 50,000 cells/well in a 96-well plate and incubated. At collection time, 10 µL/well of tetrazolium salt (WST) reagent was added to each well and incubated at 37°C for 4 hours. Then the plate was shaken for 1 minute, and the absorbance was measured at 420 nm with the reference wavelength of 650 nm on a plate reader. Bicinchoninic acid (BCA) assay was performed to normalize to total protein amount.

Clonogenic cell survival assay

Cells were transiently transfected with respective siRNAs. After 24 hours, cells were plated and treated (1 µmol/L for carboplatin) in 60 mm dishes (500 cells/dish). After 24 hours, cultured cells were washed twice in complete growth medium and maintained for 10 to 14 days. The media were then discarded, washed with PBS, and stained with crystal violet. Surviving colonies were counted and imaged to determine plating efficiency.

Statistical analysis

All experiments were performed in triplicates, and the data are presented as mean ± SD. Statistical significance was determined using the unpaired Student t test. A P value of <0.05 was considered significant. Data analyses were performed using SAS 9.4 (SAS Institute Inc.) and Microsoft Excel 2007.

Results

Increased glycolytic features in PC-3 cells compared with LNCaP cells

Cancer is characterized by increased glycolysis, regardless of oxygen availability, and the glycolytic rate generally correlates with tumor aggressiveness. Prostatic adenocarcinoma and SCNC are two histologically distinct tumor types with different clinical courses. The former usually pursues an indolent course while the latter is highly aggressive and invariably lethal. We therefore aimed to determine whether glucose metabolism may be different between the two tumor types. We used LNCaP cells and PC-3 cells to model adenocarcinoma and SCNC, respectively, based on our previous characterization of the two cell lines (15). A glycolytic phenotype is characterized by increased glucose consumption and lactate secretion along with reduced ROS levels (10). Thus, we measured the glucose and lactate concentrations of media from cultured LNCaP and PC-3 cells. Consistent with our hypothesis, we found that media from cultured PC-3 cells contained lower glucose levels and higher lactate levels, suggesting that PC-3 cells consume more glucose and secrete more lactate than LNCaP cells (Fig. 1A and B). With increased metabolic flux toward the glycolytic pathway, ROS levels decrease, due to reduced mitochondrial respiration. We measured ROS levels within LNCaP and PC-3 cells and found significantly lower ROS levels in PC-3 cells (Fig. 1C).

To further confirm these results, we performed metabolic profiling on extracted samples from both cells and media with glucose labeling to trace the metabolic pathways. With negative values from media samples indicating that the metabolite was consumed, we see that PC-3 cells consume more glucose than LNCaP cells, as expected (Fig. 2A). Concordantly, PC-3 media samples showed higher lactate levels indicating increased lactate secretion by these cells compared with LNCaP cells (Fig. 2B). When looking at the extracted samples from the cells with labeled glucose, we find more unlabeled (M0) tricarboxylic acid (TCA) cycle intermediates in PC-3 cells, whereas LNCaP cells show increased glucose labeling in these intermediates (Fig. 2C–F). Decreased labeling in the TCA cycle metabolites is consistent with a shift in metabolic flux away from mitochondrial respiration and toward other glycolysis branch pathways. The increased M2 labeling in citrate levels in PC-3 cells suggests that citrate is being used to fuel other biosynthetic pathways for rapid cell proliferation (Fig. 2G). All of these observations are consistent with our hypothesis that glycolysis is favored by the highly aggressive tumor cells of prostatic SCNC compared with the relatively slow-growing and indolent tumor cells of adenocarcinoma.

Expression of PFKFB4 in prostatic adenocarcinoma and SCNC

Several metabolic enzymes involved in glycolysis have shown to be potential therapeutic targets for certain cancers. One enzyme of importance is 6-phosphofructo-2-kinase/fructose-2,6-biphosphatase 4 (PFKFB4), an isofrom of phosphofructokinase 2
(PFK2), which is required for prostate cancer cell survival and balancing glycolytic activity and antioxidant production in prostate cancer cells (16). We thus wanted to investigate if the expression of PFKFB4 is differentially expressed in adenocarcinoma versus SCNC.

We first determined the expression levels of PFKFB4 protein in various prostate cancer cell lines. Western blot results showed that, compared with naïve AdenoCa LNCaP and VCaP cells, or to castration-resistant AdenoCa C4-2 and CWRR1 cells, prostatic SCNC PC-3 and DU145 cells express relatively higher levels of PFKFB4. Of note, PFKFB4 protein level is significantly higher in DU145 cells (Fig. 3A).

We then performed immunohistochemistry using multiple TMAs containing benign prostate (n = 73), prostatic adenocarcinoma (n = 60), and prostatic SCNC (n = 13). PFKFB4 was differentially expressed in benign and malignant prostate tissues (Fig. 3B). Most cases of benign prostate tissue were negative for PFKFB4, with only occasionally scattered luminal secretory cells being positive (Fig. 3C). The adenocarcinoma cells were weakly or moderately stained (Fig. 3D). In contrast, most of the SCNC cells were strongly and diffusely positive for PFKFB4 (Fig. 3E). Detailed scoring of the staining and statistical analysis showed that the expression of PFKFB4 in benign prostate tissue was lower than that of prostatic adenocarcinoma (P < 0.001), and its expression was significantly higher in SCNC than in adenocarcinoma (P < 0.001).

**CD44 is a regulator of glucose metabolism and PFKFB4 in prostate cancer cells**

CD44 has been implicated in controlling glucose metabolism, and it is also expressed in prostatic SCNC but not adenocarcinoma (12). Similarly, our lab has shown that CD44 is expressed in PC-3 cells and not in LNCaP cells (15). Therefore, we further hypothesized that CD44 may be a regulator of glucose metabolism in prostatic SCNC. We performed CD44 knockdown studies in PC-3 cells and CD44 overexpression studies in LNCaP cells.
studies in LNCaP cells to study changes in glucose metabolism. When CD44 was knocked down in PC-3 cells, glucose consumption levels decreased significantly compared with that in the parental cells (P < 0.001; Fig. 4A). Conversely, overexpression of CD44 in LNCaP cells led to an increase in glucose consumption (P < 0.001; Fig. 4B). These findings suggest that CD44 expression in prostatic SCNC is involved in the tumor’s increased glycolytic activity.

As mentioned previously, suppressing mitochondrial respiration in malignant tumor cells reduces the production of ROS and leads to tumor cells’ resistance to various therapies (17). We therefore investigated the relationship between CD44 expression and intracellular ROS levels in prostate cancer cells. After CD44 knockdown, we observed significantly increased ROS levels in PC-3 cells (Fig. 4C). Similarly, overexpression of CD44 decreased ROS production in LNCaP cells (Fig. 4D). As a glycolytic regulator, we examined whether CD44 affected PFKFB4 expression because PFKFB4 levels showed a positive correlation with SCNC. CD44 ablation in PC-3 cells showed a reduction in both protein and mRNA levels of PFKFB4 (Fig. 4E and F). Overall, the data show that CD44 plays a key role in altering glucose metabolism and affecting glycolytic enzymatic activity in prostatic SCNC.

Figure 2.
Differential metabolic profiling of LNCaP and PC-3 cells and media. Metabolite levels were measured in the spent media for glucose (A) and lactate (B). TCA cycle metabolite isotopomer levels were measured from cell samples with uniformly labeled glucose: alpha-ketoglutarate (C), fumarate (D), malate (E), succinate (F), and citrate (G). *, P < 0.05; **, P < 0.01; ***, P < 0.001.
CD44 knockdown inhibits cell proliferation and sensitizes SCNC tumor cells to carboplatin

Because glycolytic activity often correlates with cell proliferation, we examined the role of CD44 in the regulation of cell proliferation. Our study showed that the proliferation of PC-3 cells was significantly inhibited after CD44 knockdown (Fig. 5A). Given that glucose consumption is necessary for the survival and proliferation of cancer cells, and CD44 knockdown results in a decrease in glucose consumption and proliferation of PC-3 cells as described above, we hypothesized that the expression of CD44 may have therapeutic implications in SCNC. This hypothesis would be consistent with a previous report that the expression of CD44 in human cancer cell lines induces a more invasive and drug-resistant phenotype (11).

Conventional and newer therapeutic prostate cancer drugs targeting the AR signaling pathway are ineffective for SCNC as these tumor cells do not express AR. Because of this, many clinicians treat SCNC with carboplatin, a cisplatin analogue, because pulmonary small cell neuroendocrine carcinoma is sensitive to this drug. However, the therapeutic benefit of carboplatin for prostatic SCNC is usually limited (18). Our evidence suggests that CD44 and PFKFB4 may be involved in the drug-resistant phenotype observed in SCNC. This phenotype is associated with glycolysis and low ROS levels. Reports have shown that manipulating the metabolic flux within tumor cells, by shifting to mitochondrial respiration, increasing ROS levels, or inactivating the PPP, can contribute to increased drug sensitivity (10). In addition, CD44-expressing cancer cells have shown chemoresistance in previous studies. We therefore investigated whether CD44 or PFKFB4 ablation would sensitize PC-3 cells to carboplatin. First, we found that PC-3 cells with PFKFB4 knockdown also significantly inhibited cell proliferation (Fig. 5B). We then discovered that sensitivity of PC-3 cells to carboplatin treatment was significantly enhanced with knockdown of either CD44 or PFKFB4 (Fig. 5C and D), implying that inhibition of these key glycolytic proteins may alter the metabolic flux and sensitize SCNC tumor cells to carboplatin.

Discussion

The key finding in this study is that PC-3 cells, characteristic of SCNC, have increased glycolytic activity compared with LNCaP cells which is of clinical relevance. Most prostate cancer patients are initially diagnosed with adenocarcinoma with tumor cells expressing luminal differentiation markers AR and PSA. These tumors are highly curable when they are low grade and localized.
Advanced adenocarcinomas are treated with androgen ablation therapy (e.g., Lupron). Novel drugs inhibiting intratumoral androgen synthesis (abiraterone) or better inhibiting AR (enzalutamide) can be used after conventional hormonal therapy has failed. Importantly, a significant proportion of patients will recur in the form of SCNC after the above therapies (2).

SCNC is extremely aggressive, rapidly fatal, and does not respond to the above therapies

We have previously shown that LNCaP cells possess important features of adenocarcinoma with cells expressing AR and PSA dependent on androgen for proliferation (15). Similar to adenocarcinoma, LNCaP cells do not express CD44 and have wild-type p53. In contrast, PC3 cells are characteristic of SCNC with tumor cells being negative for AR and PSA but expressing high levels of neuroendocrine markers. Similar to SCNC, PC-3 cells express CD44 and contain mutant p53 (15). These findings validate the efficacy of performing the studies mentioned in LNCaP and PC-3 cell lines.

In this study, we demonstrate that glucose metabolism may be an important difference between the two tumor types that could potentially be targeted for the treatment of prostatic SCNC. This conclusion is based on biochemical studies in cell line models as well as immunohistochemical studies in human cancer tissues. Our study provides a metabolic basis for the difference of biologic behavior of the two tumors, revealing novel strategies to target the metabolic pathway in SCNC. Furthermore, we showed that CD44, a cell-surface molecule expressed in SCNC but not adenocarcinoma, appears to be a critical regulator of glucose metabolism, specifically in SCNC which displayed a glycolytic phenotype. CD44 ablation in PC-3 cells inhibits glucose consumption and increases ROS production, possibly through changes in activity and expression of important glycolytic enzymes. In addition, knocking down CD44 inhibits cell proliferation and increases sensitivity of PC-3 cells to carboplatin, a chemotherapeutic agent currently used to treat aggressive prostatic SCNC. Overall, our evidence suggests that CD44 may be an important and selective molecular target for prostatic SCNC.

**Figure 4.** CD44 modulates glucose metabolism, intracellular ROS levels, and PFKFB4 expression. A, PC-3 cells were transfected with siRNA targeting CD44, and glucose consumption was measured. Glucose consumption decreased with siCD44 (*, P = 0.003). B, lentivirus-mediated overexpression of CD44 protein or control in LNCaP cells. CD44 overexpression led to increased glucose consumption (*, P < 0.001). C, intracellular ROS increased after siRNA-mediated CD44 ablation in PC-3 cells (*, P < 0.001). D, intracellular ROS decreased after lentivirus-mediated CD44 overexpression in LNCaP cells (*, P < 0.001). E, CD44 knockdown in PC-3 cells results in reduced PFKFB4 protein levels (*, P < 0.001). F, CD44 knockdown in PC-3 cells results in reduced expression of PFKFB4 mRNA (*, P < 0.001).
These findings are consistent with previous reports. For example, Singh and colleagues showed that HIV-1 120-kDa glycoprotein variant IIIB (gp120-IIIB) causes apoptosis of PC-3 and DU145 cells through its interaction with CXCR4, and the inhibitory function is associated with downregulation of CD44 (19). Another report showed that antitumor activity of hyaluronic acid synthesis inhibitor 4-methylumbelliferone in prostate cancer cells is also associated with a downregulation of CD44 (20).

The function of CD44 in tumor invasion and metastases, especially in bone metastases, has been reported (21–23). Recent studies revealed that CD44 may be involved in metabolic regulation in colorectal cancer and breast cancer cells. Our studies demonstrate that in prostate cancer, CD44 is involved in the aggressive behavior of the tumor cells when there is a phenotypic switch from adenocarcinoma to SCNC which likely occurs through regulation of glucose metabolism.

PFKFB4 is an important enzyme in regulating glucose metabolism that catalyzes the synthesis and degradation of fructose-2,6-biphosphate (24,25). Silencing of PFKFB4 should divert glucose-6-phosphate toward the glycolytic pathway, thereby depleting the PPP (16). Knockdown of CD44 reduces PFKFB4 expression, which can lead to decreased levels of NADPH. NADPH is required to maintain cellular stores of glutathione, an antioxidant preventing ROS accumulation. This may provide an explanation for the increased ROS levels observed after CD44 ablation in PC-3 cells (Fig. 6). The above studies involving PFKFB4 manipulation are consistent with results obtained in a previous study (16).

Platinum-based chemotherapy is the first-line therapy for small cell lung cancer (26,27); however, it has limited efficacy in advanced prostate cancer (18,28), and CD44-expressing cancer cells have shown chemoresistance (10). Certain anticancer drugs, including cisplatin, kill cancer cells by inducing ROS generation which causes apoptosis (29). With evidence that CD44 depletion altered ROS levels, these measured increases in ROS in PC-3 cells should enhance sensitivity to carboplatin. We demonstrated that CD44 knockdown increased sensitivity of PC-3 cells to carboplatin, likely through a metabolic shift toward mitochondrial respiration and ROS production. This finding has important clinical implications. CD44 is considered a marker of cancer stem cells including prostate cancer stem cells (30). We have demonstrated that the bulk tumor cells in prostatic adenocarcinoma are negative for CD44, whereas the tumor cells of SCNC are positive (12). Interestingly, PC-3 cells, but not LNCaP cells, express CD44 (15,31), and the PC-3 human prostate carcinoma cell holoclones contain self-renewing tumor-initiating cells with high levels of CD44 (32). Our studies further support the concept that CD44 may be a therapeutic target for aggressive SCNC, due at least in part to its modulation of glucose metabolism. In conclusion, SCNC, the
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Development of methodology: W. Li, A. Cohen, D. Braas, J. Huang

Figure 6.
A potential pathway for CD44 moleculeally regulating cell proliferation and survival through PFKFB4 in human prostatic SCCN.

More aggressive form of prostate cancer shows increased glycolytic activity in comparison with adenocarcinoma, and CD44 may be playing a role in regulating this altered metabolic phenotype.

Disclosure of Potential Conflicts of Interest
No potential conflicts of interest were disclosed.

Authors’ Contributions
Conception and design: W. Li, A. Cohen, Y. Sun, G. Li, J. Huang
Development of methodology: W. Li, A. Cohen, D. Braas, J. Huang

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