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ORIGINAL ARTICLE

The differential effects of prostate stromal cells derived from different zones on prostate cancer epithelial cells under the action of sex hormones

Qi Jiang, Bang-Min Han, Fu-Jun Zhao, Yan Hong and Shu-Jie Xia

It is well known that prostate cancer (PCa) occurs predominantly in the peripheral zone (PZ), whereas benign prostatic hyperplasia (BPH) typically develops in the transition zone. To identify possible mechanisms underlying zonal differences, we compared the effects of prostate stromal cells derived from the peripheral zone (PZsc) and the transition zone (TZsc) on a PCa epithelial cell line (PC3) in the presence of sex hormones. First, we observed that androgen receptor (AR) mRNA was more highly expressed in PZsc than TZsc when the cells were treated with dihydrotestosterone (DHT) and β -oestradiol (E2) (*P*<0.05). By ELISA, we looked for differences in the secretion of peptide growth factors from PZsc and TZsc. We found that keratinocyte growth factor (KGF) secretion increased with increasing concentrations of DHT (*P*<0.01) and was higher in PZsc than TZsc. Under treatment with DHT plus E2, PZsc secreted more transforming growth factor- β 1 (TGF- β 1) than TZsc, but this pattern was reversed when the cells were treated with E2 only. With increasing concentrations of DHT, insulin-like growth factor-1 (IGF-1) secretion increased in PZsc but decreased in TZsc. To further characterize the effects of PZsc and TZsc on PC3 cells, we developed a coculture model and performed MTT assays, Western blot analysis and real-time RT-PCR. We found that PZsc promoted PC3 cell proliferation and progression better than TZsc, particularly when treated with 10 nmol I⁻¹ DHT plus 10 nmol I⁻¹ E2. In conclusion, our data suggest that PZsc may have a greater capacity to induce PCa development and progression than TZsc *via* growth factors regulated by sex hormones. These findings provide possible mechanisms underlying zonal differences in prostate diseases, which may aid the search for novel therapeutic targets for PCa. *Asian Journal of Andrology* (2011) **13**, 798–805; doi:10.1038/aja.2011.22; published online 18 July 2011

Keywords: androgen; growth factor; oestrogen; prostate cancer; stromal cell

INTRODUCTION

Prostate cancer (PCa) and benign prostatic hyperplasia (BPH) are among the most common diseases that affect elderly men. PCa occurs predominantly in the peripheral zone (PZ) of the prostate, whereas BPH typically develops in the transition zone (TZ).¹⁻⁴ The molecular mechanisms underlying this phenomenon are largely unknown. Several studies have shown that the expression of certain genes differs between these two zones.^{5–9} Feneley *et al.*¹⁰ reported that the epithelial content was significantly greater in the TZ than in the PZ. Conversely, concentrations of androgen receptor (AR) activity were higher in the PZ epithelium than in the TZ epithelium, but higher concentrations of oestrogen receptor activity were detected in the TZ zone stroma.¹⁰

The prostate is a hormone-responsive organ. Androgens play a vital role in prostate development, maintenance of tissue function and the pathogenesis of prostate disease. The stroma is a key mediator of androgenic effects on the epithelium and can regulate both the fate of the epithelial cells and potentially the initiation and progression of PCa. Prostatic epithelial cells are dependent on androgen-responsive stromal cells for development, differentiation and survival.¹¹ Thus, alterations in the stromal microenvironment are sufficient to promote malignant transformation of human

prostatic epithelial cells. When the non-tumourigenic human prostatic epithelial cell line BPH-1 was recombined with urogenital sinus mesenchyme (UGM) or stromal cells from PCa, BPH-1 cells were induced to undergo malignant transformation.^{12,13} Many research groups have also shown that a dynamic stromal-epithelial interaction plays a critical role in the progression and metastasis of PCa.¹⁴⁻²⁰ Growth factors play a central role in mediating the interactions between epithelial and stromal cells.²¹ The stromal-epithelial interaction is also critical for mediating the effects of these molecules on epithelial cells. Sex hormones regulate epithelial proliferation via a paracrine mechanism. The action of androgens on cellular proliferation and death is mediated indirectly through the actions of various growth factors via an intricate stromal-epithelial interaction. Stromal cells express certain growth factors that control the proliferation and differentiation of the prostate epithelium, demonstrating a critical role for the stroma in epithelial growth and homeostasis.

In the present study, we hypothesized that the mechanisms involved in the stromal–epithelial interaction are different in the PZ and the TZ and that this difference may be responsible for the distinct zonal localisation of prostate diseases. The study of such zone-specific roles in tumourigenesis and the progression of PCa may provide novel

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Received: 29 November 2010; Revised: 11 January 2011; Accepted: 1 March 2011; Published online: 18 July 2011

therapeutic strategies for the control and treatment of PCa. To elucidate the roles of zone-specific stromal–epithelial interactions in prostate diseases, we established a coculture model and compared the effects of prostate stromal cells derived from different zones on PCa epithelial cells in the presence of sex hormones. Our data suggest that stromal cells from the PZ may have a greater capacity to induce PCa development, growth and progression than TZsc *via* growth factors regulated by sex hormones.

MATERIALS AND METHODS

Cell culture

Fresh human prostate specimens were obtained post-mortem from consenting normal organ donors between the ages of 30 and 45 years at Shanghai First People's Hospital (Shanghai, China) with the approval of the Shanghai First People's Hospital Medical Ethics Committee. Histopathological analysis confirmed the zonal-specific and normal morphology of the prostate tissue (tissues with BPH or PCa were excluded). Prostate specimens, cut according to McNeal's zonal anatomy,²² were minced (1 mm³ pieces) and then digested for 8 ± 2 h at 37 °C in RPMI 1640 medium (Gibco, Rockville, MD, USA) with collagenase I (200 U ml⁻¹). Stromal cells were separated from epithelial cells by discontinuous gradient centrifugation.²³ The stromal cells derived from the PZ (PZsc) and stromal cells derived from the TZ (TZsc) were identified by immunocytochemistry (IHC)⁸ and cultured with RPMI 1640 supplemented with 10% foetal bovine serum (Gibco, Melbourne, Australia) and antibiotics (100 mg ml⁻¹ streptomycin and 100 IU ml⁻¹ penicillin) (Gibco BRL, Grand Island, NY, USA) at 37 °C under 5% CO2 and a humidified atmosphere. The stromal cells were used at passages 3-5. The PCa cell line PC3 was obtained from the American Type Culture Collection (Rockville, MD, USA) and cultured in RPMI 1640 medium with 10% foetal calf serum (Gibco, Melbourne, Australia).

Coculture model

A coculture model of PZsc or TZsc with PC3 cells was constructed in six-well plates (for Western blot and RT-PCR) or 24-well plates (for MTT) using cell inserts with 0.4-µm polyethylene terephthalate (PET) membranes (Corning, Lowell, MA, USA) (**Figure 1**). PC3 cells were cultured alone or cocultured with the stromal cells. PZsc or TZsc cells were seeded on six-well plates at a density of 1×10^5 cells per well or on 24-well plate inserts at a density of 1×10^5 cells per insert or on 24-well plate inserts at a density of 1×10^5 cells per insert or on 24-well plate inserts at a density of 1×10^5 cells per insert or on 24-well plate inserts at a density of 1×10^5 cells per insert or on 24-well plate inserts at a density of 1×10^4 cells per insert. They were all cultured in RPMI 1640 medium supplemented with 10% charcoal-stripped foetal bovine serum and antibiotics at 37 °C under 5% CO₂. Twenty-four hours after seeding, cells were cultured either in the absence of dihydrotestosterone (DHT) (Wako, Tokyo, Japan) and



Figure 1 Coculture model. PC3 cells were seeded on the insert chamber with PET bottom (the upper chamber of the model). The stromal cells (PZ or TZ) were seeded on regular plate (the lower chamber of the model). PET, polyethylene terephthalate; PZ, peripheral zone; TZ, transition zone.

 β -oestradiol (E2) (Sigma, St Louis, MO, USA) (as a control) or in medium containing E2 (10 nmol l^{-1}) and a series of concentrations of DHT (0, 1, 10 or 100 nmol l^{-1}). Media were changed daily.

ELISA analysis

Approximately 2×10^5 PZsc or TZsc cells were plated per well in sixwell plates. After overnight attachment and growth, the cells were washed twice with phosphate-buffered saline and fed with serum-free RPMI 1640 containing no hormones (control) or E2 (10 nmol l⁻¹) and increasing concentrations of DHT (0, 1, 10 or 100 nmol l⁻¹). Cell culture supernatants were collected after 24 h, centrifuged at 750g for 20 min and stored at -80 °C until use. The concentrations of transforming growth factor- β 1 (TGF- β 1), keratinocyte growth factor (KGF) and insulin-like growth factor-1 (IGF-1) were measured by ELISA according to the manufacturer's protocol (R&D Systems, Minneapolis, MN, USA). Each experiment was performed in triplicate. Cells were counted after supernatants had been removed, and measurements were normalized by cell number. Results were given in pg ml⁻¹ per 1×10⁵ cells.

Cell proliferation assay

The effect of stromal cells on the growth of prostatic epithelial cells was determined using a tetrazolium assay in which mitochondrial enzymes in metabolically active cells decompose a tetrazolium salt, MTT, to a coloured formazan product with a maximum absorption at 595 nm. As described in our coculture model, 5×10^3 PZsc or TZsc cells were seeded per well in 24-well plates, and 1×10^4 PC3 cells were seeded per well in 24-well plate transwell inserts. After 24 h of attachment, cells were cultured in the presence of E2 and DHT as described above. Cells were then incubated for 24, 72, 120 or 168 h. At the end of the incubation period, 100 μ l of MTT (5 g l⁻¹) was added to each well, and cells were further incubated for 4 h at 37 °C under 5% CO₂ and a humidified atmosphere. After carefully removing the medium, we added 600 µl of DMSO to each well, shook them gently and transferred the solution to 96-well microtiter plates. Absorbance at 595 nm was determined by spectrophotometry. Wells containing medium but no cells served as blanks, while wells with untreated cells were used as normal controls. The results were expressed as the mean of six replicates per condition.

Western blot analysis

The protein expression levels of proliferating cell nuclear antigen (PCNA) and Bcl2 in PC3 cells after coculture with PZsc or TZsc were investigated by Western blot analysis. Cells were plated $(1 \times 10^5$ cells per well) in the coculture model and treated as described above. After a 48-h treatment, PC3 cells were harvested, and total cellular protein was extracted from the cell pellets by homogenisation in RIPA buffer containing 10 mmol l⁻¹ Tris-HCl (pH 8.0), 150 mmol l⁻¹ NaCl, 0.1% sodium dodecyl sulphate, 0.8% Triton X-100 and a protease inhibitor cocktail (Roche, Mannheim, Germany) at 4 °C for 30 min. Protein samples (50 µg protein per sample) were subjected to electrophoretic analysis in 10%/4% sodium dodecyl sulphate-stacked polyacrylamide gels and were subsequently electrotransferred onto PVDF membranes (Bio-Rad, Hercules, CA, USA). Membranes were blocked for 1 h in Tris-buffered saline containing Tween 20 (10 mmol l⁻¹ Tris-HCl, pH 7.5, 100 mmol l^{-1} NaCl and 0.1% Tween 20) with 5% non-fat milk powder. Membranes were washed and probed at 4 °C overnight with either mouse anti-PCNA IgG (1:3000 dilution) or rabbit anti-Bcl2 IgG (1:3000 dilution). After washing, membranes were incubated for 1 h at room temperature with the horseradish peroxidase-linked



secondary antibody (1:5000 dilution). Immunoreactive bands were detected by the Immobilon Western Chemiluminescent HRP Substrate Kit (Millipore, Billerica, MA, USA). The intensity of each band was quantified by ImageJ software. All bands were normalized to beta-actin.

Quantitative real-time RT-PCR

PZsc/PC3 or TZsc/PC3 coculture models were constructed and treated as described above. After a 48-h treatment, total cellular RNA was extracted from PC3 cells using TRIzol (Invitrogen, Carlsbad, CA, USA) according to the manufacturer's recommended protocol. Two micrograms of total RNA were used to generate cDNA, and the real-time reaction was performed following the manufacturer's protocol (Promega, Madison, WI, USA). The reverse transcription product (1 µl) was used as a template in a 20-µl PCR reaction containing SYBR Premix Ex Taq PCR Mastermix (TaKaRa, DaLian, China). After an initial denaturation at 95 °C for 10 min, the samples were subjected to 45 cycles of a denaturation step at 95 °C for 30 s, an annealing step at 55-60 °C for 30 s and an extension step at 72 °C for 30 s. Primers were designed using the online tool Primer3 (http:// frodo.wi.mit.edu/primer3/); the sequences are shown in Table 1. As an internal control, cDNA for glyceraldehyde 3-phosphate dehydrogenase was amplified. Relative expression of the target genes was estimated using the method.²⁴

Statistical analysis

Unless otherwise stated, the results are expressed as the mean \pm s.d. Statistical analysis was conducted by analysis of variance or by paired *t*-test when only two values were to be compared. The data were analyzed by using SPSS software version 11.0 (SPSS Inc., Chicago, IL, USA). A *P* value of less than 0.05 was considered statistically significant.

RESULTS

Morphology of PZsc and TZsc

Figure 2 shows the morphology of PZsc and TZsc. We also analysed these stromal cells by IHC using vimentin and prostate-specific antigen as markers. PZsc and TZsc were all vimentin-positive (**Figure 2c and d**) and prostate-specific antigen-negative (**Figure 2e and f**). We also examined AR expression in PZsc and TZsc by IHC, but it was below the detection limit. These results suggest that the cells we cultured were stromal cells.

AR mRNA expression differs between PZsc and TZsc

We next measured *AR* mRNA expression levels in PZsc and TZsc (**Figure 3a**). *AR* was more highly expressed in PZsc than in TZsc in the presence of 10 nmol l^{-1} E2 plus 1–100 nmol l^{-1} DHT (*P*<0.05).

Table 1 Primers used for real-time RT-PCR

When the cells were treated with DHT plus E2, AR expression increased with increasing DHT concentrations in PZsc (P<0.05), but not in TZsc. Interestingly, when cells were treated with E2 (10 nmol l⁻¹) alone, TZsc expressed higher levels of AR than PZsc (P<0.05).

Differences in peptide growth factor secretion from PZsc and TZsc

We used ELISA to measure the concentrations of several important growth factors in the culture supernatants of PZsc and TZsc after the sex hormone treatments described above (Figure 3). The level of KGF in the supernatants increased with increasing concentrations of DHT (P < 0.01) and was higher in PZsc than TZsc (0 nmol l⁻¹ DHT, P < 0.01; 1 nmol l⁻¹ DHT, P < 0.05; 10 nmol l⁻¹ DHT, P < 0.01) except the treatment with 100 nmol l^{-1} DHT (**Figure 3b**). Interestingly, with increasing concentrations of DHT $(0-10 \text{ nmol } l^{-1})$, IGF-1 secretion increased in PZsc but decreased in TZsc (Figure 3c). In PZsc supernatants, IGF-1 levels were significantly higher than that in TZsc when the cells were treated with 10 nmol l⁻¹ DHT plus 10 nmol l^{-1} E2 (P<0.01), and decreased when DHT was applied at 100 nmol 1^{-1} . Additionally, we found that TGF- β 1 levels were significantly higher in the supernatants of PZsc and TZsc when the cells were treated with 10 nmol l^{-1} E2 alone than when they were treated with both E2 and DHT (P<0.01). TGF-β1 secretion was higher in TZsc supernatants than in PZsc supernatants when the cells were treated with 10 nmol l^{-1} E2 alone (*P*<0.01), but this pattern was reversed when cells received 10 nmol l^{-1} E2 plus 1 nmol l^{-1} DHT (P<0.01) or 10 nmol l^{-1} E2 plus 10 nmol l^{-1} DHT (*P*<0.01) (**Figure 3d**).

Different effects of PZsc and TZsc on the proliferation of PC3 cells under the action of sex hormones

After 168-h coculture, the proliferation of PC3 cells cocultured with PZsc was significantly higher than control groups (0 nmol l^{-1} DHT, P < 0.05; 1 nmol l^{-1} DHT, P < 0.01; 10 nmol l^{-1} DHT, P < 0.05; 1 nmol l^{-1} DHT, P < 0.01; 10 nmol l^{-1} DHT, P < 0.01; 100 nmol l^{-1} DHT, P < 0.01; 100 nmol l^{-1} DHT, P < 0.01; 10 nmol l^{-1} DHT plus 10 nmol l^{-1} E2 when the cells were cocultured with PZsc (P < 0.01) after 168-h coculture (Figure 4a). Indeed, the effect on proliferation increased as the DHT concentration was raised from 0–10 nmol l^{-1} , but decreased under the 100 nmol l^{-1} DHT treatment.

PCNA and Bcl2 protein expression

The protein expression levels of PCNA and Bcl2 in PC3 cells were measured by Western blot analysis. As shown in **Figure 5**, the PCNA level in PC3 cells cocultured with PZsc was higher than that of cells

Target gene	Forward primer (5' –3')	Reverse primer (5' –3')	Product size (bp)	Annealing temperature (°C)
AR	TGTCCATCTTGTCGTCTTC	CCTCTCCTTCCTCCTGTAG	103	55
KGF	CACCAGGCAGACAACAGACAT	GTAAGTTCAGTTGCTGTGACGCT	97	55
TGF-β1	CGACTACTACGCCAAGGAGG	CGGAGCTCTGATGTGTTGAA	120	55
EGF	GCCAGCTGCACAAATACAGA	AGTGGTGGTCATCTTCCCTG	121	57
IGF-1	CCTCCTCGCATCTCTTCTACCTGC	TGCTGGAGCCATACCCTGTG	165	55
bFGF	CCGTTACCTGGCTATGAAGG	ACTGCCCAGTTCGTTTCAGT	158	60
hTERT	CTGTAGAAGACGAGGCCCTG	GGCATAGCTGGAGTAGTCGC	128	57
GAPDH	GAAGGTGAAGGTCGGAGTC	GAAGATGGTGATGGGATTTC	226	55

Abbreviations: *AR*, androgen receptor; *bFGF*, basic fibroblast growth factor; *EGF*, epidermal growth factor; *GAPDH*, glyceraldehyde 3-phosphate dehydrogenase; *hTERT*, human telomerase reverse transcriptase; *IGF-1*, insulin-like growth factor 1; *KGF*, keratinocyte growth factor; *TGF-β1*, transforming growth factor-β1.





Figure 2 Morphology of PZsc (stromal cells derived from the PZ) and TZsc (stromal cells derived from TZ). Morphology and identification of primary human normal prostate stromal cells from the peripheral zone (PZ or the transition zone (TZ) by microscopy and immunocytochemistry (IHC). (a) PZsc, scale bar=200 μ m. (b) TZsc, scale bar=200 μ m. Vimentin expression was positive in PZsc (c) (scale bar=200 μ m) and TZsc (d) (scale bar=200 μ m), but prostate-specific antigen (PSA) expression was negative in PZsc (e) (scale bar=200 μ m) and TZsc (f) (scale bar=200 μ m).

cultured with TZsc (P < 0.01). PCNA expression increased further when PC3 cells cocultured with PZsc were treated with DHT plus E2. There were no significant differences in Bcl2 expression between PC3 cells cocultured with PZsc and PC3 cells cocultured with TZsc.

Relative mRNA expression by real-time RT-PCR

To evaluate the differential effects of PZsc and TZsc on PCa development as judged by growth factor expression patterns, we measured the mRNA expression levels of basic fibroblast growth factor (βFGF), KGF, epidermal growth factor (EGF), IGF-1, TGF- β 1 and human telomerase reverse transcriptase (hTERT) in PC3 cells after coculture with PZsc or TZsc. Treatment with 1 or 10 nmol l⁻¹ DHT and 10 nmol 1^{-1} E2 induced higher expression of *bFGF* in PZsc–PC3 coculture groups than in TZsc-PC3 coculture groups (Figure 6a, P<0.05). However, when the cells were treated with 10 nmol l^{-1} E2 alone, the PZsc-PC3 coculture groups expressed lower levels of bFGF than the TZsc-PC3 coculture groups (Figure 6a, P<0.01). Treatment with 1 or 10 nmol l⁻¹ DHT and 10 nmol l⁻¹ E2 led to higher expression of KGF and EGF in PZsc-PC3 coculture groups than in TZsc-PC3 coculture groups (Figure 6b and c). Under DHT stimulation (1, 10 or 100 nmol l⁻¹) and 10 nmol l⁻¹ E2, IGF-1 expression increased more in PZsc-PC3 coculture groups than in TZsc-PC3 coculture groups (Figure 6d). As shown in Figure 6e, $TGF-\beta 1$ expression was also higher in PZsc-PC3 coculture groups than TZsc-PC3 coculture groups treated with 10 nmol l^{-1} E2 and DHT (1 nmol l^{-1} , *P*<0.01; 10 nmol l^{-1} , *P*<0.05; 100 nmol l^{-1} , P<0.01). All the five growth factors were expressed at the lowest level among PZsc groups under 100 nmol l^{-1} DHT. We conclude that 10 nmol l^{-1} DHT plus 10 nmol l^{-1} E2 represents the optimal condition to increase growth factor expression in PZsc-PC3 coculture groups. However, all doses of DHT $(1, 10 \text{ and } 100 \text{ nmol } l^{-1})$ promoted the expression of hTERT in PZsc-PC3 coculture groups (Figure 6f, *P*<0.05).





Figure 3 Differences between PZsc and TZsc in *AR* mRNA expression levels and the secretion of peptide growth factors under the action of sex hormones. *AR* mRNA expression was determined by Q-PCR (**a**). Supernatant concentrations of KGF (**b**), IGF-1 (**c**) and TGF- β 1 (**d**) were detected by ELISA. The mean value of three independent measurements is shown. Cells were counted after supernatants were removed, and measurements were normalized for cell number. Results are given in pg ml⁻¹ per 1×10⁵ cells. **P*<0.05, ***P*<0.01. AR, androgen receptor; DHT, dihydrotestosterone; E2, β-oestradiol; EGF, epidermal growth factor; IGF-1, insulin-like growth factor 1; KGF, keratinocyte growth factor; PZsc, stromal cells derived from the peripheral zone; Q-PCR, quantitative real-time RT-PCR; TGF- β 1, transforming growth factor- β 1; TZsc, stromal cells derived from transition zone.





Figure 4 (a) The differential effects of PZsc and TZsc on PC3 cell proliferation under the action of sex hormones. As shown in **Figure 1**, PC3 cells were seeded on the insert chamber with a PET bottom (the upper chamber of the model). The stromal cells (PZsc or TZsc) were seeded on a regular plate (the lower chamber of the model). PZsc–PC3: PC3 cells cocultured with PZsc. TZsc–PC3: PC3 cells cocultured alone. (b) 0: treatment with 0 nmol I^{-1} DHT plus 10 nmol $^{-1}$ E2. (c) 1: treatment with 1 nmol I^{-1} DHT plus 10 nmol I^{-1} E2. (d) 10: treatment with 10 nmol I^{-1} DHT plus 10 nmol I^{-1} E2. (e) 100: treatment with 100 nmol I^{-1} DHT plus 10 nmol I^{-1} E2. (h the absence of DHT and E2 (ethanol) as a control treatment. **P*<0.05, ***P*<0.01. DHT, dihydrotestosterone; E2, β -oestradiol; PET, polyethylene terephthalate; PZsc, stromal cells derived from the peripheral zone; TZsc, stromal cells derived from transition zone.

DISCUSSION

Among prostate diseases, PCa mostly occurs in the PZ, whereas BPH occurs in the TZ. In elderly men, the incidences of PCa and BPH are increasing, but the pathogenesis of PCa and BPH is still poorly understood. Sex hormones and stromal–epithelial interactions are key factors involved in prostate growth, development and differentiation.^{25–28} In humans, the prostatic DHT level decreases with age, but the oestrogen level remains unchanged throughout life.^{29,30} Therefore, the ratio of E2 to DHT (E2/DHT) in the prostate increases with age. The period of development of PCa and BPH coincides with this elevation in the E2/DHT ratio, indicating that the balance between the two hormonal

factors may be critical to disease progression.³¹ In our study, we utilized direct stromal–epithelial coculture models to demonstrate the influence of primary prostate stromal cells on the properties of metastatic PC3 cells under an increasing E2/DHT ratio.

The growth-promoting effect of androgens on the epithelium during development is mediated through an AR-positive stroma.³² Blanchere *et al.*³³ reported that both androgen and oestrogen increased AR expression in cultured prostatic stromal cells. Our data show that androgen treatment promoted AR expression to a greater degree in PZsc than in TZsc, but that oestrogen had the reverse effect, stimulating more AR expression in TZsc than in PZsc. In PZsc, AR





Figure 5 Western blot analysis of PCNA and Bcl2 protein expression in PC3 cells. PZsc–PC3: PC3 cells cocultured with PZsc. TZsc–PC3: PC3 cells cocultured with TZsc. Actin was used as an internal control. **P<0.01, comparison between PZsc–PC3 and TZsc–PC3. DHT, dihydrotestosterone; E2, β -oestradiol; PCNA, proliferating cell nuclear antigen; PZsc, stromal cells derived from the peripheral zone; TZsc, stromal cells derived from transition zone.

expression increased with increasing androgen. This result indicates that PZsc might have a stronger response to androgen and that this response is dose-dependent. Conversely, oestrogen may play important roles in the TZ and in the development of BPH. Androgen action induces the production of growth factors from stroma that in turn can regulate the growth, development, and differentiation of the epithelium.³² Growth factors use paracrine pathways to signal to epithelial cells in the microenvironment. In our study, we detected three important growth factors (KGF, TGF-β1 and IGF-1) in secretions from prostate stromal cells (PZsc or TZsc). KGF, also called FGF-7, is positively regulated by androgens and stimulates the growth and proliferation of epithelial cells.^{34–39} We found that KGF secretion was higher in PZsc than TZsc, generally increasing as the E2/DHT ratio decreased. KGF might therefore be a factor involved in zonal differences. Androgen negatively regulates TGF-β1,⁴⁰ a finding with which

our results are consistent (Figure 2b). Additionally, we found that PZsc secreted more TGF-β1 than did TZsc under the action of androgen, but that this pattern reversed under the action of oestrogen alone. TGF-B1 is a pluralistic factor that can inhibit proliferation, induce apoptosis, promote angiogenesis and suppress immunity, all of which can enhance tumour development, progression and invasiveness.⁴¹⁻⁴⁶ Therefore, our findings indicate TGF-B1 as another possible factor involved in zonal differences. IGF-1 stimulates cell proliferation and inhibits apoptosis⁴⁷⁻⁵⁰ and has also been reported as a potent angiogenic factor.⁵¹ Stromal IGF-1 mediates tumour-stromal interactions in PCa to accelerate tumour growth.⁵² PCa epithelia are sensitive to surrounding IGF-1 levels, regardless of their androgen sensitivity status.¹⁸ Androgen treatment had opposing effects on IGF-1 secretion from the two stromal cell types, namely, it boosted IGF-1 secretion from PZsc but inhibited IGF-1 secretion from TZsc. The combination of 10 nmol l^{-1} E2 and 10 nmol l^{-1} DHT promoted IGF-1 secretion from PZsc to the greatest extent. We suggest that there might be different hormone/IGF-1 signalling pathways in PZsc versus TZsc and that IGF-1 might be a key factor in the development of prostate diseases. These differences between PZsc and TZsc indicate that growth factors differentially regulated by sex hormones may represent an important mechanism underlying zonal differences in prostate diseases.

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To further understand the effects of prostate stromal cells derived from different zones on epithelial cells and the role of sex hormones in this interaction, we developed a stromal-epithelial coculture model. We then used this model to compare the abilities of PZsc and TZsc to induce growth of the human PCa cell line PC3 in vitro. We found that PZsc have a greater capacity to promote PC3 cell proliferation than TZsc, especially in the presence of 10 nmol l^{-1} E2 plus 10 nmol l⁻¹ DHT. PZsc increased PC3 cell numbers, but TZsc had little effect. Moreover, PCNA was more highly expressed in PC3 cells cocultured with PZsc than in PC3 cells cultured with TZsc, although there was no significant difference in Bcl2 expression. These results indicate that the differential impacts of PZsc and TZsc manifested in epithelial proliferation rather than apoptosis. The enhancement of PC3 cell proliferation by sex hormones in the PZsc-PC3 coculture groups might be attributed to their modulation of a variety of growth factors. In addition, our data show that hTERT mRNA levels were higher in PC3 cells cocultured with PZsc compared with PC3 cells cocultured with TZsc, suggesting that PZsc promoted basal telomerase activity. Iczkowski et al.⁵ demonstrated that hTERT expression levels in human PCa tissue samples positively correlated with tumour grade and stage, indicating that hTERT may play a role in PCa progression. In this sense, our results suggest that PZsc may have a stronger ability to promote PCa progression. Several groups have reported that sex hormones can regulate hTERT expression,⁵⁴⁻⁵⁶ but the mechanism is unknown. We also found that E2 increased hTERT mRNA expression. Nanni et al.57 showed that oestrogen can increase the expression and activity of hTERT in normal human prostate epithelial cells. Indeed, E2 may be an initiating factor in PCa.⁵⁸ We also examined the expression of several important growth factors (bFGF, EGF, KGF, IGF-1 and TGF-β1) that act as vital stimulants for proliferation, immunosuppression, differentiation, invasiveness, angiogenesis and metastasis in PCa.⁵⁹ Following DHT plus E2 treatment, most of these growth factors were expressed at higher levels in PC3 cells cocultured with PZsc than in PC3 cells cocultured with TZsc. Furthermore, the highest levels were observed in the presence of 10 nmol l^{-1} E2 and 10 nmol l^{-1} DHT; expression decreased when DHT was increased to 100 nmol l⁻¹. The higher expression



а b PZsc-PC3 PZsc-PC3 Z TZsc-PC3 Relative expression of *bFGF* mRNA T o v v o o Z TZsc-PC3 Relative expression of KGF mRNA ** E PC3 40 D PC3 30 20 10 n - DHT (nmol I⁻¹⁾ Ω 1 10 100 0 DHT (nmol I⁻¹) 10 100 n 0 1 10 10 10 0 10 0 10 10 ÈE2 (nmol I⁻¹) E2 (nmol I-1) 10 10 с PZsc-PC3 ■ PZsc-PC3 d Relative expression of ^C HD EGF mRNA L1 0 5 01 51 Relative expression of IGF-1 mRNA Z TZsc-PC3 ** Z TZsc-PC3 30 T PC3 D PC3 25 20 15 10 5 0 DHT DHT n 1 10 100 n n 1 10 100 0 (nmol I⁻¹) (nmol I⁻¹) 10 10 10 10 0 10 10 10 10 0 E2 (nmol I⁻¹) E2 (nmol I⁻¹) PZsc-PC3 е 2 TZsc-PC3 PZsc-PC3 Relative expression of $TGF-\beta1$ mRNA Relative expression of hTERT mRNA ** D PC3 ** Z TZsc-PC3 2.5 4 2.0 PC3 3 1.5 2 1.0 0.5 1 0 0 DHT (nmol l⁻¹) DHT 0 100 0 10 100 0 1 10 0 1 (nmol l⁻¹) E2 (nmol l⁻¹) 10 10 10 0 10 10 10 0 10 10 (nmol I⁻¹)

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Figure 6 Relative mRNA expression levels in PC3 cells under different treatments (a-f). PZsc-PC3: PC3 cells cocultured with PZsc. TZsc-PC3: PC3 cells cocultured with TZsc. PC3: PC3 cells cultured alone. The absence of DHT and E2 (ethanol) is taken as a control treatment. The relative expression of the target genes was estimated using the method. The relative gene expression was calculated by assigning the reference value of 1 to the value for PC3 cells cultured alone and treated in the absence of DHT and E2. Each value represents the mean of three different experiments. *P<0.05, **P<0.01, comparision between PZsc-PC3 and TZsc-PC3. bFGF, basic fibroblast growth factor; DHT, dihydrotestosterone; E2, β-oestradiol; EGF, epidermal growth factor; hTERT, human telomerase reverse transcriptase; IGF-1, insulin-like growth factor 1; KGF, keratinocyte growth factor; PZsc, stromal cells derived from the peripheral zone; TGF- β 1, transforming growth factor- β 1; TZsc, stromal cells derived from transition zone.

of growth factors in PC3 cells cocultured with PZsc demonstrates that PZsc may have a greater capacity to induce PC3 cell development, differentiation, invasiveness and metastasis under the action of sex hormones. The mixture of 10 nmol l^{-1} E2 and 10 nmol l^{-1} DHT might be the best microenvironment to promote PC3 growth. Therefore, regulating the prostate microenvironment at different hormone levels may allow clinicians to effectively control the occurrence and development of PCa.

In conclusion, our data suggest that PZsc and TZsc represent different stromal microenvironments, as PZsc appear to more strongly induce PCa development, growth and progression than TZsc, likely via differential secretion of growth factors regulated by sex hormones. These findings offer a plausible explanation for the predominance of PCa development in the PZ. The present study provides several insights into the molecular mechanisms of zonal differences of prostate diseases, which may help illuminate novel therapeutic targets for PCa.

AUTHOR CONTRIBUTIONS

QJ designed and performed experiments, analysed data and wrote the paper; BMH designed the experiments and performed sample collection; FJZ performed cell culture and sample collection; YH performed cell culture and Western blot analysis; and SJX designed the experiments and wrote the paper.

COMPETING FINANCIAL INTERESTS

The authors declare no competing financial interests.

ACKNOWLEDGMENTS

This work was supported by a grant from the National Natural Science Foundation of China (No. 30672109).

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