

# Hypoxia of PC-3 prostate cancer cells enhances migration and vasculogenesis *in vitro* of bone marrow-derived endothelial progenitor cells by secretion of cytokines

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**Abstract.** Hypoxia is a key inducer of neovascularization which is essential for tumor growth, invasion and metastasis. It has been proposed that the recruitment of bone-marrow-derived endothelial progenitor cells (BM-EPCs) is pivotal and requires the participation of several tumor-derived cytokines. However, it is not known whether prostate cancer (PCa) cells contribute to the recruitment and vasculogenesis of EPCs in PCa progression. In the present study, we demonstrated that all conditioned medium (CM) of PC-3 PCa cells promoted proliferation and migration, and augmented the vasculogenesis capacity of BM-EPCs, and 24-h hypoxia (24H)-CM presented stronger ability compared to 24-h normoxia (24N)-CM and 48H-CM. Human cytokine antibody array with 174 anti-cytokine antibodies revealed the changes of cytokine in CMs. Twenty-five types of cytokines significantly increased in 24H-CM compared with 24N-CM. Eleven types of cytokines (5 factors increased and 6 decreased) were significantly different between 48H-CM and 48N-CM. Twelve types of cytokines (4 factors increased and 8 decreased) were signifi-

cantly different between 48H-CM and 24H-CM. Furthermore, according to the Gene Ontology analysis, all altered cytokines were involved in proliferation, chemotaxis, cell motility, cell migration, vasculogenesis and angiogenesis. Of note, the changed regularity of cytokines in the 24H-CM and 48H-CM of PC-3 cells was in concert with the functional changes of BM-EPCs treated by different CM of PC-3 cells in enhancing the proliferation, migration and vasculogenesis potential of BM-EPCs. These findings suggest that PCa cells may have the potential to modulate their microenvironment and facilitate BM-EPC migration and vasculogenesis by secretion of cytokines in the early stage of hypoxia.

## Introduction

Bone is the most common site of prostate cancer (PCa) metastasis and once tumors metastasize to bone, they are essentially incurable and result in significant morbidity prior to patient mortality (1,2). Therefore, to develop more effective therapeutics, a more thorough understanding of the key complex processes that are central to metastasis may reveal more robust and less redundant therapeutic targets.

Neovascularization is essential for tumor growth and it also facilitates tumor invasion and metastasis (3). Furthermore, in cancer patients with metastasis, neovascularization-mediated progression of micrometastasis to lethal macrometastasis is the major cause of mortality (4). Emerging evidence implicates hypoxia as a key inducer of neovascularization in tumors (5). Moreover, hypoxia-induced pathological neovascularization mediates tumor cell dissemination, invasion and metastasis in an animal model (6). However, the mechanisms underlying pathological neovascularization in relation to hypoxia in tumor invasion and metastasis remain elusive.

The process of hypoxia-induced neovascularization was formerly attributed to the migration and proliferation of pre-existing, fully differentiated endothelial cells (ECs), known as angiogenesis (7). However, it is now well established that tumors can acquire their vasculature by various mechanisms including postnatal vasculogenesis, a process during which circulating bone-marrow-derived endothelial progenitor cells (BM-EPCs) home to sites of neovascularization and diffe-

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**Abbreviations:** PCa, prostate cancer; Hx, hypoxia; Nx, normoxia; BM-EPC, bone marrow-derived endothelial progenitor cell; CM, conditioned medium; H-CM, hypoxic conditioned medium; N-CM, normoxic conditioned medium; NM, normal medium; 24N, 24-h normoxia; 24H, 24-h hypoxia; 48N, 48-h normoxia; 48H, 48-h hypoxia

**Key words:** bone marrow-derived endothelial progenitor cell, prostate cancer cell, hypoxia, cytokine, migration, vasculogenesis

rentiate into ECs (8). Although there has been controversy concerning the time during neovascularization at which the proposed contribution of EPCs to neovascular vessels might occur (9), it has been proposed that the recruitment of BM-EPCs is pivotal for tumor vasculogenesis (10-12). Notably, recent studies have shown that BM-EPCs are recruited to the angiogenic switch in tumor growth and metastatic progression (4,13,14) and blood levels of EPCs tend to increase in cancer patients and correlate with the stage of the malignant disease (15). In prostate cancer, although microvessel density is related to clinical stage, progression, metastasis and survival (16-18), it is not known whether EPCs contribute to the bone metastasis of PCa.

The mobilization, homing and incorporation of EPCs into tumors are multi-step and multi-factor events during tumor vasculogenesis. This complex process requires the participation of numerous cytokines. A series of studies have shown that several factors, such as VEGF (20), SDF-1 (20), G-CSF (21), GM-CSF (22), PIGF (23), IGF2 (24), E-selectin (25) and angiopoietin-2 (26), play an important role in inducing EPC homing and vasculogenesis. Although it is not very clear how circulating EPC home from the bone marrow specifically into the tumor microenvironment, the mobilization of EPC from the bone marrow into the bloodstream, migration to the tumor site, and subsequent integration into the vascular network, are all presumably controlled by tumor-derived factors (11). However, in PCa progression, it is not clear whether the above-mentioned tumor-derived factors are involved in the regulation of EPC migration and vasculogenesis in hypoxia, and whether other cytokines play roles in these processes. Since a recent study found that PCa cells which metastasize to bone competed with haematopoietic stem cells for the niche and colonize in the niche (27) and haematopoietic stem cell niches within the bone marrow are hypoxic (28,29), we speculated that PCa cells in the hypoxic microenvironment could recruit BM-EPCs and induce vasculogenesis by releasing hypoxia-induced cytokines.

In this *in vitro* study, we found that conditioned medium (CM) of PC-3 cells, derived from bone metastasis of PCa, by hypoxia for 24 h promoted proliferation and migration, and augmented the vasculogenesis capacity of BM-EPCs. Human cytokine antibody array revealed that under early hypoxia, the PC-3 cells secreted several cytokines involved in BM-EPC functions, including proliferation, migration and vasculogenesis. These findings suggest that PCa cells may have the potential to modulate their microenvironment and facilitate BM-EPC migration and vasculogenesis in the early stage of hypoxia.

## Materials and methods

**Cell culture and hypoxic treatment.** PC-3 cells were purchased from the American Type Culture Collection (ATCC, Manassas, VA, USA) and maintained in F-12 culture medium (HyClone) supplemented with 10% fetal bovine serum (HyClone). For normoxia experiments, the PC-3 cells were cultured at 37°C with 5% CO<sub>2</sub>/95% air in a humidified incubator, and for hypoxia experiments, the PC-3 cells were cultured at 37°C with 5% CO<sub>2</sub>, 94% N<sub>2</sub>, and 1% O<sub>2</sub> in a multigas incubator (Juji Field, Tokyo, Japan).

**Conditioned medium collection.** PC-3 cells were cultured in hypoxia and normoxia for 24 and 48 h and CM was collected. Briefly, PC-3 cells were seeded at a density of 2.5x10<sup>5</sup> cells/ml in F12 (HyClone) complete media into a 10-cm cell culture dish and grown to ~80% confluency. Cells were washed twice with phosphate-buffered saline (PBS) and incubated in serum-free medium for 24 and 48 h. CM was collected and centrifuged at 2,000 x g at 4°C for 10 min to remove cell debris and the supernatant was used.

**Isolation and cultivation of BM-EPCs.** Bone marrow was collected from male patients with lumbar degenerative diseases (age range, 27-72 years; mean age 52.4±13.9 years; disc herniation with lumbar spondylolisthesis, degenerative lumbar spondylolisthesis and lumbar spinal stenosis with instability) at the time of operation. Informed consent was obtained from the patients for bone marrow blood collection, and all procedures were performed in accordance with the guidance and approval of a research Ethics Committee at the First Affiliated Hospital of Sun Yat-sen University.

Mononuclear cells were collected by density gradient centrifugation using Ficoll-Paque™ Premium (1.077; GE Healthcare, San Francisco, CA, USA) according to the manufacturer's instructions (30). Briefly, the isolated cells were cultivated in dishes coated with fibronectin and induced by EGM-2 MV Single Quots (Cambrex) at 37°C with 5% CO<sub>2</sub>/95% air in a humidified incubator at a density of 3-5x10<sup>6</sup>/cm<sup>2</sup>. After 3 days in culture, non-adherent cells were removed by washing with new medium and changed. After 7 days in culture, BM-EPCs were identified by immunofluorescence staining and flow cytometry. Quantitative fluorescence-activated cell sorting (FACS) was performed on a FACSVantage SE flow cytometer (Becton-Dickinson, San Jose, CA, USA). The Weibel-Palade body in cultivation cells was visualized by transmission electron microscope (TEM).

**BM-EPC proliferation assay.** EPC proliferation assay was performed to assess the viability of BM-EPCs treated with CM of PC-3 cells. The effect of CM on BM-EPC proliferation was determined by 2-(2-methoxy-4-nitrophenyl)-3-(4-nitrophenyl)-5-(2,4-disulfophenyl)-2H-tetrazolium, monosodium salt (WST-8) assay kit (CKK-8, Dojindo, Tokyo, Japan). Briefly, BM-EPCs were plated in 96-well plates at a concentration of 5x10<sup>3</sup> cells/well in 100 µl CM of PC-3 cells. Following incubation for 36 h by CM, CKK-8 was used according to the manufacturer's instructions. WST-8 was added into each well for 4 h prior to the measurement. The absorbance at 450 nm was measured using a microplate reader.

**BM-EPC migration assay.** Cell migration was evaluated using the Transwell system (Corning Costar, Acton, MA, USA) with 6.5-mm diameter polycarbonate filters (8-µm pore size). Briefly, BM-EPCs were seeded onto chemotaxis filters in 100 µl F-12 medium. CM of PC-3 cells cultured under normoxia and hypoxia for 24 and 48 h was collected and then added to the lower chamber with a normal medium (NM; F-12) as control medium. After the 24-h migration period, non-migrating cells were completely removed from the top surface of the membrane and attached cells were fixed with 95% ethanol for 10 min and stained with 0.1% crystal violet for

25 min. The plate was immersed in fresh tap water to remove excess dye. The adherent cells were visualized under light microscopy at x200 magnification and quantified in triplicate by counting adherent cells in 5 randomly selected fields per well (Axiovert 100; Carl Zeiss Microimaging, Thornwood, NY, USA). The results are representative of 4 independent experiments.

**In vitro BM-EPC vasculogenesis assay.** *In vitro* EPC vasculogenesis assay was performed with In Vitro Angiogenesis Assay kit (Chemicon, Temecula, CA, USA) according to the manufacturer's instructions. Briefly, EC Matrix™ solution was thawed on ice overnight, mixed with 10X EC Matrix diluents and placed in a 96-well tissue culture plate at 37°C for 1 h to allow the matrix solution to solidify. BM-EPCs were plated in normoxic or hypoxic CM at a concentration of  $1 \times 10^4$  cells/well on the top of the solidified matrix solution (1% FBS was added to CM and NM). Tube formation was inspected under an inverted light microscope at x200 magnification. Tube formation was defined as a structure exhibiting a length four times its width. Five independent fields were assessed for each well, and the average number of tubes/x200 field was determined.

**Human cytokine antibody array.** Cytokine concentrations in CM were quantified by using human cytokine antibody arrays. RayBio Human Cytokine Antibody Array C Series 2000 kit was purchased from RayBiotech (Norcross, GA, USA). This array consisted of 174 different antibodies spotted in duplicate onto three membranes. CM was collected from normoxic or hypoxic PC-3 cells for 24 and 48 h with an NM (F-12) as control medium. Experiments were performed by KangChen Bio-tech (Shanghai, China) as recommended by the manufacturer. The signal intensities can be quantified by densitometry, and relative expression levels of cytokines can be acquired by comparing them. Positive control can be used to normalize the results from different membranes being compared.

**Gene Ontology analysis.** The Gene Ontology (GO) project provides a controlled vocabulary to describe gene and gene product attributes in any organism (<http://www.geneontology.org>). The ontology covers three domains: biological process, cellular component and molecular function. Fisher's exact test is used to find if there is more overlap between the DE list and the GO annotation list than would be expected by chance. The P-value denotes the significance of GO terms enrichment in the DE genes. The lower the P-value, the more significant the GO Term (P-value  $\leq 0.05$  is recommended). GOID, the ID of the Gene Ontology term used in Gene Ontology Project. Term, the name of the gene Ontology Term. Ontology, the ontology the GOID belongs to, i.e., 'biological process', 'cellular component' or 'molecular function'.

**Statistical analysis.** The experimental data are presented as the means of each condition  $\pm$  SEM. One-way ANOVA was performed for comparing more than two groups, and paired Student's t-test was performed when comparing two conditions (SPSS for Windows® version 16.0).  $P < 0.05$  (\* $P < 0.05$ , \*\* $P < 0.01$ ) was considered to indicate a statistically significant difference.

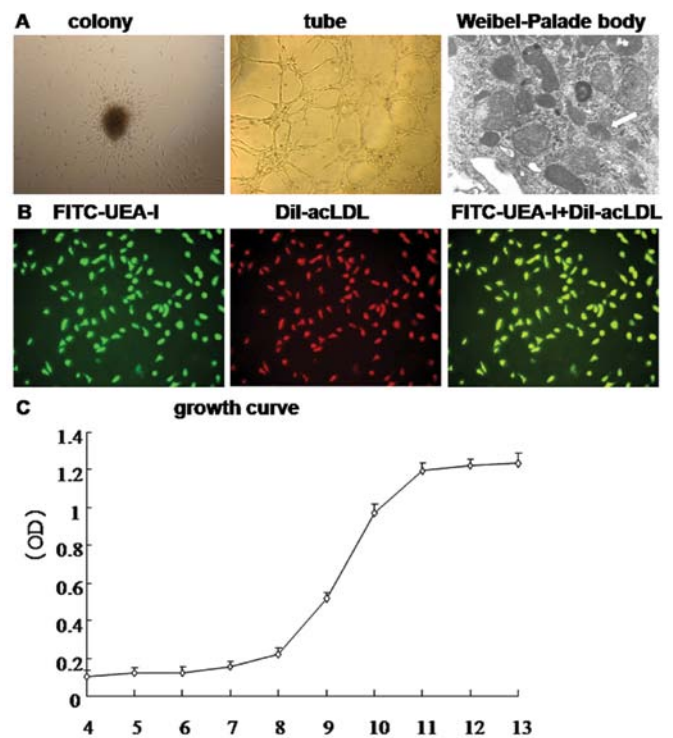


Figure 1. The growth processes of BM-EPCs. (A) The morphology of BM-EPCs in 7-day-cultured BM-EPCs formed colonies, tubes on the ECM *in vitro* and Weibel-Palade body (arrow) in the endochylema by TEM. (B) Immunofluorescence characterization of BM-EPCs. Binding to FITC-UEA-I, endocytosis of Dil-acLDL, and double positive cells appearing yellow in the overlay were identified as differentiating BM-EPCs. (C) The growth curve of BM-EPCs is shown.

## Results

**The growth processes of BM-EPCs.** Mononuclear cells were isolated from human bone marrow blood and cultivated in EGM2 on fibronectin-coated dishes. Cells demonstrated round morphology with different sizes as assessed by light microscopy during the first 3 days. With a tendency of ~20% of the mononuclear cells transferred to culture plates grew as adherent cells during 7 days of cultivation and clusters or colonies formed. Furthermore, in endochylema under TEM, the Weibel-Palade body was also observed and BM-EPCs had active angiogenic potential (Fig. 1A). Following cultivation for 7 days, the adherent cells exhibited strong ability to uptake Dil-acLDL, and FITC-UEA-I and the double positive rate was  $(94.3 \pm 4.1\%)$  (Fig. 1B), which were identified as differentiating EPCs. The growth curve is shown in Fig. 1C. The percentage of expression of CD133, CD34, KDR, VE-cadherin, E-selectin and vWF was determined by flow cytometry at Day 7 and Day 14 of culture. The expression pattern of surface markers in BM-EPCs changed toward a more mature endothelial cell phenotype from Day 7 to Day 14 (Table I).

**H-CM of PC-3 cells promotes proliferation of BM-EPCs.** The proliferation of EPCs, as a crucial step of vasculogenesis, is triggered and modulated by a variety of stimuli including hypoxic microenvironment. To demonstrate whether H-CM for 24 and 48 h promoted proliferation of BM-EPCs, the proliferation assay was performed to assess the viability of

Table I. The expression pattern of surface markers of BM-EPCs at Day 7 and Day 14.

| Time    | CD133     | CD34       | KDR       | VE-cadherin | E-selectin | vWF       |
|---------|-----------|------------|-----------|-------------|------------|-----------|
| Day 7   | 19.23±5.0 | 41.49±9.01 | 65.97±7.2 | 21.27±6.9   | 8.02±5.53  | 5.15±2.72 |
| Day 14  | 4.04±3.52 | 32.94±8.37 | 82.57±6.2 | 25.23±4.4   | 45.08±7.77 | 30.3±7.81 |
| P-value | 0.013     | 0.295      | 0.039     | 0.448       | 0.003      | 0.002     |

BM-EPCs treated with CM of tumor cells. WST-8 assay was used to examine the changes in cell proliferation. We exposed BM-EPCs to 24 and 48 H-CM, 24/48 N-CM and NM as a control medium for 36 h. All CM induced BM-EPC proliferation and 24H-CM promoted proliferation of BM-EPCs more evidently compared to 24N-CM (Fig. 2). However, 48N-CM promoted proliferation of BM-EPCs more apparently compared to 48H-CM. Furthermore, 24H-CM had stronger effects on enhancing BM-EPC proliferation than 48H-CM. These results suggested that hypoxia induced PC-3 cells to release some factors which promoted the proliferation of BM-EPCs.

*H-CM of PC-3 cells promotes migration of BM-EPCs.* The mobilization of EPC from the bone marrow into the bloodstream, migration to the tumor site, and subsequent integration into the vascular network are all presumably controlled by tumor-derived factors and those factors are released to the tumor microenvironment that is CM of tumor cells *in vitro*. To determine if CM collected from hypoxia-treated PC-3 cells (H-CM) for 24/48 h increased migration of BM-EPCs compared to CM collected from normoxia-treated PC-3 cells (N-CM), we used a Transwell assay to estimate the ability of EPCs to migrate. All CM enhanced BM-EPC migration compared with NM as a control (Fig. 3A and C). Also, 24H-CM enhanced migration of BM-EPCs more apparently than 24N-CM. However, 48N-CM increased migration of BM-EPCs more evidently than 48H-CM. Moreover, 24H-CM enhanced migration of BM-EPCs more apparently than 48H-CM. These observations suggested that hypoxia induced PC-3 cells to release some factors, which promoted the migration of BM-EPCs.

*H-CM of PC-3 cells augments in vitro vasculogenesis of BM-EPCs.* One of the most prominent effects of hypoxia is the induction of tumor neovascularization (5). To determine whether CM of PC-3 cells augmented the vasculogenesis of BM-EPCs *in vitro*, we exposed BM-EPCs to 24/48H-CM, 24/48N-CM and NM as a control medium for 18 h on Matrigel. All CM induced BM-EPC tube formation (Fig. 3B and D). The 24H-CM augmented the tube formation of BM-EPCs more evidently than 24N-CM. However, 48N-CM augmented the tube formation of BM-EPCs more apparently than 48H-CM. At the same time, 24H-CM had stronger effects on enhancing BM-EPC tube formation than 48H-CM. These observations indicated that hypoxia induced PC-3 cells to release some factors which augmented the *in vitro* vasculogenesis capacity of BM-EPCs.

*Significant alterations of the cytokines in H-CM.* In order to investigate tumor-derived cytokines of PC-3 cells which effect the functions of BM-EPCs, we used human cytokine antibody

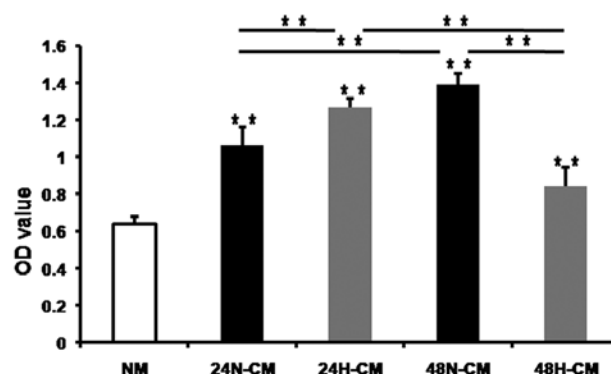


Figure 2. The effects of conditioned medium (CM) collected from hypoxia-treated PC-3 cells on the proliferation of BM-EPCs (\*\*P<0.01, respectively). All CM enhanced BM-EPC proliferation compared with normal medium (NM) as a control (\*\*P<0.01, respectively).

array to reveal the changes of cytokines in H-CM compared with N-CM. The cytokine profile for each of the CM was performed using three independent biological samples on protein arrays printed with 174 anti-cytokine antibodies in replicate. Relative expression levels of cytokines could be acquired by comparing the signal intensities. Significantly differentially expressed cytokines were selected as following standards: fold-change >1.3-fold and P<0.05 in H-CM compared with N-CM, and 24H-CM compared with 48H-CM. The results of cluster analysis are shown in Fig. 4. Compared with 24N-CM, most cytokines in 24H-CM were upregulated, but compared with 48N-CM, most cytokines in 48H-CM were downregulated. Compared with 24N-CM, most cytokines in 48N-CM were upregulated, however, compared with 24H-CM, most cytokines in 48H-CM were downregulated. These data indicated that in the early stage, hypoxia induced expression of most cytokines in PC-3 cells, but in the later stage, hypoxia repressed expression of several cytokines.

Twenty-five types of cytokines were significantly different between 24H-CM and 24N-CM, and all these factors increased in 24H-CM (Fig. 5A). Eleven types of cytokines were significantly different between 48H-CM and 48N-CM (Fig. 5B), and, among them, only 5 factors increased and 6 decreased in 48H-CM. Twelve types of cytokines were significantly different between 48H-CM and 24H-CM, and, among them, 4 factors increased and 8 decreased in 48H-CM (Fig. 5C). Forty-four types of cytokines were significantly different between 24N-CM and 48N-CM, and, among them, 43 factors increased in 48N-CM (Fig. 5D).

Some cytokines increased the expression in 24H-CM compared with that in 24N-CM, but these changes were not

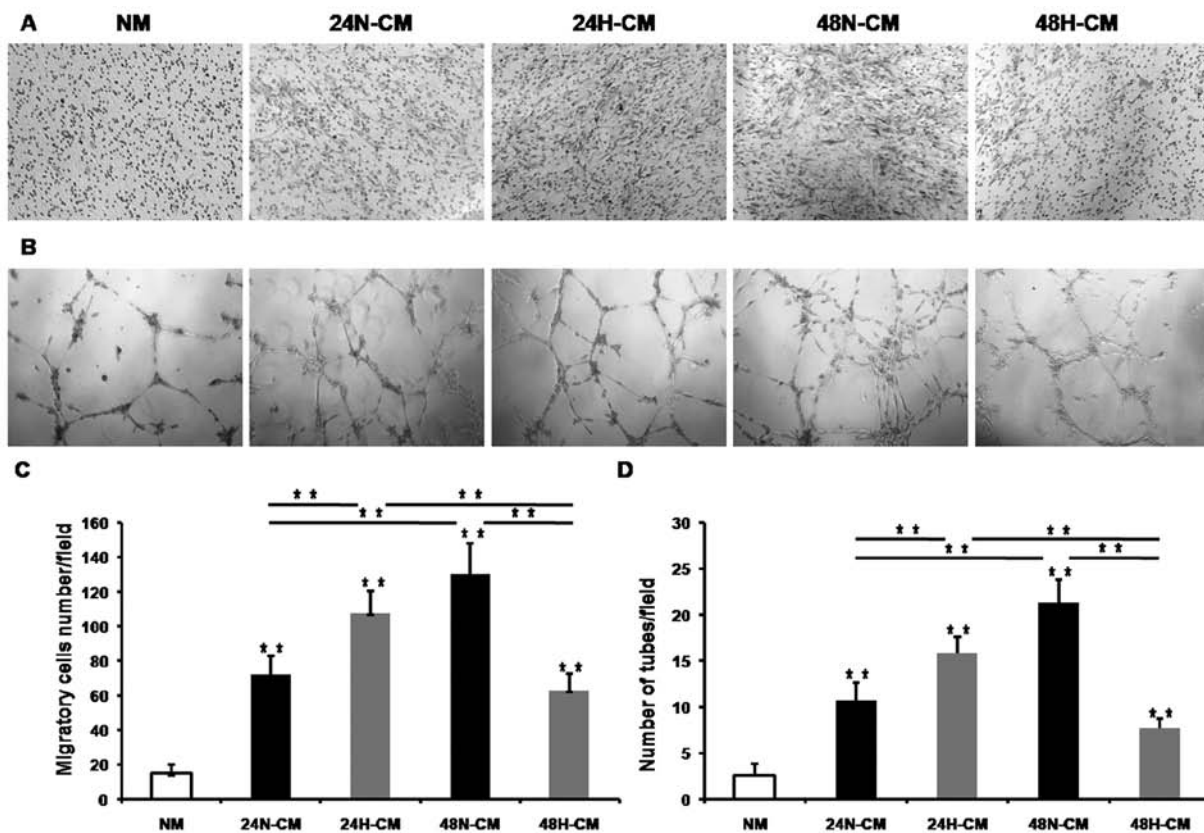


Figure 3. The effects of conditioned medium (CM) collected from hypoxia-treated PC-3 cells on migration and *in vitro* vasculogenesis of BM-EPCs. All CM enhanced migration and *in vitro* vasculogenesis of BM-EPCs compared with normal medium (NM) as a control (\*\* $P < 0.01$ , respectively). (A and C) Migration capacity of BM-EPCs treated with CM of PC-3 cells (\*\* $P < 0.01$ , respectively). (B and D) *In vitro* vasculogenesis capacity of BM-EPCs treated with CM of PC-3 cells (\*\* $P < 0.01$ , respectively).

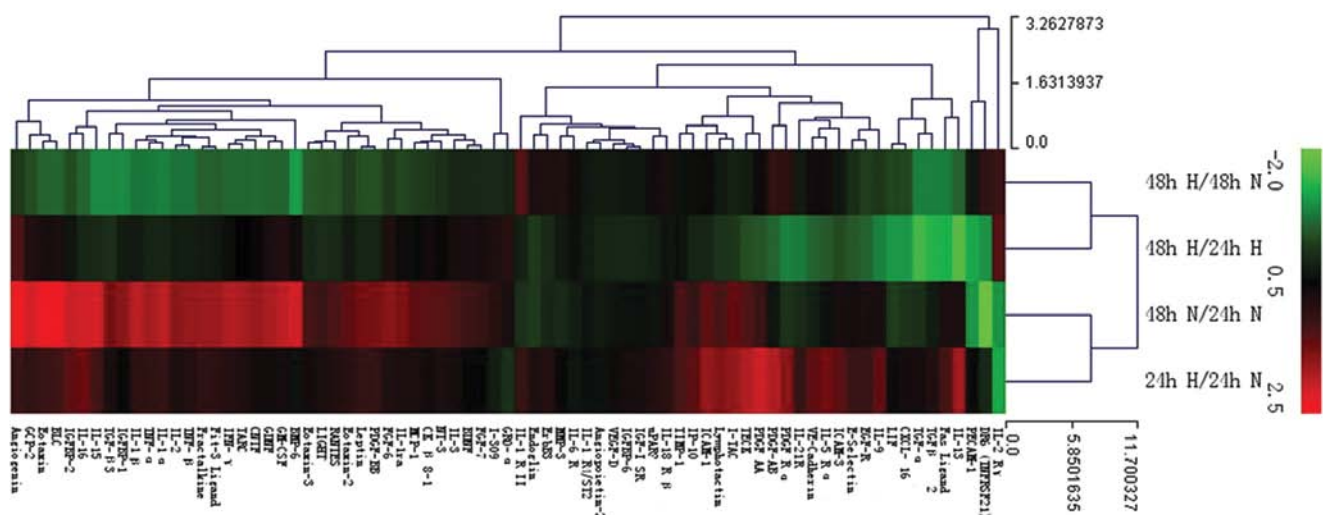


Figure 4. Cluster analyses of all significantly differentially expressed cytokines in hypoxic (H) and normoxic (N) conditioned medium (CM) (fold-change  $> 1.3$ ,  $P < 0.05$ ).

statistically significant, which play an important role in inducing EPCs homing and vasculogenesis according to the results in literature (20,23,24), including VEGF (fold-change=2.47,  $P=0.083$ ), SDF-1 (fold-change=1.43,  $P=0.429$ ), IGF-II (fold-change=1.84,  $P=0.053$ ), PIGF (fold-change=1.79,  $P=0.125$ ).

*Functions of the cytokines with significant alterations in H-CM are involved in proliferation, migration and vasculogenesis.* Differentially expressed cytokines selected as standards were analyzed by GO analysis. The GO project provides a controlled vocabulary to describe gene and gene

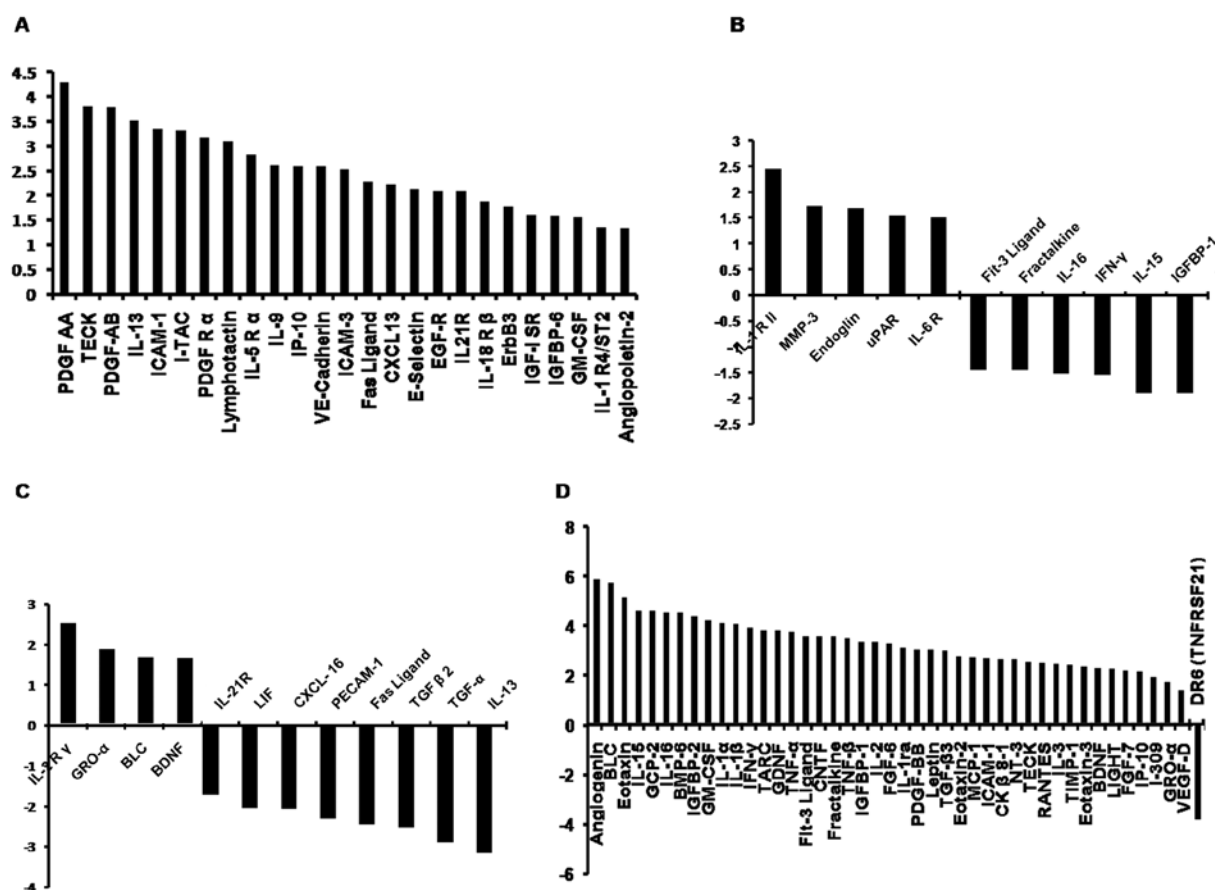




Table II. Gene Ontology analysis results of differentially expressed cytokines in conditioned media which related with proliferation, migration and angiogenesis.

|   | 24H/24N   | 48H/48N  | 48H/24H  | 48N/24N  |
|---|---|--|--|--|
| Positive regulation of cell proliferation (GO:0008284)    | Upregulated<br>CSF2, CXCL10, EGFR, FASLG, IGF1, IL9, IL13, PDGFA, PDGFB, PDGFRA | Upregulated<br>IL6R<br>Downregulated<br>IFNG, IL15               | Downregulated<br>FASLG, IL13, LIF, TGFA, TGFB2                 | Upregulated<br>ANG, CCL2, CCL5, CCL11, CCL24, CCL26, CNTF, CSF2, CXCL10, FGF7, FGF6, FIGF, IFNG, IL1B, IL2, IL3, IL15, PDGFB, TIMP1, TNF                   |
| Negative regulation of cell proliferation (GO:0008285)    |   | Upregulated<br>ENG<br>Downregulated<br>FNG, IL15                 | Upregulated<br>CXCL1<br>Downregulated<br>LIF, TGFB2            | Upregulated<br>ANG, CCL23, CXCL1, IFNG, IL1A, IL1B, IL2, IL15, LTA, TGFB3, TNF   |
| Chemotaxis (GO:0006935)                                   | Upregulated<br>ANGPT2, CCL25, CXCL10, CXCL11, CXCL13, EGFR, PDGFA, PDGFB, XCL1  | Upregulated<br>ENG, IL6R, PLAUR<br>Downregulated<br>IL16, CX3CL1 | Upregulated<br>CXCL1, CXCL13<br>Downregulated<br>CXCL16, TGFB2 | Upregulated<br>CCL1, CCL2, CCL5, CCL11, CCL17, CCL23, CCL24, CCL25, CCL26, CXCL1, CXCL6, CXCL10, CXCL13, CX3CL1, FGF7, FIGF, GDNF, IL1B, IL16, NTF3, PDGFB |
| Cell migration (GO:0016477)<br>Cell motility (GO:0048870) | Upregulated<br>ANGPT2, CCL25, EGFR, ICAM1, IGF1, PDGFA, PDGFB, PDGFRA, SELE     | Upregulated<br>ENG, IL6R<br>Downregulated<br>CX3CL1, IL16        | Downregulated<br>CXCL16, ECAM1, TGFB2                          | Upregulated<br>ANG, CCL2, CCL5, CCL11, CCL24, CCL25, CCL26, CX3CL1, FGF7, FIGF, GDNF, ICAM1, IL1B, IL16, PDGFB, TNF  |
| Vasculogenesis (GO:0001570)                               |   | Upregulated<br>ENG   |  |  |
| Positive regulation of angiogenesis (GO:0045766)          |   | Downregulated<br>CX3CL1  |  | Upregulated<br>IL1A, IL1B, CCL11, CCL24, CX3CL1  |
| Negative regulation of angiogenesis (GO:0016525)          |   |  | Downregulated<br>FASLG, LIF                                    |  |
| Angiogenesis (GO:0001525)                                 |   | Upregulated<br>ENG<br>Downregulated<br>CX3CL1                    | Downregulated<br>FASLG, LIF, TGFB2                             | Upregulated<br>ANG, CCL2, CCL11, CCL24, CX3CL1, FGF6, FIGF, IL1A, IL1B   |

preceding vessel formation, followed by differentiation into endothelial cells and luminal incorporation into a subset of sprouting tumor neovessels (12). These findings suggested that BM-EPCs might contribute to the development of micrometastasis of PCa in bone marrow in the early stage of colonization of metastatic PCa cells.

The migration of EPCs induced by cytokines released from tumor cells is one of the key mechanisms in the process of vascular development of hypoxia-induced vasculogenesis and one of the rate-limiting steps in the homing of EPCs (10-12). Several studies have demonstrated that factors such as VEGF (20), SDF-1 (20), G-CSF (21), PIGF (23) and IGF2 (24) are pivotal in the regulation of homing of EPCs. In the present study, the results showed that hypoxia did not induce statistically significant high expressions of the above-mentioned cytokines in PC-3 cells.

However, the expression levels of GM-CSF, angiopoietin-2, PDGF-AB, and E-selectin were induced significantly by 24-h hypoxia and 2 types of cytokines, endoglin (ENG) and IL6, in 48-h hypoxia. Previous studies have demonstrated that GM-CSF (22), E-selectin (25), angiopoietin-2 (26), PDGF-AB (33), ENG (34) and IL6 (5) are the important factors for the hypoxia-induced homing of BM-EPCs in the vasculogenesis process. Therefore, we speculated that these cytokines might play a more important role in the hypoxia-induced homing of EPCs in PCa bone metastasis compared to VEGF, SDF-1, G-CSF, PIGF and IGF2. We also found that the expression levels of several cytokines significantly upregulated, including CXCL13 (BLC), CXCL10 (IP-10), CCL25 (TECK), CXCL11 (I-TAC), lymphotactin (XCL1), ICAM1, EGFR, IGF1R in CM of PC-3 cells by 24-h hypoxia. These cytokines are involved in chemotaxis, cell motility and cell

migration according to the gene ontology analysis. Some of them have also been proved to be involved in EPC homing, such as ICAM1 (36,37), IGF1R (38). Those cytokines may be other factors that recruit EPCs to the tumor. Therefore, EPC homing might be a combined effect of several cytokines in the vasculogenesis process of bone metastasis of PCa.

EPCs that are incorporated into the tumor vasculature could constitute as much as 50% (39), but it is not known if the proliferating capabilities of EPCs are controlled by the tumor microenvironment factors. In the present study, the results showed that CM of PC-3 cells by 24-h hypoxia promoted EPC proliferation and hypoxia significantly induced PC-3 cells releasing several cytokines of positive regulation of cell proliferation. According to the Gene Ontology analysis, those cytokines included GM-CSF, IL5, IL13, EGFR, IGF1R, IL9, CXCL10 in 24-h hypoxia, and IFNG, IL15, IL6 in 48-h hypoxia, among them, GM-CSF (40) was proved to have the function of promoting proliferation of EPCs. Some of those cytokines have also been reported to be involved in EPC homing, such as GM-CSF (22), IL6 (35) and IGF1R (38). Therefore, the proliferating capabilities of EPCs may be controlled by the tumor microenvironment.

Our results showed that 24H-CM of PC-3 cells augmented the *in vitro* vasculogenesis capacity of BM-EPCs and hypoxia significantly induced PC-3 cells releasing angiopoietin-2. Since angiopoietin-2 was proved to promote vasculogenesis of EPC (41), it was suggested that angiopoietin-2 might play an important role in the vasculogenesis of BM-EPCs in PCa progression. The results also showed that ENG and IL6 significantly increased in 48H-CM of PC-3 cells and ENG (34) and IL6 (35) have been proved to be involved in vasculogenesis. It was suggested that ENG and IL6 might play a role in vasculogenesis in 48-h hypoxia.

In conclusion, our study detected 174 cytokines in CM of PC-3 cells from bone metastasis of PCa and found that 24-h hypoxia significantly induced the secretion of several cytokines. Those released cytokines in the CM of PC-3 cells by 24-h hypoxia promoted proliferation and migration and augmented vasculogenesis *in vitro* of BM-EPCs. Our data suggested that PCa cells may have the potential to modulate their microenvironment and facilitate BM-EPC migration and vasculogenesis in the early stage of hypoxia and BM-EPCs might contribute to the development of micrometastasis of PCa in bone marrow in the early stage of colonization of metastatic PCa cells. However, aside from the cytokines found in this study, other cytokines may be involved in these processes. Furthermore, it also is necessary to detect the functions of these cytokines in future studies. The characterization of tumor-associated BM-EPCs may provide valuable insight for more specific anti-angiogenesis therapy and/or tumor diagnosis.

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