Research Paper

Effects of Hypoxia, Surrounding Fibroblasts, and p16 Expression on Breast Cancer Cell Migration and Invasion

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Abstract

Cancer cell migration and invasion play essential roles in the metastatic cascade that transforms the local, noninvasive confined tumor cells to the motile, metastatic cancer cells moving through the extracellular matrix and basement into the circulation. Accumulated evidences suggest that intratumoral hypoxia, a characteristic of fast-growing solid tumors, promotes cancer cell motile and invasive abilities. In this study, we investigated the effects of hypoxia, surrounding fibroblasts, and p16 expression on the migration and invasion of breast cancer cells. We found that hypoxia promoted breast cancer cell migration and invasion, and cocultured fibroblasts stimulated invasiveness of breast cancer cells. Moreover, by using a Tet-on inducible system, we found that p16 is capable of inhibiting hypoxia-induced cell migration and invasion of breast cancer cells, and suppressing cocultured fibroblast-stimulated invasiveness of breast cancer cells. These results suggest that p16, in addition to its well-known anti-tumor proliferation function, has novel anti-cancer properties capable of suppressing hypoxia-mediated cancer cell migration and invasion. This study may provide important validation for p16-mediated cancer therapy either by gene therapy or pharmacological activation of internal p16 gene that is usually inactive due to hypermethylation in the tumor cells.

Key words: breast cancer, fibroblasts, hypoxia, invasion, migration, p16.

Introduction

One key characteristic of fast-growing solid tumors is the development of intratumoral hypoxia, which promotes malignant tumor progression. Cancer cell migration and invasion play important roles in the metastatic cascade. Cell migration is the transition process from the local, noninvasive confined tumor cells to the motile, metastatic cancer cells when the cells obtain the ability to dissociate from intracellular adhesions and become motile (1,2). Invasion, on the other hand, is the process by which malignant cells move through the basement membrane and gain access to blood vessels and lymphatic channels. As cancer cell migration and invasion are essential prerequisites for tumor metastasis, identifying agents that effectively block hypoxia-induced cancer migration and invasion would promote development of effective anti-cancer therapeutic approaches.

Hypoxia-inducible factor-1 (HIF-1), a master transcriptional factor in response to hypoxia, activates a group of downstream genes that promote tumor malignant progression. HIF-1 is composed of an inducible subunit, HIF-1α and a constitutively ex-
pressed subunit, HIF-1α (3). Heterodimerization of HIF-1α and HIF-1β is required to form the transcriptionally active HIF-1 that binds to the hypoxic response elements within the promoter regions of target genes (4). HIF-1α degrades quickly under normal oxygen tension (5) but it is inducible and stabilized under low oxygen tension (hypoxia) (6). Increased HIF-1α levels have been shown to correlate with decreased patient survival in many cancers including breast cancer (BCa) (7). The activation of HIF-1α stimulates a group of HIF-1α-regulated genes including vascular endothelial growth factor (VEGF) (3,8). As a result, the cells are converted towards malignant progression. HIF-1 activity is mainly dependent on the level of HIF-1α protein, the inducible and regulatory subunit of the HIF-1 dimer complex (9, 10).

We have previously shown that p16, a tumor suppressor gene and cyclin D kinase inhibitor and a negative cell cycle regulator (11), can neutralize the transactivation ability of HIF-1α on its target gene VEGF (12). In this study, we evaluated whether hypoxia stimulates breast cancer migration and invasion, and whether ectopic expression of p16 is capable of modulating hypoxia-mediated cell migration and invasion. We also evaluated whether cocultured stromal cells (fibroblasts) promotes BCa cell invasion and whether p16 inhibits that effect, as well as whether HIF-1α of fibroblasts specifically has a role in promoting cocultured BCa cell invasiveness.

Materials and Methods

Cell lines, cell culture conditions, and reagents

Human breast cancer (BCa) cell line MDA-MB-231 was obtained from American Type Culture Collection (ATCC, Manassas, VA). Human breast cancer cell lines LM2, a MDA-MB-231 derivative line that has high lung metastatic ability (15), was a generous gift from Dr. T Seagroves Azuma of Osaka Medical College, Osaka, Japan (13). Mouse MEF fibroblasts expressing wild-type (WT) and knockout (KO) HIF-1α were from Dr. T Seagroves of University of Tennessee Health Science Center (TN).

MDA-MB-231 cells were grown in RPMI-1640 (Gibco, Life technologies Corporation, Grand Island, NY) with 10% fetal bovine serum (FBS) (Gibco), LM2 cells were grown in Dulbecco’s Modified Eagle medium (DMEM) (Cellgro, Mediatech, Inc, Manassas, VA) with 10% FBS. JygMC(A) cells were grown in DMEM (Cellgro) with 10% FBS. Both MEF lines were grown in DMEM medium with 25 mM HEPES (Gibco) with 10% FBS. All cell lines were grown in medium containing 100 units/ml penicillin, and 100 µg/ml streptomycin. The cell cultures were incubated at 37°C either under normoxia (5% CO2, 21% O2) or hypoxia (5% CO2, 1% O2, balanced with N2) conditions.

Generation and screen of stably transected, Dox inducible (Tet-on) p16-expressing breast cancer cell lines

The construction of the Tet-on p16 lentiviral system Lenti-Tet-on p16 (pLenti-Tet-p16-pgkpuro and pcFUW-rtTA3-ires-puro) was described previously (12). Breast cancer cells (MDA-MB-231, LM2, and JygMC(A), respectively) were co-transduced by pLenti-Tet-p16-pgkpuro and pcFUW-rtTA3-ires-puro at multiplicity of infection (moi) of 10 each, together with 6 µg/ml polybrene (Millipore, Bedford, MA). The stably transected cells were enriched in medium containing 1 µg/ml puromycin (Clontech, Mountain View, CA). The resultant stably transected cells are named as MDA/Tet-on-p16, JygMC(A)/Tet-on-p16, and LM2/Tet-on-p16, respectively. For induction of p16 transgene expression in Tet-on-p16 stably transected cells, 1 µg/ml doxycyline (Dox) (Clontech) was used in the medium for incubation.

Immunohistochemistry

The procedure followed the method as described previously (12). Briefly, for immunohistochemical (IHC) staining, cultured cells were grown on SlideFlasks with bottom detachable slides (Nalge Nunc, Naperville, IL) that could be used for IHC staining directly later. The samples (slides) were first incubated with 1% H2O2 for 30 min. The samples were then incubated with first antibody against human p16 (Santa Cruz Biotechnology, Inc., Santa Cruz, CA) for 16 h at 4°C, then by corresponding second antibody and the Universal Elite ABC Kit (Vector Laboratories, Inc., Burlingame, CA) according to the manufacturer’s protocol. The reaction was visualized with DAB solution (75 mg 3,3’-Diaminobenzidine and 30 ml 50% H2O2 in 150 ml PBS) for 3-10 min.

Cell migration assay

The breast cancer MDA-MB-231 cells stably transfected with inducible Tet-on p16 (MDA/Tet-on p16) were incubated in the absence or presence of 1 µg/ml Dox for 72 h. The cells were then harvested and used for the following migration assay. The cell migration was measured by a modified Boyden’s chamber method using BD Falcon Cell Culture Inserts incorporating polyethylene terephthalate (PET) track-etched membrane with a pore size of 8.0 µm (BD)
The BCa cells were examined by the invasion assay under normoxia by using commercially available invasion inserts. For each Dox inducible BCa/Tet-on p16 cell line, the cells were either incubated in medium with or without 1 μg/ml Dox for 72 h. The cells were then harvested and used for the following *in vitro* invasion assay in the 6-well-plate BD BioCoat Matrigel Invasion Chambers (BD Biosciences Bio-Coat Matrigel Invasion Chamber) according to the manufacturer’s procedure. Briefly, the chamber was first rehydrated with serum-free medium (SFM) for 2 hr at 37°C. After rehydration, the chambers were placed in the lower compartment loaded with medium containing 5% FBS. Meanwhile, the above-mentioned cells were suspended and adjusted to 1.25×10^5 cells/ml in SFM with or without 1 μg/ml Dox. The cell suspension (2 ml or 2.5×10^5 cells per well) was immediately added to the upper compartment of the chamber. The cells were then allowed to invade through the matrigel for 22 h at 37°C, and the noninvading cells were removed by scrubbing the upper surface with a wet cotton swab. The filters were stained with Diff-Quick stain kit (Dade Behring Inc., Newark, DE), drained and counted under the microscope.

### In vitro invasion assay with self-coated matrigel

The BCa LM2 cells, which express a green fluorescent protein (GFP) gene (15), were transfected with Lenti-Tet-on p16 to generate LM2/Tet-on-p16 line. LM2/Tet-on-p16 cells were incubated in the absence or presence of 1 μg/ml Dox for 72 h before the experiment, and the samples with Dox induction were maintained throughout the following assay. All the cells (for simplicity, we used LM2 here referring LM2 or LM2/Tet-on-p16, and MEF referring MEF/HIF-1α WT or MEF/HIF-1α KO) were grown in SFM overnight and then harvested for the following invasion assay. The 24-well size Falcon Cell Culture Inserts (Corning Incorporated, Corning, NY) were precoated on the upper-surface (inside the inserts) with 100 μl matrigel (BD Matrigel, Bedford, MA) per insert and let them solidify at 37°C overnight. For experiment with BCa cells alone, cell suspension of 5×10^5 LM2 cells in 100 μl SFM was seeded into the upper chamber of the insert; for experiment with coculture of BCa and fibroblasts, cell suspension of 5×10^5 LM2 cells and 5×10^5 MEF cells in 100 μl SFM was seeded into the upper chamber of the insert. For the lower chamber of a 24-well culture plate, 650 μl SFM with 5% BSA (alternatively, with 5% FBS) was placed. The cells were then incubated at either normoxic or hypoxic conditions (see above) at 37°C for 18 h. The inserts were then removed, the matrigel (containing the unpenetrated cells) in the upper chamber of the inserts was carefully wiped off by using a wet cotton-tipped applicator. The resultant inserts were placed on a clean 24-well plate, and the matrigel-penetrated/invaded BCa LM2 cells on the membrane of the inserts were measured by using the SpectraMax M2 microplate reader (Molecular Devices, Sunnyvale, CA) for counting the GFP fluorescence intensity (RFU). Because LM2 line expresses GFP, this approach would only count the invaded BCa LM2 (GFP-expressing) cells.

### Results

#### Hypoxia stimulates BCa cell migration and p16 inhibits this hypoxia-induced migration

Cell migration is an important aspect of the tumor metastatic process. HIF-1α was implied, by others’ studies, in stimulation of cancer cell migration (16,17) and hypoxia induces HIF-1α (14). Thus, we were interested in analyzing whether hypoxia has effect on breast cancer cell migration and whether p16, a biological agent that appears to neutralize HIF-1α transcriptional activity (12), has ability to suppress hypoxia-mediated cancer cell migration.

To accurately examine the effects of p16 on BCa cell migration, we designed an inducible p16 expres-
sion so that p16 expression and effects can be studied in the same cells with or without inducer. A lentivirus expressing inducible human wild-type (WT) p16 under the control of a Tet-on promoter (Lenti-Tet-on p16) was constructed and stably transfected into MDA-MB-231 cells. The resultant stably infected line (MDA/Tet-on-p16) demonstrated a tightly regulated and inducible p16 expression under inducer Dox treatment (+Dox, top panel) (Fig. 1). Importantly, MDA-MB-231 cells, like other BCa cell lines (including JygMC(A) and LM2) we have examined, do not express endogenous p16 protein in the absence of Dox (-Dox) induction (Fig. 1, top panel). In other words, MDA/Tet-on-p16 cells in the absence of Dox behave just as control parental MDA-MB-231 cells.

MDA/Tet-on-p16 line was subsequently used to detect potential effects that p16 had on cell migration. A modified Boyden’s chamber assay using transwell precoated with fibronectin (a component of extracellular matrix) onto the insert undersurface of the membrane was used as described in the M & M for the migration measurement. As shown in Fig. 1 (bottom panel), MDA/Tet-on p16 cells under hypoxia had more than two-fold (>100%) increase in cell migration than that of normoxia (compare first two columns on the left). Importantly, inducible p16 expression by Dox significantly inhibited this hypoxia-induced cell migration (column 4) with a 36.0% inhibition in comparison to cells under hypoxia without p16 expression (column 2), indicating that p16 effectively inhibits hypoxia-induced migration in MDA-MB-231 cells. In contrast, p16’s effect on cell migration under normoxia is marginal. It should be noted that the inhibition of migration is not due to p16-mediated effect on cell proliferation, as the migration assay was conducted within 3 h and the MDA-MB-231 cell line’s doubling time is 41.9 h. In addition, Dox treatment per se did not have any effect on migration as evidenced by Dox treatment on BCa cells stably transfected with inducible GFP (not shown). These combined results (Fig. 1) strongly demonstrate that hypoxia stimulates BCa cell migration and p16 significantly inhibits this hypoxia-induced migration.

**p16 inhibits invasiveness ability of BCa cells**

Invasion is the process by which malignant cells move through the basement membrane and gain access to blood vessels and lymphatic channels. The classic and well-established *in vitro* invasion assay aims to measure cancer cells’ ability to penetrate the matrigel (a resemble mixture of basement membrane matrix).

p16’s effect on cancer cell invasiveness has never been studied before. To investigate the effects of p16 on BCa invasive ability, we used the *in vitro* invasion assay to analyze three BCa lines’ abilities to penetrate matrigel, in the presence or absence of p16 expression, respectively. Briefly, BCa cells stably transfected with inducible Tet-on p16 were incubated with or without 1 µg/ml Dox for 3 days, and then the treated BCa cells were placed on the matrigel invasion chamber (BD Biosciences Bio-Coat Matrigel Invasion Chamber). After 22 hours incubation at 37°C, the noninvasive cells were carefully removed from the upper surface of the membrane, and the penetrated cells (i.e., the invasive cells) going through matrigel were counted under the microscope. We found that all three BCa cell lines had significantly reduced invasiveness in the presence of p16 (Dox induced), at a decrease of 85.8% (MDA-MB-231), 82.7% (JygMC(A)), and 80.4% (LM2), compared to their corresponding untreated (p16 non-expressing) counterparts, respectively (Fig. 2).

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Figure 1. Inducible p16 expression in MDA-MB-231 cells and p16’s effect on hypoxia-induced migration. (Top panel) IHC staining for p16 protein in MDA-MB-231 cells stably expressing Tet-on inducible p16 (MDA/Tet-on-p16): MDA/Tet-on-p16 cells were treated without or with 1 µg/ml Dox for 72 h. The cells were then IHC stained by primary anti-p16 antibody. The dark brown color indicates p16 protein (+Dox). (Bottom panel) p16 inhibits hypoxia-induced cell migration: MDA/Tet-on-p16 cells were incubated with or without 1 µg/ml Dox for 72 h, followed by cell migration assay as described in M & M section under normoxic and hypoxic conditions. The results represent the data from at least two independent experiments, each performed in triplicate.
Effects of surrounding fibroblasts, and p16’s neutralization, on cocultured BCa cell invasiveness

Emerging evidence implies that tumor-surrounding areas contribute to tumor growth and progression; however, the key attributes from tumor-surrounding cells that clearly display promotion for tumor progression as well as the relevant mechanisms of action remain unknown. To investigate effects of tumor-surrounding cells such as fibroblasts on BCa progression and invasion, we used the in vitro invasion assay to analyze effects of MEF fibroblasts on cocultured BCa LM2 cells’ ability to penetrate the matrigel. After 18 hours incubation at 37°C of the LM2 alone or cocultured LM2 and MEF on the upper chamber of the insert that were precoated with matrigel (see details of M & M section), the matrigel (containing the noninvasive cells) was carefully removed from the upper surface of the membrane, the invaded cells (i.e., the matrigel-penetrated cells) remaining on the membrane were counted by a fluorescence microplate reader. Because LM2 cells express a GFP reporter gene (15), the fluorescence intensity (RFU) readout on the penetrated cells would be reflected only from BCa cells, not MEFs. As shown in Fig. 3, by comparing to LM2 cells alone, MEF-cocultured LM2 cells had a 4.3-fold increased invasiveness, demonstrating that surrounding fibroblasts stimulated cocultured BCa cell invasion.

To examine whether induced p16 could also inhibit the MEF-stimulated invasiveness, LM2/Tet-on-p16 with or without Dox induction were used for a similar assay. Consistently, p16 induction in the cocultured LM2 cells also significantly decreased invasiveness of the cocultured BCa cells (Fig. 4A). To exclude the possibility that Dox incubation per se might alter the BCa cell invasiveness, we also used parental LM2 cells cocultured with MEF for the same assay; we did not observe any effect from Dox treatment alone on the change of LM2 cell invasiveness (Fig. 4B).
Hypoxia and cocultured fibroblasts have additive effects on stimulation of BCa cell invasion ability

To examine whether hypoxia stimulates BCa cell invasiveness, we analyzed the matrigel-penetrating ability of LM2 cells in the presence or absence of cocultured MEF, under hypoxia or normoxia conditions. We found that hypoxia stimulated LM2 invasive ability at either LM2 growing alone (2.7-fold increase, compare column 1 and 2, Fig. 5) or coculturing with MEF (1.3-fold increase, compare column 3 and 4, Fig. 5). Consistent with what we observed earlier, MEF significantly simulated BCa cell invasion ability, either at normoxic condition (4.4-fold increase, compare column 1 and 3, Fig. 5) or hypoxia (2.2-fold increase, compare column 2 and 4, Fig. 5). The combination of hypoxia and cocultured MEF had apparent additive effects on stimulation of BCa LM2 invasive ability (5.9-fold increase, compare column 1 and 4).

HIF-1α from fibroblasts contributes to BCa cell invasion in coculture

Tumor-surrounding stromal cells (such as fibroblasts) are integral components of the tumor environment that together as an entity plays critical roles in the tumor progression. As both hypoxia and cocultured fibroblasts MEF stimulated BCa LM2 invasiveness (see above), we further investigated whether HIF-1α from tumor-surrounding fibroblasts cells plays a critical role for BCa cell invasion. We used a similar coculture setting as described above, with both HIF-1α WT MEF and HIF-1α KO MEF lines, respectively, to coculture with LM2 cells for the invasion assay. We found that HIF-1α WT MEF had a higher stimulation on cocultured LM2 invasiveness than its HIF-1α KO counterpart did (Fig. 6); by compared with LM2 alone, HIF-1α WT MEF had a stimulation of 10.9-fold increase of LM2 invasiveness, whereas HIF-1α KO MEF had 7.1-fold increase. These results implied that at least part (the difference of about 3.8-fold, Fig. 6) of the stimulation of LM2 invasion from the cocultured MEF was from HIF-1α effect of the surrounding MEF cells.

Discussion

Despite the fact that p16 has been studied extensively on its anti-tumor cell proliferation properties and related function as a negative cell-cycle regulator, the link between p16 and hypoxia-stimulated tumor progression remains unclear. Specifically, exploration of p16’s effects on cancer cell migration and invasion has barely begun. In this report, we investigated novel properties of p16’s functions on its anti-migration and anti-invasion abilities of BCa cells. We found that p16 is capable of inhibiting hypoxia-induced cell migration and invasion of BCa cells, and suppressing cocultured fibroblast-stimulated invasiveness of BCa cells.

We have previously reported that p16 downregulates VEGF gene expression and BCa angiogenesis by neutralizing HIF-1α transactivation on VEGF pathway (12,14). Consistent with our finding, a recent publication (18) revealed that p16 appears to repress the migration/invasion abilities of breast stromal fibroblasts and their VEGF-dependent angiogenesis through inhibition of Akt, a component of PI3K/Akt signaling pathway that regulates HIF-1α (19,20). Moreover, silencing HIF-1α in malignant gliomas effectively inhibited cell migration and invasion under hypoxic environment, suggesting that HIF-1α expression induced by hypoxia is an essential event in the activation of glioma cell motility and invasion (21).
Furthermore, a recently identified HIF-1α-direct target gene, HEF1 (human enhancer of filamentation 1), was shown to mediate hypoxia-induced migration of colorectal cancer cells (22). HEF1 was originally found as a scaffolding protein but recent data implicated it also acts as a key player in cancer cell migration and invasion (22-25). Elevated levels of HEF1 were found in various human cancers including BCa (26) and its expression proportionally correlates to tumor grade (27) and metastasis ability (28). Our unpublished results revealed that p16 inhibited hypoxia/HIF-1α stimulated HIF1 promoter transactivation (not shown). One possibility is that p16 may block the axis of “Hypoxia-HIF-1α-HEF1-migration/invasion” by neutralizing HIF-1α’s transactivation on HEF1 gene. We are in the process to further elucidate the mechanism of action of p16-mediated inhibition on BCa cell migration and invasion.

Although several studies suggested that tumor-surrounding microenvironment contribute to tumor growth and progression, the key attributes from tumor-surrounding cells that clearly display promotion for tumor progression have yet to be specified. While many studies have reported that HIF-1α protein in cancer cells plays an important role in cancer progression and various aspects of the tumor metastatic cascade, no study has yet reported on whether HIF-1α expression from tumor-surrounding cells contributes to tumor progression. We demonstrate here, for the first time, that HIF-1α from cocultured fibroblasts plays a critical role in promoting BCa cell invasion (Fig. 6). Consistent with our finding that HIF-1α from tumor-surrounding fibroblasts plays a critical role in promoting tumor cell invasion (Fig. 6), our recent data, in a 3-D culture system containing BCa 4T1 cells cocultured with HIF-1α WT or KO fibroblasts, revealed that HIF-1α from fibroblasts plays a critical role under hypoxia for the 3-D BCa cells’ protrusion formation (not shown). Protrusion formation is an established marker for invasive, aggressively malignant tumor type that start to invade the surrounding matrix in 3-D culture (29,30). The increased invasiveness of BCa may be partially due to the increased secreted growth factors (such as VEGF) that is upregulated by HIF-1α in the tumor microenvironment from surrounding HIF-1α-WT fibroblasts.

The development of intratumoral hypoxia is a common characteristic for fast-growing solid tumors including breast cancer. The intratumoral hypoxia activates HIF-1α in the tumor cells and its surrounding stromal cells, which in turn stimulate a group of HIF-1α-downstream genes responsible for tumor malignant progression including migration and invasion. Current chemotherapy for BCa treatment, including targeting the HIF-1α pathway, is not ideal due to toxic side effects of chemoagents and cancer resistance to them. One of the more promising approaches is the development of biologically based therapies to thwart the progression of metastatic disease (31). Specifically, a biological therapeutic approach that targets hypoxia-induced malignancy and directly neutralizes HIF-1α activity and its downstream genes mediating oncogenic pathways greatly deserves investigation. This study validates p16 as an effective biological agent to suppress hypoxia-promoted migration and invasion in BCa cells. The results of this study suggest that elevation and restoration of p16 in cancer microenvironment, either by gene therapy or pharmacology approaches, should provide a therapeutic mean to efficiently suppress BCa malignant progression.

Abbreviations

Breast cancer, BCa; Doxycycline, Dox; green fluorescent protein, GFP; hypoxia-inducible factor-1alpha, HIF-1α; immunohistochemistry, IHC; knockout, KO; tetracycline-induced regulated promoter, Tet-on promoter; vascular endothelial growth factor, VEGF; wild-type, WT.

Competing Interests

The authors have declared that no competing interest exists.

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