N-cadherin promotes epithelial-mesenchymal transition and cancer stem cell-like traits via ErbB signaling in prostate cancer cells

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Abstract. N-cadherin has been reported to be upregulated and associated with metastasis and poor prognosis in prostate cancer patients, however the underlying mechanism still remains puzzling. In the present study, we found that upregulation of N-cadherin enhanced, while downregulation of N-cadherin impaired the invasion, migration, and epithelial to mesenchymal transition (EMT) of prostate cancer (PCa) cells. Overexpression of N-cadherin increased the efficiency of colony and tumor spheroid formation and the stemness factor expression (including c-Myc, Klf4, Sox2 and Oct4), and vice versa. Furthermore, microarray analysis and western blot analysis mechanistically proved that N-cadherin activated ErbB signaling pathway by upregulating the expression of Grb2, pShc and pERK1/2. Importantly, the regulation of N-cadherin on EMT and stemness was counteracted by lapatinib, a specific ErbB signaling pathway inhibitor. Collectively, these findings demonstrate that N-cadherin regulates EMT and stemness of PCa cells via activating ErbB signaling pathway, which indicates the pivotal role of N-cadherin/ErbB axis in the metastasis of prostate cancer.

Introduction

Prostate cancer (PCa) is the second most frequently diagnosed cancer and bone metastasis is the principal issue, accounting for as many as 90% of patients with advanced PCa (1). The main therapeutic option for bone metastasis in hormone-responsive PCa is androgen deprivation therapy. Despite initial response rates of 80-90%, virtually all treated patients progress to androgen-insensitive disease, a state referred to as metastatic castration-resistant PCa (mCRPC). Although these agents effectively palliate symptoms and prolong life, mCRPC remains incurable (2,3). Therefore, an increased understanding of the mechanisms of PCa bone metastasis and metastatic castration-resistance is needed to develop novel therapeutic approaches.

Epithelial to mesenchymal transition (EMT), as a transient phenomenon involving in the process of metastasis of cancers, plays a key role in tumor cells invasion and metastasis (4). Cancer stem cells (CSCs) are a rare subpopulation of cells with stem cell-like properties which have been found in solid malignancies (5-7), and also are thought to be responsible for cancer relapse and metastasis (8,9). Recent evidence has showed that EMT can generate cancer cells with stemness properties (10). The study of Ribeiro and Paredes revealed that P-cadherin expression, which has been already identified as a breast cancer stem cell marker and invasive promoter, was probably able to identify an intermediate EMT state associated with a metastatic phenotype (11). This important finding implies a direct link between EMT and properties of CSCs. Therefore, unveiling the molecular mechanisms responsible for EMT and CSCs would be helpful to develop new promising therapies for PCa patients (12).

N-cadherin, as a marker of ongoing EMT, is not expressed in normal epithelial cells, but its expression has been demonstrated in several types of carcinomas (13-15). Recent study showed that FGFR signaling was responsible for the initiation of N-cadherin-driven EMT and stemness properties in breast cancer cells (16). In PCa, simultaneous upregulation of N-cadherin and downregulation of E-cadherin have been
found in more aggressive PCa lines, primary and metastatic PCa. Importantly, aberrant N-cadherin expression has been reported as crucial in PCa progression not only to metastasis, but also to castration resistance (13,17). Furthermore, in xenografts of castration-resistant PCa, a monoclonal antibody that targeted the ectodomain of N-cadherin inhibited androgen-independent growth, local invasion and metastasis. However, the underlying mechanism of N-cadherin in promoting PCa progression is not fully understood. As a switch from E-cadherin to N-cadherin plays a critical role in EMT and progression of PCa and high mortality (18), we hypothesize that N-cadherin positively regulates metastatic abilities of PCa cells by modulating EMT and stemness properties of PCa cells.

The ErbB family tyrosine kinases consists of four members, ErbB1-4, and ErbB1 and ErbB2 also known as EGFR and HER2, respectively. EGFR was the first receptor evidenced with a relationship between receptor overexpression and epidermoid carcinoma (19). Amplification and mutation of EGFR has been proved to be associated with poor prognosis in cancers, such as glioma (20-22), lung cancer (23-26) and breast cancer (27,28). HER2 has been also reported to be amplified and been widely studied in breast cancer. HER2 serves as an important prognosis maker and therapy target for breast cancer (29-31). Furthermore, it has been reported that HER2 could induce EMT in both mammary epithelial cells (32) and breast cancer cells (33,34). Furthermore, activation of ERBB2 and ERBB3 was able to mediate glioblastoma cancer stem-like cell resistance to EGFR-targeted inhibition (35) and activation of EGFR was reported to promote acquisition of stem cell-like properties in head and neck squamous cell carcinoma (36). Currently, the underlying mechanism between ErbB signaling and EMT and stemness of cancer cells is poorly understood, and the role of ErbB signaling in PCa is largely unknown.

In the present study, we reported that N-cadherin positively regulated invasion, migration, EMT and stemness properties of PCa cells. Importantly, through microarray analysis and further test, we found that overexpression of N-cadherin activated ErbB signaling, but not FGF signaling. Furthermore, our results demonstrated that N-cadherin regulated EMT and stem cell-like property linked with ErbB signaling in PCa cells. Taken together, N-cadherin might serve as a novel potential therapeutic target in PCa.

Materials and methods

Cells and cell culture. The brain metastatic cell line DU145 and the bone metastatic PCa cell line PC-3 were purchased from the American Type Culture Collection (ATCC) and grown in DMEM culture medium (Hyclone) and Ham’s F-12 culture medium (Hyclone) respectively, supplemented with 10% fetal bovine serum (Hyclone). Cells were grown at a humidified atmosphere of 5% CO2 at 37°C. Labatinib was purchased and prepared as a 10 mM concentrated stock solution in dimethyl sulfoxide (Fisher Scientific).

Vectors and retroviral infection. N-cadherin gene was amplified from cDNA by RT-PCR and cloned into the pMSCV-EF2 lentiviral vector. Two human N-cadherin-targeting shRNA sequences were cloned into pSuper-retro-puro to generate pSuper-retro-N-cadherin RNAi (s) and the sequences of RNAi#1 and RNAi#2 are GCTGAAAGAACTGAAGCATTT and AATGCTTTAGTTTCTTCAGC, respectively. Retroviral production and infection were performed as previously described (37). Stable cell lines expressing N-cadherin or N-cadherin shRNAs were selected for 10 days with 0.5 mg/ml puromycin.

Microarray analysis. Total RNA from PC-3/vector, PC-3/N-cadherin-RNAi and N-cadherin-overexpression were extracted. Total RNA from each sample was quantified by the NanoDrop ND-1000 and RNA integrity was assessed by standard denaturing agarose gel electrophoresis (38). Microarray analysis was performed commercially by the Shanghai Biochip Corp, according to standard Agilent protocol. Briefly, integrity and concentration of RNA was assessed after RNA extraction and prior to sample labeling. Total RNA of each sample was used for labeling and array hybridization with the following steps: i) reverse transcription with Invitrogen Superscript ds-cDNA synthesis kit; ii) ds-cDNA labeling with NimbleGen one-color DNA labeling kit; iii) array hybridization using the NimbleGen Hybridization system and followed by washing with the NimbleGen wash buffer kit; iv) array scanning using the Axon GenePix 4000B microarray scanner (Molecular Devices Corp.). Data were extracted and normalized using NimbleScan v2.5 software. Results are provided in the NimbleScan Generated Data Folder. Further data analysis was performed using Agilent GeneSpring GX v11.5.1 software. Bioinformatics analysis and visualization of microarray data were performed with the MeV v4.4 program (http://www.tm4.org/mev/) (39).

Quantitative reverse transcription-PCR. The procedure was performed according to the instruction of All-in-One™ miRNA qRT-PCR Detection kit (GeneCoopoeia, USA), as described previously (40). The relative expression levels from three independent experiments were counted following the 2-ΔΔCt method of Livak and Schmittgen (41). The qRT-PCR primers for N-cadherin, c-Myc, OCT-4, SOX2, Klf4 and glyceraldehyde-3-phosphate dehydrogenase (GAPDH) were designed by the Primer Express version 2.0 software (Applied Biosystems). N-cadherin forward, 5’-GGCATACACCATGGCATCCTTT-3’; reverse, 5’-GTGCGATAGGACACGAGCTCT-3’; c-Myc forward, 5’-CAAGCTTCATTACAGGCTTGAT-3’; reverse, 5’-GCTGCTTTAGACGCCGTTGATT-3’; OCT-4 forward, 5’-TCTCCAGGGTCCTCCACT-3’; reverse, 5’-GTGAGAGAGTGCAACAACA-3’; SOX2 forward, 5’-GTCTTCTTCTCTCAATGTTGTT-3’; reverse, 5’-ATCGGCAAGGATTTTCGTAAC-3’; Klf4 forward, 5’-CCCCGGTGTGTTACCGATG-3’; reverse, 5’-GAGTTCCGAAATGCGGTAAGC-3’; GAPDH forward, 5’-ACATCCCTGACCCACAAA-3’; reverse, 5’-TAGCCAAATCATACTGCTCGTC-3’.

Western blotting. For the analysis of expression of related proteins, western blot assay was performed according to a standard method, as described previously (42). The following primary antibodies were used: mouse anti-vimentin, mouse anti-E-cadherin (CST, cell signal technique); mouse anti-fibroconnectin, mouse anti-N-cadherin (BD Biosciences); mouse anti-Grb2, anti-pShc and anti-pERK1/2 (Abcam). Blotting
membranes were stripped and re-probed with anti-tubulin antibody (Sigma) as a loading control. Nuclear extracts were prepared using the Nuclear Extraction kit (Active Motif), according to the manufacturer's instructions.

Wound healing assay. PCa cells were cultured on 6-well plates with DMEM containing 10% FBS to 90% confluency and scratched with a sterile 10 µl pipette tip to create artificial wounds. After scratching, the detached and damaged cells were carefully washed with phosphate-buffered solution (PBS) and maintained in 10% fetal bovine serum media. Progression of migration was observed and photographed at 24 h after wounding. Images of the cells migrating into the wound were taken at the time points of 0, 6 and 12 h by an inverted microscope (x40).

Invasion assays. Cell invasion assays were performed using Transwell chambers consisting of 8-mm membrane filter inserts (Corning; Corning Inc.) coated with Matrigel (BD Biosciences). Briefly, the trypsinized PCa cells were resuspended in serum-free medium and seeded in the upper chamber. Then, the lower chamber of the Transwell was filled with 1 ml Ham's F-12 medium or T-medium supplemented with 10% FBS to 90% confluency and seeded in the upper chamber. The trypsinized PCa cells were resuspended in serum-free medium and seeded in the upper chamber. Then, the lower chamber of the Transwell was filled with 1 ml Ham's F-12 medium or T-medium supplemented with 10% FBS. After incubation for 24-48 h at 37°C in 5% CO₂, cells passing through the coated membrane to the bottom side of the inserts were fixed with 4% paraformaldehyde and stained with hematoxylin. The non-migratory cells on the upper chamber were removed with cotton swabs, and the migratory cells were stained, photographed, and quantified by counting them in 5 random high-power fields.

Colony formation assay. The cells were trypsinized as single cells and suspended in the media with 10% FBS. Indicated cells (300 cells per well) were seeded into of 6-well plate for ~10-14 days. Colonies were stained with 1% crystal violet for 10 min after fixation with 10% formaldehyde for 5 min. Plating efficiency = number of colonies (≥50 cells per colony)/per input cells x 100%. Different colony morphologies were captured under a light microscope (Olympus).

Self-renewing spheroid formation assay. Indicated cells (500 cells/well) were seeded into 6-well Ultra Low Cluster plate (Corning) and were cultured in suspension in serum-free DMEM/F12 (BioWhittaker), supplemented with 2% B27 (Invitrogen), 20 ng/ml EGF (BD Biosciences), 20 ng/ml bFGF (PeproTech), 5 µg/ml insulin (Sigma) and 0.4% bovine serum albumin (Sigma). After 10-12 days, the number of cell spheres (tight, spherical, non-adherent masses >50 µm in diameter) were counted, and image of the spheres were captured under inverse microscope. Sphere formation efficiency = colonies/input cells x100%.

Luciferase reporter assay. Luciferase assays were carried out in 293FT cells that were co-transfected with miRNAs and luciferase reporter plasmids in 24-well plates and cultured for 48 h before the cells were harvested and lysed for luminescence detection. Subsequent processing and detection were performed by using the luciferase assay kit (Promega) according to the manufacturer’s protocols. Renilla luciferase was activated to emit primary luminescence, and firefly luminescence was used for normalization. Each test was repeated in triplicate.

Statistical analyses. All statistical analyses were carried out using SPSS 17.0 statistical software package. Means ± SD was calculated and two-tailed Student's t-test or one-way ANOVA was performed using data analysis tools provided in the software package. In all cases, P<0.05 was considered statistically significant.

Results
N-cadherin promotes invasion, migration and EMT of PCa cells. To investigate the biological function of N-cadherin in PCa cells, we transduced PC-3 and DU145 PCa cells with constructed pMSCV/N-cadherin-derived retrovirus to establish N-cadherin-overexpressing stable cell lines. As N-cadherin was not highly expressed in DU145 human prostate cell lines (43,44), N-cadherin-specific RNA interference (RNAi) oligonucleotides were cloned into a retroviral transfer vector pSuper-retro-puro to establish N-cadherin-low expressing stable cell lines only in PC-3 (Fig. 1A). Western blot analysis suggested that N-cadherin expression is positively related with the expression of mesenchymal makers fibronectin and vimentin, but negatively related with the expression of epithelial marker E-cadherin in PC-3 and DU145 cells (Fig. 1B). On the contrary, the low expression of N-cadherin significantly reduced the expression of fibronectin and vimentin and increased the expression of E-cadherin in PC3 cells (Fig. 1B). The result suggested that N-cadherin expression might be involved in EMT of PCa cells. Firstly, upon knockdown of N-cadherin in PC-3 cells, we noted that PC-3 cells underwent a marked change in morphology, from spindle like morphology to epithelial transition (Fig. 1C). Moreover, Transwell matrix penetration assay revealed that overexpression of N-cadherin promoted, while silencing N-cadherin strongly repressed the invasive ability compared to vector cells (Fig. 1D). In addition, wound healing assay showed that upregulation of N-cadherin increased, while knockdown of N-cadherin decreased healing speed of the scratch in transduced cells (Fig. 2). These data demonstrated that N-cadherin was able to modulate the invasion and migration properties of PCa cells.

N-cadherin promotes colony, spheroid formation and Sox2, c-Myc, Oct4 and Klf4 expression in PCa cells. Numerous studies suggested that cancer stem cells are involved in tumor metastasis (10,45). Firstly, colony formation assays was used to explore whether N-cadherin had an effect on viability of PCa cells. As N-cadherin was not highly expressed in DU145 human prostate cell lines (43,44), N-cadherin-specific RNA interference (RNAi) oligonucleotides were cloned into a retroviral transfer vector pSuper-retro-puro to establish N-cadherin-low expressing stable cell lines. As N-cadherin was not highly expressed in DU145 human prostate cell lines (43,44), N-cadherin-specific RNA interference (RNAi) oligonucleotides were cloned into a retroviral transfer vector pSuper-retro-puro to establish N-cadherin-low expressing stable cell lines only in PC-3 (Fig. 3A). As shown (Fig. 3B), the number of colonies (% plating efficiency) was 85.7% in PC-3/N-cadherin versus 55.9% in DU145/vector. Colonies with different morphologies were classified as holoclones, meroclones and paraclones (46). Holoclones are generally classified as holoclones, meroclones and paraclones (46).
Figure 1. N-cadherin promotes invasiveness and EMT in PCa cells. (A) Expression of N-cadherin was detected by real-time PCR and western blotting (*p<0.01). (B) N-cadherin expression is positively related with the expression of mesenchymal makers fibronectin and vimentin, but negatively related with the expression of epithelial marker E-cadherin in PC-3 and DU145 cells. (C) Downregulation of N-cadherin converted a stick-like or long spindle-shaped mesenchymal profile to a cobblestone-like or a short spindle-shaped epithelial morphology in PC-3 cells. (D) Overexpression of N-cadherin increased invasion in PC-3 and DU145 cells, while downregulation of N-cadherin efficiently reduced invasion in PC-3 cells (*p<0.01).
more round and tightly packed; paraclones are irregular in composition and often contain more elongated or flattened cells; and meroclonies are an intermediate phenotype (Fig. 3C). We did not find typical holoclones in PC-3 and DU145 cells. The proportion of meroclonies was 87.6% in PC-3/N-cadherin, 77.2% in PC-3/vector and 78.1% in PC-3/scramble, 52.2% in PC-3/N-cadherin-RNAi#1 and 55.4% in PC-3/N-cadherin-RNAi#2 cells, and 84.7% in DU145 cells of N-cadherin overexpression versus 75.3% in DU145/vector. Overexpression of N-cadherin significantly increased the proportion of meroclonies of PCa cells (p<0.01, Fig. 3D). Moreover, sphere formation assays suggested that upregulating N-cadherin enhanced the number and size of tumor spheroids in both PC-3 and DU145 cells, while downregulating N-cadherin inhibited the number and size of tumor spheroids in PC-3 cells (Fig. 4A). The spheroid formation efficiency was 7.1% in PC-3/N-cadherin versus 4.2% in PC-3/vector, 4.4% in PC-3/scramble versus 2.0% in PC-3/N-cadherin-RNAi#1 cells and 2.2% PC-3/N-cadherin-RNAi#2, and 6.7% in DU145 cells overexpressing N-cadherin versus 4.0% in DU145/vector (Fig. 4B). Real-time PCR was performed to examine the mRNA level of pluripotency-associated markers including Sox2, c-Myc, Oct4 and Klf4. The result suggested that CSC markers were significantly higher expressed in N-cadherin-transduced cells than control group and significantly lower expressed in N-cadherin-RNAi cells (Fig. 4C). Thus, our results suggested that overexpression of N-cadherin promotes prostate CSC-like traits.

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Figure 2. N-cadherin promotes the ability of wound healing in PCa cells. (A) Overexpression of N-cadherin increased, while downregulation of N-cadherin efficiently suppressed migration in PC-3 cells. (B) Overexpression of N-cadherin increased migration in DU145 cells.
N-cadherin promotes EMT and CSC-like traits of PCa cells via ErbB signaling pathway. To understand the underlying mechanism of N-cadherin in regulation of EMT and stemness of PCa cells, we performed microarray analysis on PC-3/vector and PC-3/N-cadherin cells. As shown in Fig. 5, the microarray data showed a comprehensive activation of ErbB in PC-3/N-cadherin cells compared to PC-3/vector cells, which indicated that N-cadherin might achieve its function via ErbB signaling. To further confirm the data, downstream ERBB signaling component Grb2 and phosphorylation of components Shc and ERK were examined by western blot analysis. We found that overexpression of N-cadherin enhanced the expression of Grb2, pShc and pERK1/2 (Fig. 6A).

In order to explore whether the stem cell-like phenotype in PCa cells is associated with ErbB signaling activation, we also analyzed the impact of blocking ErbB signaling on the stem cell population capability of N-cadherin transduced PC-3 and DU145 cell lines. We used lapatinib, a potent ATP-competitive inhibitor, to inhibit both EGFR and HER2 (100 nM). Western blot analysis showed that lapatinib successfully reduced downstream Grb2, pShc and pERK1/2 in PC-3 and DU145 cells (Fig. 6B). Furthermore, western blot analysis showed that fibronectin and vimentin expression were suppressed in the lapatinib treated PCa cells, while expression of E-cadherin was increased (Fig. 6C). Colony formation assays suggested that the number of colonies were inhibited in the treatment cells compared with N-cadherin-transduced cells (Fig. 6D). Real-time PCR analysis showed that the mRNA expression level of ‘stemness’ factors c-Myc, Oct4, Klf4, and Sox2 were downregulated in the lapatinib-treated cells compared with...
Figure 4. N-cadherin enhanced tumor sphere formation and stemness factor expression. (A and B) Overexpression of N-cadherin increased, while downregulation of N-cadherin efficiently suppressed spheroid formation (*p<0.05; **p<0.01). (C) Overexpression of N-cadherin increased, while downregulation of N-cadherin significantly reduced the mRNA expression levels of C-Myc, Oct4, Sox2 and Klf4 (*p<0.01).

Figure 5. Expression profiling of signaling pathway in N-cadherin overexpressing PC-3 cells.
N-cadherin-transduced cells (Fig. 6E). All the above data revealed that N-cadherin regulates EMT and CSC-like traits of PCa cells via ErbB signaling pathway. *N-cadherin does not mediate the function of miR-145 in PC-3 cells.* Analysis using the publicly available algorithms (e.g., TargetScan, miRANDA) indicated that the N-cadherin-
3'-UTR region is the theoretical conserved target of miR-145 (Fig. 7A). The expression level of the N-cadherin mRNA and protein was significantly decreased in miRNA-145-transfected cells (Fig. 7B-D). However, upon transfection with miR-145, we did not find strong reduction in luciferase activity from pGL3-N-cadherin-3'-UTR (Fig. 7E). Our results indicated that miR-145 did not directly target N-cadherin in PC-3 cells.

Discussion

In this study, we present a pivotal finding that overexpression of N-cadherin promoted, while knockdown of N-cadherin repressed EMT and stem cell-like property in PCa. Microarray analysis and further test mechanistically demonstrated that N-cadherin achieve EMT and stemness promoting function by activating the ErbB signaling. These findings provide strong evidence that, for the first time, elevated N-cadherin promotes EMT, stemness and metastatic ability via ErbB signaling pathway in PCa cells, which further supported that N-cadherin could be a therapeutic target in PCa.

N-cadherin, a mesenchymal cadherin associated with epithelial-to-mesenchymal transition, has been widely studied in various tumor types. Hazan and colleagues reported that N-cadherin promotes adhesion between invasive breast cancer cells and the stroma (47) and overexpression of N-cadherin correlates with invasiveness in breast carcinoma (48). In PCa, N-cadherin has also been demonstrated to be elevated and associated with poor prognosis (13,49). An important finding by Tanaka and colleagues demonstrated that monoclonal antibody targeting of N-cadherin could inhibit PCa growth, metastasis and castration resistance (17). Here, we found that increasing N-cadherin enhanced, while silencing N-cadherin impaired the invasion and migration of PCa cells. Western blot analysis showed that overexpression of N-cadherin enhanced the expression of mesenchymal cell markers, fibronectin and vimentin, and decreased the expression of epithelial cell markers, E-cadherin, and downexpression of N-cadherin inhibited the expression of fibronectin and vimentin, and increased the expression of E-cadherin. Moreover, upregulation of N-cadherin also enhanced the stem cell-like property of PCa cells as indicated by higher tumor spheroids and colony formation efficiency and increased the expression of stem cell property-associated factors, including Sox2, c-Myc, Oct4 and Klf4, and vice versa. All together, the above presented the pivotal role of N-cadherin in promoting metastasis in PCa.

Although a recent study showed that N-cadherin-driven EMT and stemness properties depend on FGFR activation, ERK activity, matrix metalloproteinase 9 production and on selective inhibition of the AKT3 isoform in many solid tumors (14,16,50), our results showed that the mechanism via which N-cadherin regulates invasion, migration, EMT and stem cell-like property in PCa is different from that in breast cancer. The present findings indicated that N-cadherin-driven EMT and stemness properties at least partially depend on ErbB signaling in PCa cells. Furthermore, accumulating evidence also shows that ErbB signaling pathway is associated with poor prognosis in various cancers. In glioma, ErbB1, known as EGFR, extracellular missense mutations as a novel mechanism for oncogenic EGFR activation may help identify patients.

Figure 7. N-cadherin is not an authentic target of miRNA-145 in PCa cells. (A) Wild-type 3'-UTR sequence of N-cadherin was cloned into luciferase plasmids (Luc-N-cadherin-3'-UTR-wt). Solid lines indicate binding sites. (B) Expression level of the N-cadherin was detected in PC-3 cells infected with miR-145 mimics. (C) miR-145 expression was detected by real-time PCR. (D) mRNA level of N-cadherin was detected by real-time PCR (**p<0.01). (E) Luciferase assay of PC-3 cells transfected with pGL 3-N-cadherin-3'-UTR reporter with miR-145 mimic of increasing amounts (20 and 50 nM).
who can benefit from EGFR kinase inhibitors for treatment of glioblastoma (22); in lung cancer, especially non-small cell lung carcinoma (NSCLC), it has been reported that EGFR amplification is universal and EGFR mutations in lung carcinomas make the disease more responsive to treatment with tyrosine kinase inhibitors (23-25); in breast cancer, EGFR was regarded as a predictor of early recurrence and death (27), and ErbB2, also known as HER2, serves as an important prognosis maker and therapy target for breast cancer (29). Further studies have demonstrated that HER2 could induce EMT in both mammary epithelial cells (32) and breast cancer cells (33,34). Furthermore, some evidence reported that glioblastoma cancer stem-like cell resistance to EGFR-targeted inhibition was mediated by activation of multiple ERBB family receptors (35). In head and neck squamous cell carcinoma, EGFR kinase promotes acquisition of stem cell-like properties (36). Therefore, all the above-mentioned evidence indicated that ErbB signaling pathway plays an important role in metastasis of cancer.

For most PCa patients who were identified in the early stages the initial therapies mostly result in significant long-term remission (51). However, for advanced metastatic cases these treatments, such as prostatectomy, radiation and chemotherapy, show little benefit and without effective control the patients eventually die of the disease. Although androgen deprivation and chemotherapy are currently effective treatments for these patients, development of hormone ablation resistance is inevitable (52), which is termed castration-resistance PCa. Consequently, improved understanding of the mechanisms underlying mCRPC progression has contributed to the recognition of multiple molecular targets and advances in the therapeutic landscape. Recent study indicated that N-cadherin is crucial in PCa progression not only to metastasis, but also to castration resistance (17). Our results showed that N-cadherin promoted EMT and stemness of CSCs of PCa cells by upregulating ErbB signaling. Because EMT and CSCs play crucial roles during the development of castration-resistance in PCa (12), one of the important mechanism by which N-cadherin increased castration resistance of metastatic PCa cells may promote EMT and stemness of CSCs of PCa cells by upregulating ErbB signaling. Therefore, understanding the inhibition of this signaling pathway may be useful to develop new therapies for metastatic PCa.

A wide range of studies have demonstrated that many microRNAs (miRNAs), as crucial post-transcriptional regulators repressing the expression of their target genes, play a pivotal role in solid tumor metastasis via regulating migration, invasion and EMT of cancer cells and the properties of CSCs (53-58). Our previous studies found that miR-145 played an important role in inhibiting migration, invasion, EMT and stemness properties of PCa cells via different targets (40,59,60). Furthermore, recent study showed that N-cadherin was a direct target of miR-145 and promoted the invasion-metastasis cascade in gastric cancer (61). Moreover, our present results showed that N-cadherin promoted EMT and stemness properties of PCa cells. Therefore, we can suppose that N-cadherin mediates the function of miR-145 in regulating EMT and stemness properties in PCa cells. However, we used dual-luciferase reporter gene assay to check the relationship between miR-145 and N-cadherin, and found miR-145 did not targets N-cadherin in PC-3 cell. This difference may be due to the using of different tumor models. It will be of great interest to investigate the reasons why miR-145 is off-target for N-cadherin in PCa cells.

In conclusion, this study demonstrated that N-cadherin promotes invasion, migration, EMT and stemness of PCa cells, which suggest a pivotal role of N-cadherin in metastasis and castration resistance of PCa cells. Importantly, these findings exploit interesting and realistic avenues for cancer therapies using N-cadherin antagonists.

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References


