Abstract. Breast cancer cell lines and mouse models are valuable tools for investigating the biology of and developing potential therapeutics for human breast carcinoma. The PTEN-/-/NIC mouse is a genetically engineered mouse model for ErbB2/Neu-overexpressing/-PTEN deficient breast carcinoma with histopathological and molecular features relevant to the luminal subtype of primary human breast cancer. However, the PTEN-/-/NIC model develops multifocal and aggressive mammary tumors with a short life-span, which greatly impedes its preclinical usage. To complement the genetic engineering approach and to facilitate the future application of this model, in the present study, two newly established cell lines, NICP20 and NICP21, from PTEN-/-/NIC mammary tumors are described. These NICP20 and NICP21 cells retained the crucial molecular phenotype similar to the origin, as confirmed by genotyping and western blot analysis. These cells induced tumors in immunocompetent syngeneic mice by mammary fat pad injection and produced lung metastasis when injected intravenously. Tumors induced by these cells displayed luminal-like histologic morphology and hyperactivation of Akt which are similar to PTEN-/-/NIC tumors. Immunohistochemical staining also revealed that tumors induced by the NICP20 and NICP21 cells showed a high proliferative level, comparable angiogenesis and T-cell infiltration properties similar to PTEN-/-/NIC tumors. Therefore, these NICP20 and NICP21 cells represent an alternative and useful model system to enhance our understanding of the nature of ErbB2-positive breast cancers, particularly accompanying PTEN loss and to facilitate further experimental therapeutic studies.

Introduction

Breast cancer remains one of the most common malignancies and is the second leading cause of cancer-related mortality in women (1). Standard treatment modalities have improved the overall survival and the quality of life of patients. However, more personalized therapies or improvement of existing treatments are needed due to the heterogeneity of breast cancer and resistance to existing therapies. The molecular profiling of human breast cancers has identified at least 5 subtypes with distinct clinical outcomes (2-5). One of the common subgroups is the ErbB2-positive subtype, which occurs in 20-30% of breast cancer cases. It has been demonstrated that elevated expression/amplification of ErbB2 correlates with poor prognosis (6-8).

Mutation or deficiency of the tumor-suppressor phosphatase and tensin homologue (PTEN) has been reported to occur in 5-10% of human breast cancer cases (9). Loss of PTEN function results in hyperactivation of the PI3K/AKT pathway and induction of basal-like breast cancers (10,11). Furthermore, numerous studies have indicated that loss of PTEN confers resistance to trastuzumab (Herceptin), a humanized monoclonal antibody targeting ErbB2 (12-14). The above findings suggest that PTEN disruption may play a critical role in ErbB2-positive human breast cancer. To directly evaluate the effect of PTEN loss on ErbB2-induced mammary tumorigenesis and progression, Shade et al. generated the PTEN-deficient/NIC genetically engineered mouse model. This novel model utilized the murine mammary tumor virus (MMTV) promoter to drive co-expression of activated ErbB2/Neu and Cre recombinase coupling PTEN conditional depletion in the same mammary epithelial cells (15). PTEN-deficient/NIC mice exhibited rapid formation of highly metastatic mammary tumors and displayed histopathological and molecular features characteristic of the luminal subtype of primary human breast cancer (15,16). We found previously that loss of both PTEN alleles (PTEN-/-/NIC mice) resulted in significant resistance to Neu antibody treatment (17). Therefore, PTEN-/-/NIC mice represent a valuable tool for biological study and drug discovery for trastuzumab-resistant ErbB2/Neu-positive breast cancer.

However, there are several disadvantages and challenges of using the PTEN-/-/NIC model to preclinically evaluate novel...
therapeutics. First, it has a high cost, is time consuming and is labor intensive to obtain cohorts of female PTEN-/-/NIC mice. Second, it may be very difficult to quantify tumor size during preclinical investigations due to the fact that the PTEN-/-/NIC model develops multifocal and aggressive mammary tumors. To partially overcome the major drawbacks and complement the genetic engineering approach, we attempted to establish cell lines from spontaneous mammary tumors that arose in the PTEN-/-/NIC mice. In the present study, two cell lines derived from PTEN-/-/NIC mammary tumors were generated and characterized both in vitro and in vivo, providing an alternative and useful resource for future studies.

Materials and methods

**Cell lines and cell culture.** Cells were cultured in growth medium [DMEM/F12 with 10% FBS, 5 µg/ml insulin, 10 ng/ml epidermal growth factor (EGF), 1 µg/ml hydrocortisone, 35 µg/ml bovine pituitary extract and 100 U penicillin/streptomycin]. The cell number was evaluated on a cell counter. TM15 cells were established from a spontaneous ErbB2/Neu-positive/PTEN wild-type (PTEN+/+-ErbB2KI) mammary tumor (18,19). Genetic structure and general procedures

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**Mammary fat pad and tail vein injection of cells.** Confluent cells (70-80%) were digested with trypsin and washed with PBS. Cells were counted and resuspended to a final concentration. For mammary fat pad injection, 10⁶, 5x10⁵, 10⁵, 5x10⁴ or 10⁴ cells in 100 µl PBS/Matrigel (1:1) were injected into the #2 inguinal mammary gland of FVB/N female mice at 8 weeks of age. For tail vein injection, mice were anesthetized and 2x10⁴ cells in 100 µl PBS were injected using a 1-ml syringe (29 G; BD insulin syringe). FVB/N mice were obtained from the Harlan Laboratory. All animal experiments were performed according to the Guidelines for the Institutional Animal Care and Use Committee of The University of Texas M.D. Anderson Cancer Center and School of Medicine, Nanjing University.

**Polymerase chain reaction (PCR) for genotyping.** Genomic DNA was extracted from the cells at 70-80% confluency by digestion with proteinase K (Sigma) in 200 µl of lysis buffer in a 60°C incubator for 2 h, followed by heating at 95°C for 10 min to inactivate the enzyme. After centrifugation, 1 µl of the supernatant was used for PCR reaction. Primer sequences for NEU were: 5'-TTCCGGAACCCACATCAGGCC-3' and 5'-GT TTCTTCGACGAGCTACGC-3'; for CRE, 5'-TGCTCTGTGC CGTTTGCGC-3' and 5'-ACTGTTGTCCAGAACCAGGC-3'.

**Immunoblotting.** Cells in monolayer were washed with ice cold-PBS and harvested by scraping in lysis buffer (1% Triton X-100, 50 mM HEPES pH 7.4, 150 mM NaCl, 1.5 mM MgCl₂, 1 mM EGTA, 100 mM NaF, 10 mM Na pyruvate, 1 mM Na₃VO₄, 10% glycerol) containing protease inhibitors (Sigma) and phosphatase inhibitors (Roche). The lysate was incubated on ice for 20 min and centrifuged at 15,000 x g for 15 min at 4°C. The protein concentration of the supernatant was determined by BCA assay (Thermo Scientific), and 30 µg total protein was electrophoresed on 10% SDS-PAGE and then transferred onto a nitrocellulose membrane (Bio-Rad). Pten, pAKT S473 and Akt antibodies were obtained from Cell Signaling Technology. Antibodies for ErbB2/Neu and β-actin were obtained from Santa Cruz Biotechnology and Sigma, respectively. The blots were incubated with HRP-conjugated secondary antibodies and visualized by ECL (Amersham).

**Whole mount and histology.** Number 4 mammary gland was excised from 4- to 6-week-old virgin female PTEN-/-/NIC mice and spread on glass slides for overnight fixation in 4% paraformaldehyde. The next day the samples were hydrated and stained in a filtered solution of 0.2% carmine (Sigma) and 0.5% aluminum potassium sulfate for 1-2 days. Glands were then dehydrated sequentially through 70, 90 and 100% ethanol for 15 min each, precleared in toluene and stored in methylsalicylate. Lung whole mount preparations were harvested and infused with formalin. Tumor tissues and lungs were fixed in formalin, processed routinely and embedded in paraffin. For histological analysis, paraffin sections (4-µm) were stained with hematoxylin and eosin according to standard protocols.

**Immunohistochemistry.** Paraffin-embedded tumor sections (4-µm) were subjected to antigen retrieval in a pressure cooker with sodium citrate buffer (pH, 6.0) and incubated with antibodies specific for phospho-Akt S473 (1:100; Cell Signaling), Ki-67 (1:200; Dako), CD34 (1:50; eBioscience), CD3 (1:300; Epitomics) overnight at 4°C. Biotin-conjugated secondary antibodies were used. Remaining steps were performed using Vectastain ABC kits (Vector Laboratories). Slides were counterstained with hematoxylin. The stained slides were evaluated by two pathologists, and images were acquired using a Zeiss microscope with Axiosvision software (Carl Zeiss, Inc.).

**Results**

PTEN-/-/NIC mice develop multifocal and aggressive mammary tumors. Genetic structure and general procedures to generate the PTEN-/-/NIC mice are depicted in Fig. 1A. PTEN-/-/NIC mice developed tumors rapidly with a latency of 28 to 43 days (data not shown), which was consistent with previous results (15,17). As shown in Fig. 1B, a tumor mass was found in the inguinal mammary gland from a female
PTEN−/−/NIC mice developed multifocal and aggressive mammary tumors at 30-days old. All female PTEN−/−/NIC mice developed multifocal mammary tumors (Fig. 1C and D). Due to the rapid tumor growth and progression, most female PTEN−/−/NIC mice had to be euthanized three to four weeks after initial tumor detection (tumor size >15 mm in diameter). The overall survival curve of these female mice is shown in Fig. 1E with a median survival time of 50 days.

Establishment and characterization of the mammary tumor cells from PTEN−/−/NIC mice in vitro. Given the multifocal and aggressive features of the PTEN−/−/NIC tumors, it was difficult to follow tumor growth. Meanwhile, to partially overcome the drawbacks of the genetic engineered model (e.g. high cost, time consuming and labor intensive), we attempted to establish cell lines from tumors that arose in the PTEN−/−/NIC mice. According to protocols as previously described (21), we obtained two cell lines, named NICP20 and NICP21, respectively. Firstly, the NEu and CRE transgenes in the NICP20 and NICP21 cells were confirmed by PCR (Fig. 2A). As expected, western blot analysis showed no detectable PTEN expression and a high level of activated Akt both in the NICP20 and NICP21 cells (Fig. 2B). These results revealed that the NICP20 and NICP21 cells retained the critical molecular phenotype similar to the origin. Moreover, gross appearance

Figure 1. PTEN−/−/NIC mice develop multifocal and aggressive mammary tumors. (A) Genetic structure and generation of PTEN−/−/NIC mice are depicted. (B) Whole-mount staining of an inguinal mammary gland from a female PTEN−/−/NIC mouse at 30 days of age. Scale bar, 1 cm. The arrow indicates the tumor mass. (C and D) Gross observation of a female PTEN−/−/NIC mouse with multifocal mammary tumors (C) and appearance of the dissected tumors (D). (E) The overall survival curve of the female PTEN−/−/NIC mice (n=11). Median survival time was 50 days. PTEN, phosphatase and tensin homologue.

Figure 2. Establishment and characterization of the ErbB2/Neu-positive/PTEN-deficient mammary tumor cells from the PTEN−/−/NIC mice in vitro. (A) Genotyping for NEu and CRE using genomic DNA from the indicated cells. (B) Western blot analysis for Pten and phospho-Akt using lysates from the indicated cells. MT104T and TM15 cells were previously generated from PTEN−/−/ErbB2KI and PTEN+/+/ErbB2KI mammary tumors, respectively. (C) Representative image of the cultured cell lines from PTEN−/−/NIC tumors. PTEN, phosphatase and tensin homologue.
of the cultured cells displayed epithelial morphology over subsequent passages (Fig. 2C).

*Tumorigenesis and lung metastasis of the NICP20 and NICP21 cells in syngeneic mice.* After generation of the above two ErbB2/Neu-positive/PTEN-deficient cell lines, the tumorigenic ability of these cells in immune intact, syngeneic FVB/N female mice was initially evaluated. The initial round of intramammary injections utilized $10^6$, $5 \times 10^5$, $10^5$, $5 \times 10^4$ or $10^4$ cells for each cell line. By 2-5 weeks post injection, all mice had palpable tumor nodules in the mammary glands. Seven weeks post injection of $10^6$ NICP20 cells (Fig. 3A) and 8 weeks post injection of $5 \times 10^5$ NICP21 cells (Fig. 3B), the tumors were $>15$ mm in diameter and the mice were euthanized. Histological analysis revealed that the tumors induced by the NICP20 and NICP21 cell lines had large, solid nodular nests that resembled luminal-like histologic features (Fig. 3C and D) similar to the morphology of the original PTEN$^{-/-}$/NIC tumors. Expression of ErbB2/Neu and PTEN protein in the tumors induced by NICP20 and NICP21 cells was also determined by
western blotting (Fig. 3E). The weak bands of PTEN protein confirmed the expression of PTEN in the stromal tissue but not in the tumor cells. As expected, loss of PTEN increased Akt phosphorylation relative to the PTEN wild-type MMTV-NIC tumors (Fig. 3E). Meanwhile, the lung tissues of all the tumor-bearing mice (induced by NICP20 or NICP21 cells) were examined macroscopically and microscopically when euthanized, and no metastatic lesion was found. Based on this finding, tail vein injections were performed using different cell concentrations. Both cell lines produced lung metastasis when only 10⁴ cells were injected through the tail vein (Fig. 4).

Akt hyperactivation, high proliferation, angiogenesis and immune-cell infiltration of tumors formed by NICP20 and NICP21 cells. To determine whether the in vivo characteristics of the NICP20 and NICP21 cells resemble the features of the original PTEN⁻/⁻/NIC tumors, serial immunohistological analyses were further performed. Examination of phospho-Akt staining revealed that tumors formed by NICP20 or NICP21 cells displayed hyperactivation of Akt (Fig. 5A). As shown in Fig. 5B, these tumors displayed a high proliferative level. Moreover, microvessel density reflected by CD34-positive endothelium (Fig. 5C) and CD3-positive cells (Fig. 5D) per area of tumor epithelium in the cell line-induced tumors also revealed comparable angiogenesis and T-cell infiltration properties similar to the in PTEN⁻/⁻/NIC tumors (15,17).

Discussion

Genetically engineered mouse models of human breast cancer have contributed substantially to our understanding of the initiation and progression of breast cancer and have emerged as valuable tools for preclinical research (22-24). Breast cancer is a complex and heterogeneous disease that has distinct histopathological features, genetic variability and diverse clinical outcomes. Overexpression of the ErbB2 oncogene and loss of tumor-suppressor PTEN are both observed in certain human breast cancer cases. The FVB/N PTEN⁻/⁻/NIC genetically engineered mouse was originally generated to better study the role of PTEN in ErbB2/Neu-induced mammary tumorigenesis (15). Notably, the PTEN⁻/⁻/NIC model developed multifocal, highly invasive mammary tumors with histopathological and molecular features relevant to the luminal subtype of primary human breast cancer (15). Our data confirmed the aggressive features of this model with an extremely short lifespan (Fig. 1).

To facilitate our understanding and ease the future application of this well-defined model, cell lines were generated from PTEN⁻/⁻/NIC mammary tumors in the present study. With the generation of the in vitro cell lines from PTEN⁻/⁻/NIC mammary tumors and development of the syngeneic xenograft model, an additional model system for the investigation of the interplay between ErbB-2/Neu and PTEN function/signaling in mammary tumor occurrence, progression and therapies is available, possessing the advantages of easy use, trouble-free monitoring of tumor growth, relatively inexpensive cost as well as the potential for in vitro manipulation. Similarly, Sahin et al reported that syngeneic transplantation of tumors from PTEN⁻/⁻/NIC mice is an effective way to generate large cohorts to test therapeutics (25). Thus, the availability of our cell line model and the syngeneic transplantation approach may provide tractable alternatives for optimizing PTEN⁻/⁻/NIC mice as powerful preclinical models.

Given that PTEN is a negative regulator of the PI3K/Akt pathway, its loss can lead to hyperactivation of Akt. As expected, the NICP20 and NICP21 cells had high levels of activated Akt
in vitro (Fig. 2B) and immunohistochemical staining also localized the activated Akt on tumor cells in vivo (Fig. 5A). Hyperactivation of Akt has been associated with trastuzumab resistance and Akt has been proposed as a potential target to overcome trastuzumab resistance (12,26-28). Novel inhibitors can be tested and evaluated using these NICP20 and NICP21 cells in vitro and in vivo in immunecompetent mice. In addition to inhibiting oncogenic signaling of tumor cells (29) (autonomous mechanism), studies have revealed that the immune response or stroma-tumor interactions in the tumor microenvironment (non-autonomous mechanism) are involved in the therapeutic activities of trastuzumab alone or in combination with other agents (17,30-32). Since the NICP20 and NICP21 cell-induced tumors showed comparable angiogenesis and T-cell infiltration properties as those of PTEN+/−/NIC tumors (Fig. 5), they may also be useful to investigate the roles of non-autonomous mechanisms on the therapeutic response of potential treatment, by working with syngeneic wild-type or related knockout mice.

In summary, we provide an additional valuable mouse mammary carcinoma cell resource to enhance our understanding of the nature of ErbB2-positive breast cancers, particularly accompanying PTEN loss, and to facilitate experimental therapeutic studies.

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