MTDH Mediates Estrogen-Independent Growth and Tamoxifen Resistance by Down-Regulating PTEN in MCF-7 Breast Cancer Cells

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Key Words
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Abstract
Background: About 70% of human breast cancers express estrogen receptor α (ERα) and in this kind of breast cancer estrogen plays an important role. Estrogen independent growth has been reported to promote resistance to one of the selective estrogen receptor modulators (SERMs) tamoxifen which is clinically the first line treatment for patients with ERα-positive breast cancer. The resistance of tamoxifen is a major problem in the clinical management of breast cancer. Methods: We used MCF-7 cells with ectopic expression of MTDH in this study. MTT, clone formation and tumor formation in nude mice methods were utilized to confirm the role of MTDH in estrogen-independent growth and tamoxifen resistance. Flow cytometry, western blot and siRNA were used to study the detailed mechanisms. Results: We found that MTDH could mediate estrogen-independent growth and induce resistance to tamoxifen in ERα-positive breast cancer cells. MTDH could reduce the expression of PTEN, up-regulate AKT and BCL2 and inhibit the apoptosis induced by tamoxifen. Conclusion: Our study indicated that MTDH was a candidate marker to predict the clinical efficacy of tamoxifen and targeting MTDH would overcome the resistance to tamoxifen in breast cancer cells.
Introduction

Breast cancer is the most common malignant tumor diagnosed in women worldwide and has become leading cause of cancer death in economically developing countries [1]. About 70% of breast cancers express high concentration of estrogen receptor (ERα) and candidates for endocrine therapy. Over the past few decades, tamoxifen, a selective ER modulator, has been the most commonly prescribed in endocrine therapy of ERα positive breast cancer and has been recommended as a preventative drug for female at high risk of developing breast cancer [2]. Tamoxifen which was used as an adjuvant drug has significantly improved the survival of early stage breast cancer patients. Nevertheless, nearly 35% of breast cancer patients fail to adjuvant tamoxifen therapy [3]. Tamoxifen resistance has been a major problem in the endocrine therapy of breast cancer. A large number of patients exhibit de novo tamoxifen resistance despite the presence of ERα in their breast tumors. Unfortunately, the molecular mechanisms of such resistance remain a poor understanding and serious clinical problems. It was considered that molecular crosstalk between ER and various growth factor signaling pathways might play important role in resistance to tamoxifen [4]. And a recent study reported that miR-375 and its target gene MTDH also worked in mechanisms of resistance to tamoxifen [5]. These studies provided research directions for understanding of resistance to tamoxifen. However it is still far from enough.

MTDH (metadherin, also known as AEG-1 and Lyric) is a novel oncogene found in 2002 [6]. Our previously studies show that it is widely over-expressed in many breast cancer cell lines or tumors and plays a crucial role in tumor invasion, metastasis and drug resistance [7-10]. Nearly 50% ER-positive breast cancer patient express high concentration of MTDH by immunohistochemical method and was significantly correlated with an aggressive phenotype and a poor prognosis [11]. In many ER-positive breast cancer cells estrogen (E₂) plays a crucial role [12]. But it hasn’t been reported whether E₂ is still important in tumor proliferation and progression in MTDH over-expressed ER-positive breast cancers. Otherwise the potential efficacy of MTDH in the treatment of tamoxifen in breast cancer has been unknown. In this study we over-expressed MTDH in ER-positive breast cancer cell line MCF-7 and found that MTDH could promote estrogen independent growth in MCF-7 cell line and further induce tamoxifen-resistance through inhibiting apoptosis by regulating PTEN/AKT pathway.

Materials and Methods

Reagents

The 17-β estradiol was purchased from Wako Pure Chemical Industries, Ltd. (Osaka, Japan), dissolved to a final concentration of 1 mM and stored at -20°C. Tamoxifen was purchased from Sigma-Aldrich (St. Louis, MO, USA). Estradiol pellet was purchased from Innovative Research of America. Rabbit polyclone antibody against MTDH was obtained from Invitrogen (Carlsbad, CA, USA). Antibodies against ERα, BCL-2 were purchased from Dako (Glostrup, Denmark). Rabbit antibodies against PTEN, p-AKT(Ser473), GAPDH and mouse anti-AKT were purchased from Cell Signaling Technology (Beverly, MA, USA). Mouse monoclonal antibody against β-actin was purchased from Sigma-Aldrich (St. Louis, MO, USA). The HRP-labeled secondary antibodies were from KPL (Gaithersburg, MD, USA).

Cell culture

The ERα negative cell lines MDA-MB-231, MDA-MB-468 and ERα positive cell lines MCF-7 and T47D were obtained from the American Type Culture Collection (Rockville, MA, USA) and grown in Dulbecco's Modified Eagle's Medium (DMEM, Gibco-BRL, Rockville, IN, USA) with 10% fetal bovine serum (FBS, Haoyang biological manufacture Co. Ltd, Tianjin, China), 100U/ml penicillin and 100μg/ml streptomycin at 37°C 5% CO₂ incubator. And 17 β-estradiol (E₂; Sigma-Aldrich, St. Louis, MO, USA) treatment cells were plated with phenol red-free DMEM medium supplemented with 5% charcoal-stripped FBS (Biochrom AG, Berlin, Germany) with or without E₂.
Vector construction and identification

The vector construction was performed as described previously [8]. Briefly, the cDNA representing the complete open reading frame of MTDH was cloned into the BamHI-XhoI vector fragment derived from pcDNA3.1/myc-His A vector (3.1n) (Invitrogen, USA) to generate pcDNA3.1-MTDH (3.1m). The expression plasmids were verified by sequencing of both strands.

Transfection and expression of MTDH

We transfected MCF-7 cells by using Lipofectamine TM 2000 (Invitrogen, Karlsruhe, Germany) according to the manufacturer’s instructions. The cells were seeded into 24-well culture plates (Biochrom AG, Berlin, Germany) at a density of 3×10^4 cells/well. The next day, cells were washed twice with PBS. Transfection mixes were added and cells were incubated for 4h at 37°C and 5% CO2. Subsequently, the transfection medium was replaced by complete medium. Cells were passaged at a 1:10 dilution into fresh growth medium 24h after transfection and the following day G418 was added at the concentration of 600μg/ml for selecting the stable cell lines. The selected cells were maintained in DMEM with 300 μg/ml G418. The efficiency of over-expression of MTDH in this cell line was tested by western blot analysis and real-time PCR (RT-PCR). Empty 3.1n vector was used as control. The stable selected cells transfected with 3.1n was called MCF7-3.1n while cells with 3.1m called MCF7-3.1m.

Real-time RT-PCR

Total RNA was isolated using Trizol (Invitrogen Corporate, Carlsbad, CA, USA) according to the manufacturer’s protocol. RNAs were reverse transcribed by using PrimeScript® RT reagent Kit (Takara, Dalian, China). The resulting cDNA was used for quantitative real-time PCR on a StepOne Plus instrument (Applied Biosystems, Carlsbad, CA, USA) with SYBR green (Takara, Dalian, China) detection. Primers involved were the MTDH primers forward ‘-AAATAGGCGATCTACGAGTCTGGAAGGAG-G3’ and the GAPDH primers (forward 5’-GGGCTGCTTTTAACTCTGGTAAAG-3’ , reverse 5’-CCATGGGTGGAATCATATTGG-3’). StepOne Plus software was used to calculate crossing threshold (Ct) points from the fluorescence curves and the delta/delta-Ct method was applied to quantify the induction of mRNA as compared to the control sample.

Cell proliferation assay

The cell proliferation with or without E2 and drug sensitivity to tamoxifen was measured by 3-(4,5-dimethylthiazol-2-yl)-2,5- diphenyltetrazolium bromide (MTT, Amresco, Solon, OH, USA). For estrogen dependent assay, cells were plated in 96-well culture plates at the concentration of 2000 cells/well. The next day, different concentrations of E2 were added into 100ul dilution of the culture medium. Cells were incubated for 96 hours at 37°C, 5% CO2. For estrogen independent growth, 2000 cells/well were plated in 96-well culture plates in phenol red-free DMEM medium supplemented with 5% charcoal-stripped FBS. The next day all wells were incubated in E2 free medium or 10^-9 M E2 medium for 0, 2, 4 and 6 days. For drug sensitivity assay, 3000cells/well were plated in 96-well culture plates. The next day different concentrations of tamoxifen from 0-25μM were added into dilution medium for 72 hours. Before harvesting, 20μl MTT were added to each well. Cells were incubated for another 4 hours and dimethyl sulfoxide (DMSO, Sangon Biotech, Shanghai, China) were added 100μl/well to dissolve the formazan after removing the culture medium. The plates were read at wavelength of 490 nm using Microplate Reader (Bio-Rad, Hercules, CA, USA). The culture medium was regarded as the bank control while the cells without reagents were the control sample.

Clone formation for Cytotoxicity assay

The drug sensitivity to tamoxifen was also measured by clone formation assay. Cells were plated in 6-well plates at a concentration of 500 cell/well. The next day different concentrations of tamoxifen from 0-500nM were added into 2.5ml dilution medium. The medium with drug was replaced every 5 days. After three weeks cells were fixed by methanol and stained with crystal violet. Clones containing over 50 cells were numbered under the microscope.
Tumor formation in nude mice

Tumor xenografts were developed by injecting 5×10^6 cells in a Matrigel Basement Membrane Matrix subcutaneously into each flank of 4-5 weeks-old BALB/c nu/nu female mice and analyzed as described elsewhere with the following modifications. The estrogen groups were implanted subcutaneously in the neck region with a 60 day release 0.72mg estradiol pellet (Innovative Research of America) and the minus estrogen groups were implanted with placebo pellet during the entire experiment. Tumor volume was calculated every 5 days. All the investigations have been approved by the local ethical committee.

Analysis of apoptosis by Flow Cytometry

Cells with or without treatment were detected by flow cytometry using the Annexin V-FITC Apoptosis Detection Kit (BD Bioscience, San Jose, CA, USA). The experiments were performed according to the protocol described. Cells were trypsinized, collected and washed twice with ice-cold PBS at the indicated time. Then 1×10^6 cells were resuspended in 100 μl 1×binding buffer, added with 5 μl of annexin V-FITC and 5 μl of PI following incubation for 15 min at room temperature in dark. Finally 400 μl 1×binding buffer was added to each sample. The samples were analyzed by flow cytometry (Becton Dickinson, Franklin Lakes, NJ, USA). Within 1 hour using FACS calibur with Cell Quest software. Cells that stained FITC Annexin V but not PI were identified as apoptotic cells.

Western Blot

Cells were washed with ice-cold PBS and lysed with ice-cold RIPA buffer (Shennengbocai, Shanghai, China) (1×PBS, 1%NP40, 0.1% sodium dodecyl sulfate (SDS), 5mM EDTA, 0.5% sodium deoxycholate) containing protein inhibitor cocktail (Sigma, St. Louis, MO, USA) and phosphatase inhibitor sodium fluoride (NaF) and sodium vanadate (NaVO3). Proteins from each group were quantified by BCA Protein Assay Kit (Merck, Darmstadt, Germany) and separated by 5%-10% SDS-PAGE and transferred onto PVDF membrane (Millipore, Schwalbach, Germany). Membranes were blocked in 1×TBST with 5% skim milk for 1 hour and subsequently incubated with the corresponding primary antibodies overnight at 4°C. Then the membranes were washed three times with TBST and the incubated with secondary antibody for 2 hours at room temperature. After washing 3 times in TBST, each protein was detected using an ECL system (Merck, Darmstadt, Germany).

siRNA and transfection

To knockdown the expression of PTEN in MCF7-3.1n cell line, we transfected the small interfering RNA (siRNA) of PTEN and Non-specific Negative control (NC, also obtained from Genepharma, Shanghai, China) with lipofectamine 2000 (Invitrogen) according to the manufacturer’s protocol. Transiently transfected cells were harvested after 48–72h for MTT and protein analysis.

Statistical analysis

All experiments were repeated at least three independent times. All data were presented as means±SD of the mean and analyzed using SPSS 13.0 for Windows (SPSS Inc., Chicago, IL, USA). Differences between individual groups were analyzed by student t test. P values of <0.05 were considered statistically significant.

Results

Over-expression of MTDH in breast cancer cell line MCF-7

We detected the expression of MTDH in a series of breast cancer cells including ERα negative cell lines MDA-MB-231, MDA-MB-468 and ERα positive cell lines MCF7 and T47D (Fig. 1A). To determine whether MTDH played an important role in estrogen dependence and tamoxifen sensitivity in breast cancer, we designed MTDH over-expression vector 3.1m to transfect MCF-7 cells and generated the MTDH over-expressed cells. The efficacy of MTDH expression in transfected MCF-7 cells was confirmed by RT-PCR on the RNA levels (Fig. 1B) while western blot on the protein levels (Fig. 1C). Cells transfected with 3.1m showed significantly increased MTDH expression on both mRNA and protein levels compared to the control 3.1n.
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**MTDH induced estrogen independent growth of cells**

To demonstrate if MTDH induced MCF-7 cells are less dependent on estrogen, we treated cells with different concentrations of E$_2$ for 96 hours and tested relative cell viability using cells in the phenol-free DMEM with CS-FBS as control. In contrast to MCF-7-3.1n, the MTDH over-expressed cells are less dependent on E$_2$ stimulation (p<0.001). The results showed that MCF7-3.1m cells could proliferate at the low dose of E$_2$ of 10$^{-13}$M contrast to that of 10$^{-9}$M in MCF7-3.1n cells (Fig. 2A).

To verify our finding that MTDH made estrogen independent growth in MCF-7 cells, we treated the two stable selected cell lines with E$_2$ at the concentration of 10$^{-9}$M and the...
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MTDH induced cells resistance to tamoxifen

As MTDH was indicated to play an important role in E₂ independent growth, we concluded that MTDH might regulate the sensitivity to the selective estrogen receptor modulator tamoxifen. To demonstrate it we used cell proliferation assay by MTT and clone formation in different concentrations of tamoxifen to measure cells sensitivity to tamoxifen in MCF7-3.1m and MCF7-3.1n. The results showed that the cells over-expressing MTDH were more resistant to tamoxifen (p=0.009) (Fig. 3A and 3B).

We further demonstrated the resistance induced by MTDH using testing apoptosis rate. After 24h treated with tamoxifen at the concentration of 10μM to MCF7-3.1n and MCF7-
3.1m cells, the apoptosis rates were tested using the untreated cells as control. The results showed the apoptosis rates of MCF-7-3.1m cells were lower than that of MCF-7-3.1n treated with tamoxifen (p=0.005) (Fig. 3C and 3D). These indicated that MTDH could inhibit cell apoptosis induced by tamoxifen.

MTDH mediated estrogen-independent growth and tamoxifen resistance by regulating PTEN/PI3K/AKT pathway

The crosstalk of ER-E2 and non-endocrine signals are often reported to induce tamoxifen resistance [13, 14]. Evidence indicated that estrogen independent growth and tamoxifen resistance are often associated with down-regulation of ERα [15]. We examined ERα expression and found that MCF-3.1m cells showed the same level of ERα as MCF-3.1n cells. Next, we determined the status of phosphorylated AKT (p-AKT, Ser473) because AKT activation has been implicated in tamoxifen resistance [16]. The results above showed that MTDH could increase expression of p-AKT and BCL2 which might contribute to tamoxifen resistance directly (Fig. 4A).

Phosphoinositide-3 kinase (PI3K)/AKT is a key factor in the mitogen-mediated cell growth pathway. Many factors are known to regulate p-AKT. The most direct and important upstream factor is PTEN (phosphatase and tensin homolog), a well known inhibitor of phosphoinositide-3 kinase. Inhibition of PTEN would increase the expression of p-AKT. Our results showed that MTDH over-expression down-regulated PTEN expression by western blot analysis (Fig. 4A).

To clarify the mechanism that MTDH mediated resistance to tamoxifen, we further silencing the expression of PTEN in MCF-7-3.1n cells. Then MTT assay and western blot were all performed with cells transfected with siRNA of PTEN (MCF-7-3.1n-siPTEN) and its negative control (MCF-7-3.1n-con). Knockdown of PTEN resulted in un-sensitivity to tamoxifen in MCF-7-3.1n-siPTEN cells. Results of western blot exhibited that the p-AKT levels was up-regulated (Fig. 4B). So we concluded that MTDH could lead to tamoxifen-resistance by suppression of PTEN.

We also confirmed the role of PTEN in regulation of resistance to tamoxifen. After silencing the expression of PTEN, we repeated the MTT assay to test the cell sensitivity to tamoxifen. The results indicated that when PTEN was down-regulated, MCF-7-3.1n-siPTEN cells were more resistant to tamoxifen than the control cells (Fig. 4C). All the results above...
showed that MTDH mediated estrogen-independent growth and tamoxifen resistance by regulating PTEN/PI3K/AKT pathway.

Discussion

Breast cancer is the most common malignant tumor diagnosed in women worldwide and has become leading cause of cancer death in economically developing countries [1]. Tamoxifen, which can target ERα and block the action of estrogen on breast cancer, is still a first-line endocrine therapy for the management of all stages of ERα-positive breast cancer. Five years of tamoxifen use was considered the standard duration, however, 10 years treatment is recommended recently. It can reduce the risk of recurrence by 41% on average in patients with ERα-positive tumors [17, 18]. It was improved to greatly prolong disease-free survival and induce remission in over half of patients with ER-positive breast cancer [19]. Unfortunately, almost 50% of patients with advanced disease do not respond to first line treatment with tamoxifen. Furthermore, a significant percentage of patients experience tamoxifen resistance relapse, despite an initial positive drug response [18, 20]. The mechanisms of tumor resistance to tamoxifen therapy continue to pose a significant challenge to both clinicians and researchers. It might reside in the expression of specific molecules involved in different signaling pathways, which eventually could be used as predictive biomarkers of resistance. Moreover, these markers may be used to select patients might benefit from additional targeted treatments aside from ERα [21, 22].

As found in 2002, MTDH has been found over-expressed in multiple cancers such as esophageal squamous cell carcinoma, breast carcinoma, melanoma, hepatocellular carcinoma, epithelial ovarian cancer and so on [23-25]. Many experiments have been done to demonstrate MTDH as an oncogene to regulate multiple molecular and pathways including PI3K/AKT pathway, NF-kappaB pathway, MAPK pathway, Wnt pathway, vascular endothelial growth factor (VEGF), transcription factor family (such as FOXO1, FOXO3a) and so on to promote cell proliferation, migration, tumor metastasis and angiogenesis in various tumors [7, 25-30]. In breast cancer our previous studies have shown that MTDH was over-expressed in more than 40% of the tumors which could indicate poor clinical outcomes [7, 9] and MTDH could enhance the invasiveness of cells by inducing epithelial to mesenchymal transition (EMT) [8]. Besides these we found that MTDH could induce multiple drug resistance in breast cancer such as doxorubicin, paclitaxel and cisplatin [31]. However there is still little report about the role of MTDH in estrogen dependence and tamoxifen resistance. In this study we found the functions of MTDH to promote estrogen independent growth and induce tamoxifen resistance.

Estrogen (E\textsubscript{2}) is demonstrated to be an endogenous factor which plays a critical role in normal mammary functions [32]. Studies in human breast cancer cells subjected to long-term estrogen deprivation (LTED) have demonstrated that these cells first developed hypersensitivity to low-dose estrogens and then became estrogen independent. This phenomenon seemed to contribute to resistance to endocrine therapy, which might in part result by increased levels of ERα and up-regulation of the MAPK, PI-3-kinase and mTOR growth factor pathways [33]. In our studies we first found that the oncogene MTDH could promote MCF-7 cells more sensitivity to E\textsubscript{2} stimulating. Our results showed the two cell lines, MTDH over-expressed cells MCF7-3.1m and the control cells MCF7-3.1n, responded to E\textsubscript{2} stimulation. The MCF7-3.1m cells could respond to E\textsubscript{2} at a very low dose of 10^{-11}M. MTDH could promote cell proliferation in MCF-7 much more obviously than control cell lines in E\textsubscript{2} free medium. These results indicated that MTDH might induce estrogen independent growth. We further demonstrated this by xenograft tumor formation in nude mice. Tumors were smaller in nude mice injected with MCF7-3.1n cells without estrogen pellet while significantly larger with MCF7-3.1m cells. Our studies suggested MTDH could promote MCF-7 cells estrogen independent growth. Studies have indicated that ER positive breast cancer cells with estrogen independent growth were resistance to tamoxifen therapy [34,
We further tested the sensitivity to tamoxifen treatment of these two cell lines MCF7-3.1n and F7-3.1m by MTT assay. We found that MTDH over-expressed cell lines expressed resistance to tamoxifen treatment contrast to control cell lines. Tumor cell apoptosis plays an important role in both preclinical and clinical responses to tamoxifen and tumor cell evasion of apoptosis contributes to tamoxifen resistance [36, 37]. To determine whether MTDH promoted tamoxifen resistance though reducing the apoptosis after treated with tamoxifen, we found that the increased apoptosis rate induced by tamoxifen in MCF7-3.1m was lower than that in MCF7-3.1n.

As down-regulation of ERα is often associated with estrogen independent growth and tamoxifen resistance, we examined ERα expression. As shown in Fig. 4, MCF7-3.1m cells revealed little decrease at the level of ERα than MCF7-3.1n cells, suggesting that MTDH impacted tamoxifen sensitivity though other signaling pathway. Next we determined the status of phosphorylated Akt (p-AKT) because Akt activation has been implicated in estrogen independent growth and tamoxifen resistance [16]. Western blot analysis revealed that MCF7-3.1m cells expressed high level of p-AKT than MCF7-3.1n cells.

Phosphoinositide-3 kinase /Akt is a key factor in the cell growth and proliferation pathways which regulated by many factors. PTEN (phosphatase and tensin homolog), a well known inhibitor of phosphoinositide-3 kinase, is the most important and direct upstream factor. Reduction of its activity or suppression of PTEN expression would increase p-Akt levels. Down-regulation of PTEN expression in ERα-positive tumors is associated with failure to tamoxifen treatment [38]. Our study found that MTDH over-expression down-regulated PTEN expression by 38%. This suggested that MTDH induced tamoxifen resistance through inhibiting PTEN expression and activation of p-AKT. To further clarify the mechanism that MTDH mediate resistance to tamoxifen, we silenced the expression of PTEN in MCF-3.1n cells. The results showed that silencing of PTEN resulted in p-AKT over-expression by western blot and tamoxifen resistance by MTT assay. We failed to detect altered ERα and BCL-2 expression in response to suppression of PTEN, which suggests MTDH contributes to tamoxifen resistance through multiple mechanisms.

Based on our study, MTDH could mediate estrogen-independent growth and induce resistance to tamoxifen in ERα-positive breast cancer cells. MTDH could reduce the expression of PTEN, up-regulate AKT and BCL2 and inhibit the apoptosis induced by tamoxifen. PTEN contributes to MTDH induced tamoxifen resistance in breast cancer, however, MTDH contributes to tamoxifen resistance through multiple mechanisms. Our findings provided the key functions of MTDH in tamoxifen resistance of breast cancer and indicated that MTDH could be a convincing target to reverse the resistance to tamoxifen in ERα-positive breast cancer patients.

Disclosure Statement

Authors declare no conflict of interest.

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