



Original articles

Micro ribonucleic acid-448 regulates zinc finger e-box binding homeobox 1 to inhibit the growth of breast cancer cells and increase their sensitivity to chemotherapy

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HIGHLIGHTS

- MicroRNA-448 expression decreased in Breast Cancer (BC) tissues and BC cells.
- Zinc finger E-box Binding Homeobox 1 (ZEB1) was upregulated in BC tissues and cells.
- Upregulation of miR-448 expression could inhibit the malignant behaviors of BC cells.
- Upregulated miR-448 expression could enhance BC cells' sensitivity to PTX or 5-FU.

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ABSTRACT

Objective: This study aimed to investigate the effect of Zinc Finger E-box Binding Homeobox 1 (ZEB1) regulation by Micro Ribonucleic acid (miR)-448 on Breast Cancer (BC) cells and their sensitivity to chemotherapy.

Methods: miR-448 and ZEB1 mRNA levels in BC and normal tissues were detected by qPCR, and ZEB1 protein was detected by Western Blotting (WB). The correlation between miR-448 and tumor metastasis, clinical staging, and ZEB1 expression was analyzed. MCF-7 cells were transfected or co-transfected with the miR-448 mimic, oe-ZEB1, or their negative controls. Changes in miR-448 and ZEB1 expression were detected by qPCR and WB. Cell proliferation was determined by CCK-8 assays, invasion changes were analyzed by Transwell assays, and apoptosis was detected by flow cytometry.

Results: miR-448 expression in BC tissues was lower than that in normal tissues, while ZEB1 expression was increased in the former. ZEB1 expression was lower in BC patients with lymph node metastasis than in those without. In patients with clinical stage I–III BC, miR-448 expression decreased with an increase in tumor stage, which was negatively correlated with ZEB1 expression. Upregulation of miR-448 expression can suppress MCF-7 cell proliferation and invasion and promote apoptosis. Upregulation of ZEB1 expression in cells overexpressing miR-448 can partially reverse the inhibition of BC cell growth induced by miR-448. miR-448 can enhance the sensitivity of cells toward paclitaxel and 5-fluorouracil.

Conclusions: miR-448 suppresses cell proliferation and invasion and promotes apoptosis by targeting ZEB1. Moreover, it can increase the sensitivity of cells toward paclitaxel and 5-fluorouracil.

Introduction

Breast Cancer (BC) is the most common malignancy in women.¹ In China, BC-associated morbidity is high, while BC-related mortality ranks third among tumor-related diseases.² The disease develops and progresses through polygenic, multifactorial, and multi-step complex processes; these processes interact and are mutually regulated.³ Although the incidence of BC continues to increase, the mortality rate associated with it is declining.⁴ This may be largely due to the systematic

development of treatments, including chemotherapy,^{5,6} hormone therapy,⁷ radiotherapy,⁸ and targeted therapy.⁹ Because of the high cost of targeted therapy, chemotherapy is still the first choice of treatment in many developing and underdeveloped countries. However, drug resistance is the main cause of chemotherapy failure.¹⁰ Other factors, such as metastasis, are also responsible for chemotherapy failure; in addition, the long-term survival rate of BC patients is not ideal.¹¹ The discovery period is relevant to the prognosis.¹² Clinical studies have confirmed that the 5-year survival rate of early-stage BC patients after treatment is

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as high as 70%–90%, while for patients with advanced-stage BC, the rate can be as low as < 15% due to the proliferation or metastasis of cancer cells.¹³ Therefore, there is an urgent need to develop a new strategy to treat BC to lower the dosage and side effects of chemotherapy drugs and improve their efficacy.

Many studies have confirmed that miRNA expression is relevant to disease development and progression. Many studies have found that there are marked differences in miRNA expression between normal and cancerous cells, and most of these differences are related to the gene loci of cancer cells.^{14–16} In addition, miRNA is involved in the process of tumor formation and related drug resistance, which affects the growth of tumor cells and the treatment.¹⁷ As potential targets, various miRNA have been found to be involved in the sensitivity of BC chemotherapy. For example, it has been found that the miR-326 expression in BC MCF-7 cells resistant to etoposide is down-regulated, and increasing the expression by cell transfection can increase the sensitivity of drug-resistant cells to adriamycin and etoposide.¹⁷ Another study found that over-expressing miR-451 in BC cell lines can reduce the drug resistance of adriamycin, which leads to the enhancement of cell proliferation and invasion.¹⁸ In the present study, by overexpressing miR-448, changes in the sensitivity of MCF-7 BC cells to chemotherapeutic drugs were observed, and a novel target for chemosensitivity enhancement was discovered.

Materials and methods

Clinical tissues and cell lines

From January 2014 to January 2015, cancer tissues and adjacent normal tissues were collected from 38 primary BC patients admitted to the Breast Surgery Department of our hospital. Among them, one patient had invasive lobular carcinoma, while 37 had invasive ductal carcinoma. There were 25 patients with lymph node metastasis and 13 without. BC staging is based on the standards established by the American Joint Committee on Cancer. In this study, 9 patients were at stage I, 15 at stage II, 11 at stage III, and 3 at stage IV of BC. The collection of all tissue samples was approved by the ethics committee of the First Affiliated Hospital of Soochow University. All study participants provided written informed consent before participating in the study. None of the patients had a history of radiotherapy or chemotherapy before surgery. After fresh tissues were excised during surgery, they were stored in liquid nitrogen for transfer to the laboratory. The patients were followed up for 5 years. Human BC cell lines MDA-MB-231, MCF-7, T47D, DK-BR-3, and breast epithelial cell MCF-10A were provided by the Shanghai Institute of Cell Research, Chinese Academy of Sciences. Those were cultivated in RPMI1640 with 10% FBS at 37°C, 5% CO₂, and a saturated humidity incubator.

qPCR

Total RNA was separated by TRIzol reagent (Takara), and cDNA was synthesized with miScript II RT Kit. The miScript SYBR Green PCR Kit (Qiagen) was applied in real-time quantitative PCR. PCR primers were designed and synthesized by Make Research Easy Co., Ltd. The following primers were used: miR-448, F: 5'-TTATTGCGATGTTCCTTATG-3' and R: 5'-ATGCATGCCACGGGCATATACACT-3'; U6, F: 5'-CTCGCTTCGGCAGCACA-3' and R: 5'-AAGCCTTCACGAATTTGCGT-3'; ZEB1, F: 5'-GCCAATAAGCAAACGATTCTG-3' and R: 5'-TTGGCTGGATCACTTCAAG-3'; GAPDH: F: 5'-CAAGGTCATCCATGACAACCTTG-3' and R: 5'-GTCCACCACCTGTTGTGTAG-3'. U6 was used as an internal reference for miR-448, GAPDH was used as an internal reference for ZEB1, and the 2^{-ΔΔCt} method was employed for quantitative analysis. All reactions were performed with three negative controls with multiple holes and no template.

Cell transfection

The logarithmic growth phase cells were inoculated into a 6-hole plate, and miR-448 mimic and its negative control, miR-448 mimic + oe-ZEB1, mimic NC + oe-NC, and miR-448 mimic + oe-NC were transfected into MDA-MB-231 and MCF-7 cells via Lipofectamine 2000 kit.

WB

Total proteins were extracted from the tissues and cells. The concentration of each sample was measured and adjusted with deionized water to ensure the same loading amount. Next, 10% SDS separation and concentrated gels were collected. The samples were blended with sample buffer, boiled at 100°C for 5 min, cooled in an ice bath, centrifuged, and then added to each lane with a micro-sampler for electrophoresis separation. The gel protein was transferred to the nitrocellulose membrane. Afterward, the cellulose nitrate membrane was closed by 5% skimmed milk powder at 4°C all night. The primary antibody was added: ZEB1 and GAPDH (1:1000, Proteintech) were incubated overnight and washed with PBS (Phosphate Buffer) 3 times at indoor temperature, each time for 5 min. The second antibody of HRP labeled IgG (1:1000, Boster Biological Technology Co. Ltd) was incubated 1h at 37°C. At indoor temperature, PBS buffer solution was cleaned 3 times, 5 min each time. And the membrane was immersed for 1 min in an ECL reaction solution (Pierce, USA). The liquid was removed, closed by cling film, and exposed with X-Ray film in a dark environment. The results were observed after developing and fixing. GAPDH was considered as an internal reference, and the protein imprinted images were analyzed by ImageJ2x software.

CCK-8 assay

The transfected cells were collected and prepared into a 2 × 10⁴ cells/mL single-cell suspension. The suspension (100 μL/well) was inoculated into the 96-hole plate. Then, the plate was cultivated at 37°C, with 5 multiple wells in each group. Subsequently, 10 μL of CCK-8 solution was added 24h, 48h, 72h, and 96h later, and special attention was paid to avoid bubble formation during sample addition. The culture plates were incubated for 2h at 37°C. Cell absorbance was measured at 450 nm using a microplate reader. The test was conducted three times, and the data were recorded.

Drug experiment: The cells were collected and inoculated into a 96-well plate. Each group had three wells, and a blank control group was also used. Paclitaxel (PTX) or 5-Fluorouracil (5-FU) at final concentrations of 2.5, 5.0, 7.5, 10.0, and 12.5 ng/mL or 10, 20, 30, 40, and 50 μg/mL, respectively, were added after cell attachment. Seventy-two hours after cell treatment, the culture medium was discarded, and 10% CCK-8 solution was supplemented to cultivate for 2h. The absorbance value of magnetic nails at 450 nm of the microplate reader was measured. The IC50 growth inhibition rate of each group = [(average absorbance value of control hole - that of an experimental hole) / that of control hole] × 100%.

Plate cloning

The cells of each group were inoculated into a 6-well plate in a complete medium containing 10% FBS. Next, they were incubated at 37°C and 5% CO₂ in a saturated humidity incubator for 2 weeks. When visible clones appeared on the culture dish, the supernatant was discarded, cleaned twice with PBS, and immobilized for 15 min with 4% paraformaldehyde. After the cells were rinsed carefully with PBS twice, they were dyed with an appropriate amount of crystal violet staining solution (Wuhan Google Biological Co., Ltd.) for 10 min, following which the solution was washed off slowly with running water. After the plate was dried, the clone number was counted directly with the naked eye.

Transwell assay

The upper surface of the Transwell chamber basement membrane was coated with Matrigel matrix glue diluted with serum-free cell culture solution, and each group of cell suspension, which was diluted with 100 μ L of serum-free culture medium (approximately 2×10^5 cells), was added. The complete culture solution with 10% serum was supplemented to the lower orifice plate of the chamber. The culture chamber was taken out after 24-hour culture at 37°C. The matrix glue and cells were cleaned using wet cotton swabs and immobilized with 4% paraformaldehyde.

Apoptosis

The cells from each group were collected. Apoptosis was analyzed using a fluorescein isothiocyanate/Propidium Iodide (PI) kit (BD Biosciences, San Jose, CA, USA) and flow cytometry.

Dual-luciferase reporter gene assay

The miR-448 and ZEB1 binding sites were predicted using the miRNA database and identified by a dual-luciferase reporter gene assay. The ZEB1 3'-UTR was synthesized and cloned into a plasmid in the psiCHECK reporter (Promega) and verified by DNA sequencing and named Wild-Type ZEB1 (WT-ZEB1). Then, a 3'-UTR Mutant (MUT-ZEB1) was constructed and synthesized through site-directed mutagenesis. MDA-MB-231 and MCF-7 cells were co-transfected with recombinant reporter gene WT, MUT plasmid, and mimic NC, or miR-448 mimic in a 24-well

plate. The activity was tested after 48h using a dual-luciferase reporter assay (Promega).

Statistical methods

The results were assessed using SPSS 22.0, and the measurement data are represented as the mean \pm SD. The differences between the two groups were evaluated using the independent samples *t*-test, while differences between multiple groups were assessed using one-way ANOVA. Both groups were examined using Tukey's multiple comparison test. The prognosis of patients with cholangiocarcinoma was examined using Kaplan-Meier analysis. The difference was considered statistically significant at $p < 0.05$.

Results

miR-448 and ZEB1 levels in BC

qPCR and WB analysis revealed that miR-448 expression decreased and ZEB1 levels increased in BC tissues ($p < 0.0001$, Fig. 1A–1D). Moreover, miR-448 expression was lower in BC patients with lymph node metastasis than in those without ($p < 0.0001$, Fig. 1E). For patients at different stages, the expression of miR-448 decreased as the stage increased ($p < 0.05$, Fig. 1F), but stages III and IV showed no obvious differences ($p < 0.05$). The miR-448 level was negatively correlated with that of ZEB1, suggesting the potential functional relevance between ZEB1 and miR-448 (Fig. 1G). In view of the median level of miR-448, patients were randomized into high and low expression groups. miR-448 level's influence on BC patients' survival and prognosis was assessed

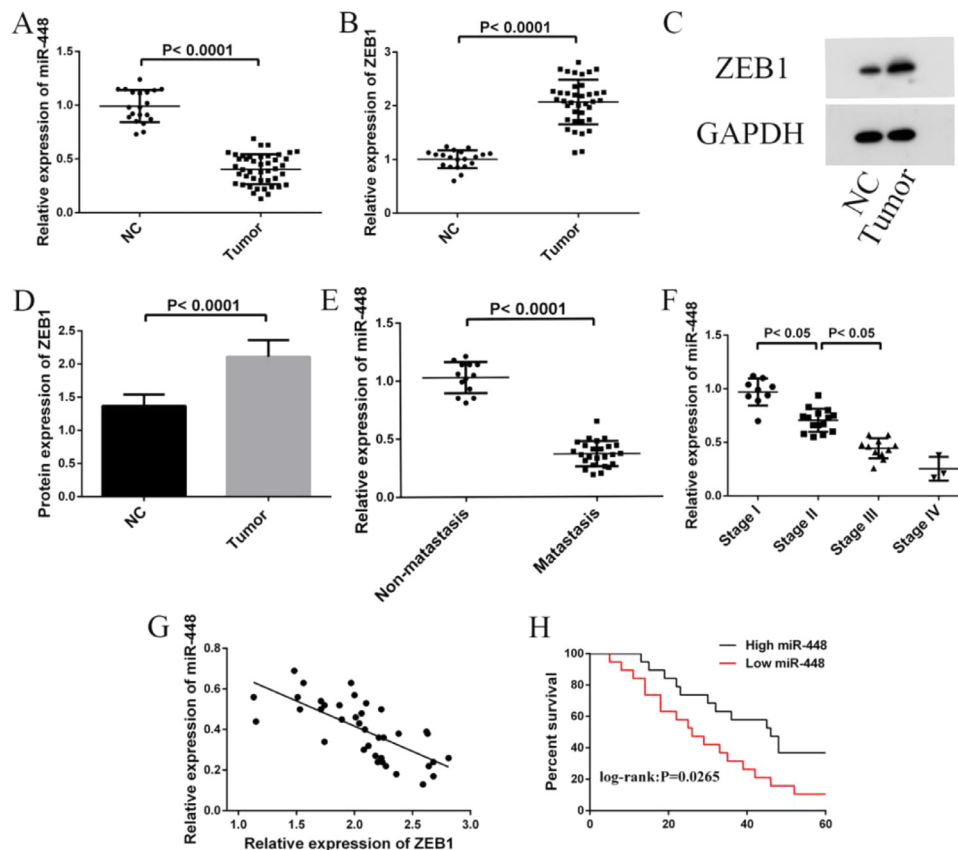


Fig. 1. miR-448 and ZEB1 expression in BC. (A) Expression of miR-448 in BC; (B) Expression of ZEB1 mRNA in BC; (C) ZEB1 protein expression; (D) Expression of ZEB1 protein in BC; (E) Expression of miR-448 in patients with or without lymph node metastasis; (F) miR-448 expression in BC patients at different stages; (G) Correlation between miR-448 and ZEB1 expression; Correlation was determined using Spearman analysis. and (H) 5-year survival rate of patients with high and low expression of miR-448.

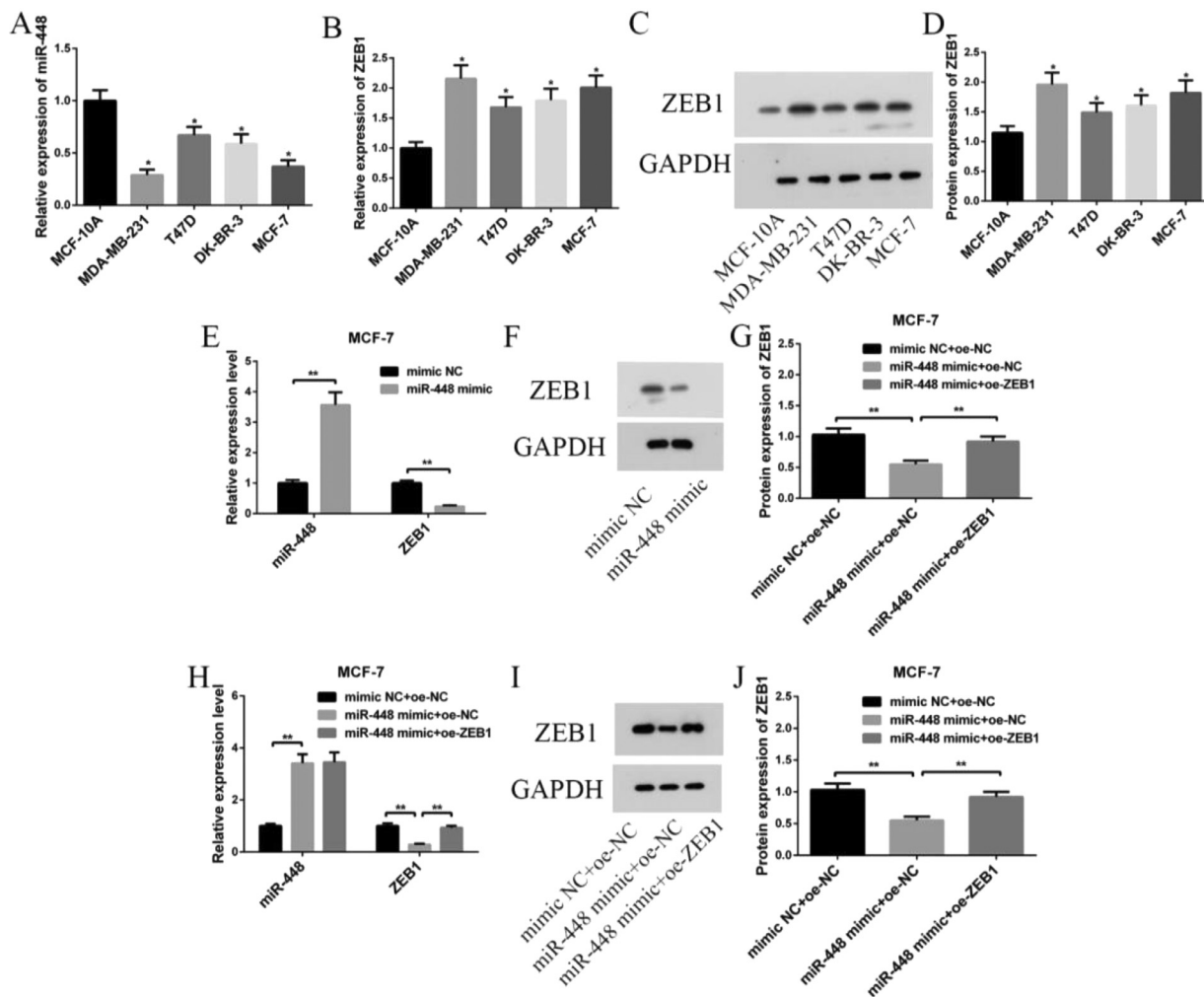


Fig. 2. miR-448 and ZEB1 expression in each cell line. (A) Expression of miR-448 in each cell line; (B) Expression of ZEB1 mRNA in each cell line; (C) ZEB1 protein band map in each cell line; (D) Expression of ZEB1 protein in various cell lines; (E) Expression of miR-448 and ZEB1 mRNA in MCF-7 cells after transfection; (F) ZEB1 protein banding pattern in transfected MCF-7 cells; (G) Expression of ZEB1 protein in MCF-7 cells after transfection; (H) Expression of miR-448 and ZEB1 mRNA in MCF-7 cells after co-transfection; (I) ZEB1 protein banding pattern in MCF-7 cells after co-transfection; (J) Expression of ZEB1 protein in MCF-7 cells after co-transfection; * Compared with MCF-10A cells, $p < 0.05$; ** $p < 0.01$.

via Kaplan-Meier. It manifested that the 5-year survival rate of those in the low-expression group was lower ($p < 0.05$, Fig. 1H).

miR-448 and ZEB1 levels in cells

Compared with human breast epithelial cell MCF-10A, the miR-448 expression in all human BC cells decreased, while the ZEB1 mRNA and protein level increased ($p < 0.05$, Fig. 2A–2D). MCF-7 cells were selected as subjects. Compared with mimic NC and mimic NC + oe-NC groups, the miR-448 expression in miR-448 mimic and miR-448 mimic + oe-NC groups increased, while that of ZEB1 mRNA and protein decreased ($p < 0.01$, Fig. 2E–2G). Compared with miR-448 mimic + oe-NC group, the miR-448 level in miR-448 mimic + oe-ZEB1 group had no obvious difference, but that of ZEB1 mRNA and protein increased ($p < 0.01$, Fig. 2H–2J).

Effect of miR-448 expression on the biological behavior of BC cells

The possible binding sites of miR-448 and ZEB1 were predicted using an online biological prediction tool and were verified using a dual-luciferase reporter gene assay. The results showed that when WT-ZEB1 and miR-448 were co-transfected into MCF-7 cells, the fluorescein activity

decreased ($p < 0.05$, Fig. 3A–3B), while this activity followed others, co-transfections did not change. Compared with the mimic NC group, cell proliferation activity ($p < 0.01$, Fig. 3C) and a number of clonal ($p < 0.01$, Fig. 3D) and invasive cells in the mimic group transfected with miR-448 was reduced ($p < 0.01$, Fig. 3E), and the apoptosis rate increased ($p < 0.01$, Fig. 3F) (Fig. 3).

Effects of miR-448 and ZEB1 expression on the biological behavior of BC cells

Compared with the mimic NC + oe-NC group, the cell proliferation activity, the number of plate clonal and invasive cells in the miR-448 mimic + oe-NC group reduced ($p < 0.05$, Fig. 4A), and the apoptosis rate rose ($p < 0.01$, Fig. 4B). Conversely, compared with miR-448 mimic + oe-NC group, the cell proliferation activity of co-transfected miR-448 mimic + oe-ZEB1 group increased ($p < 0.05$, Fig. 4C), and the number of clonal and invasive cells increased, while the apoptosis rate decreased ($p < 0.01$) (Fig. 4, Fig. 4D).

Influence of miR-448 on the sensitivity of BC cells to PTX and 5-FU

Different concentrations of PTX and 5-FU had different effects on miR-448 transfection efficiency. The results showed that the optimal

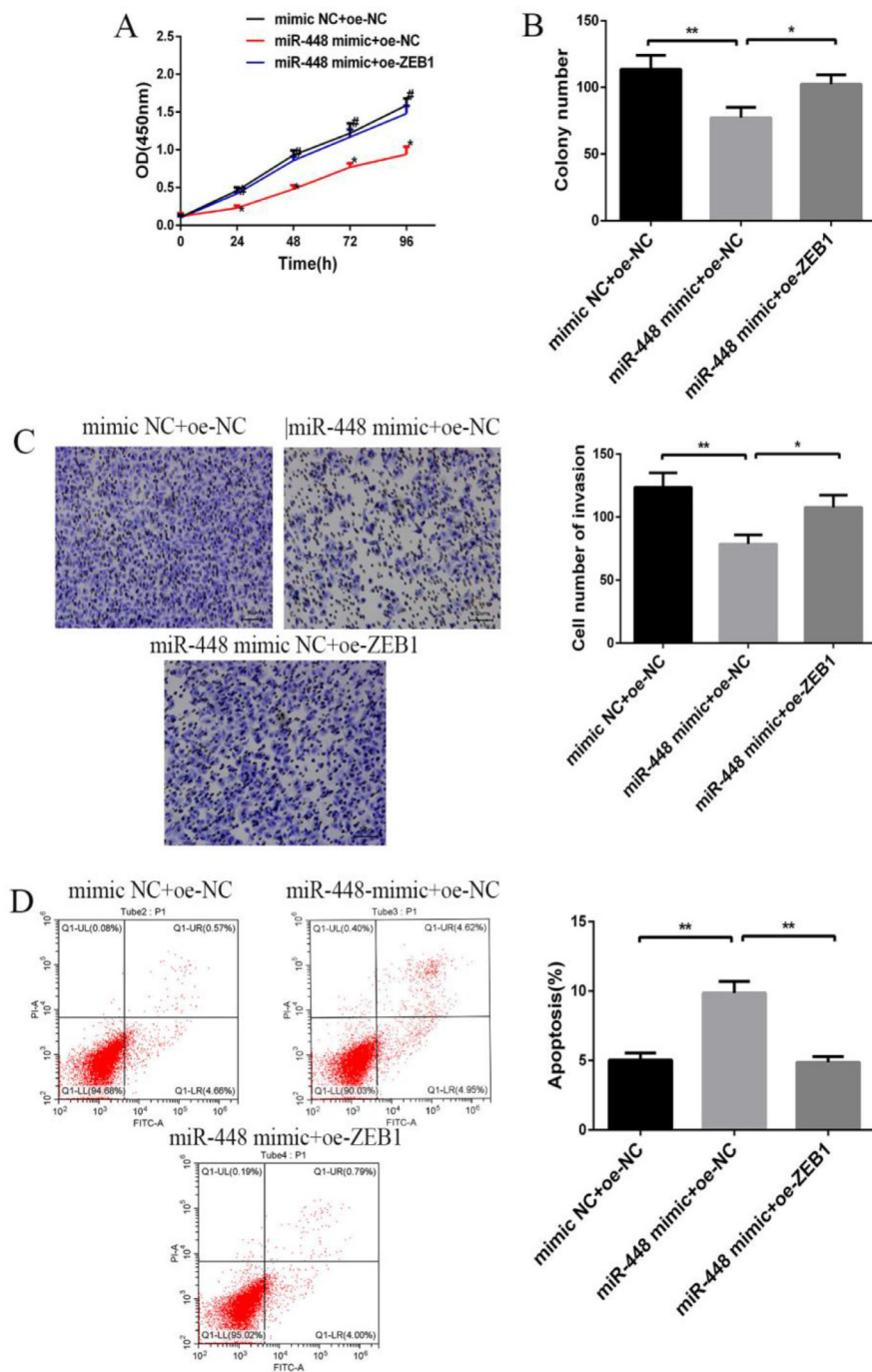


Fig. 4. Effects of miR-448 and ZEB1 expression on the biological behavior of BC cells. (A) Effects of miR-448 and ZEB1 on BC cell proliferation; (B) Effects of miR-448 and ZEB1 on plate clone number of BC cells; (C) Effects of miR-448 and ZEB1 on BC cell invasion; (D) Effects of miR-448 and ZEB1 on the apoptosis of BC cells; * Compared with the control group, # compared with the miR-448 mimic + oe-NC group, p < 0.05; ** p < 0.01.

448 expression decreased with an increase in tumor stage, which was negatively correlated with ZEB1 expression. Some studies have found that miR-448 expression is downregulated in many cancers, such as colon³⁶ and lung cancers.³³ miR-448 expression is also downregulated during EMT induced by BC chemotherapy.³⁴ A previous study reported that in BC, ZEB1 expression could reduce the expression of inflammatory cytokines,³⁵ which is similar to the findings of the present study. Upregulation of miR-448 expression can suppress MCF-7 cell proliferation and invasion and promote apoptosis. Upregulation of ZEB1 expression in cells overexpressing miR-448 can partially reverse miR-448-induced inhibition of BC cell growth. Peng et al.³⁷ found that ectopic miR-448 expression suppresses cancer cell migration and invasion by regulating the EMT. Their results were verified in the present study. miRNAs regulate the sensitivity of cancer cells toward PTX and 5-FU. Mao et al.³⁸ and

Li et al.³⁹ confirmed that PTX could increase the ratio of CD44⁺/CD24⁻ cells in BC. Collectively, the findings of these studies suggest that the enrichment of cancer stem cells induced by PTX may be a vital reason for the decrease in drug sensitivity. It has been reported that miRNAs, such as miR-1204,⁴⁰ miR-125b,⁴¹ and miR-634,⁴² can increase sensitivity to PTX. It has also been reported that miRNAs, such as miR-489⁴³ and miR-210,⁴⁴ can increase the drug sensitivity of 5-FU. The current study demonstrated that miR-448 could enhance the sensitivity of BC cells toward PTX and 5-FU.

However, there are a few limitations to the study. First, there may exist other molecules in micro ribonucleic acid-448 that regulates zinc finger e-box binding homeobox 1 to inhibit the growth of breast cancer cells and increase their sensitivity to chemotherapy which may have the same effects. Second, breast cancer cells animal models must be created

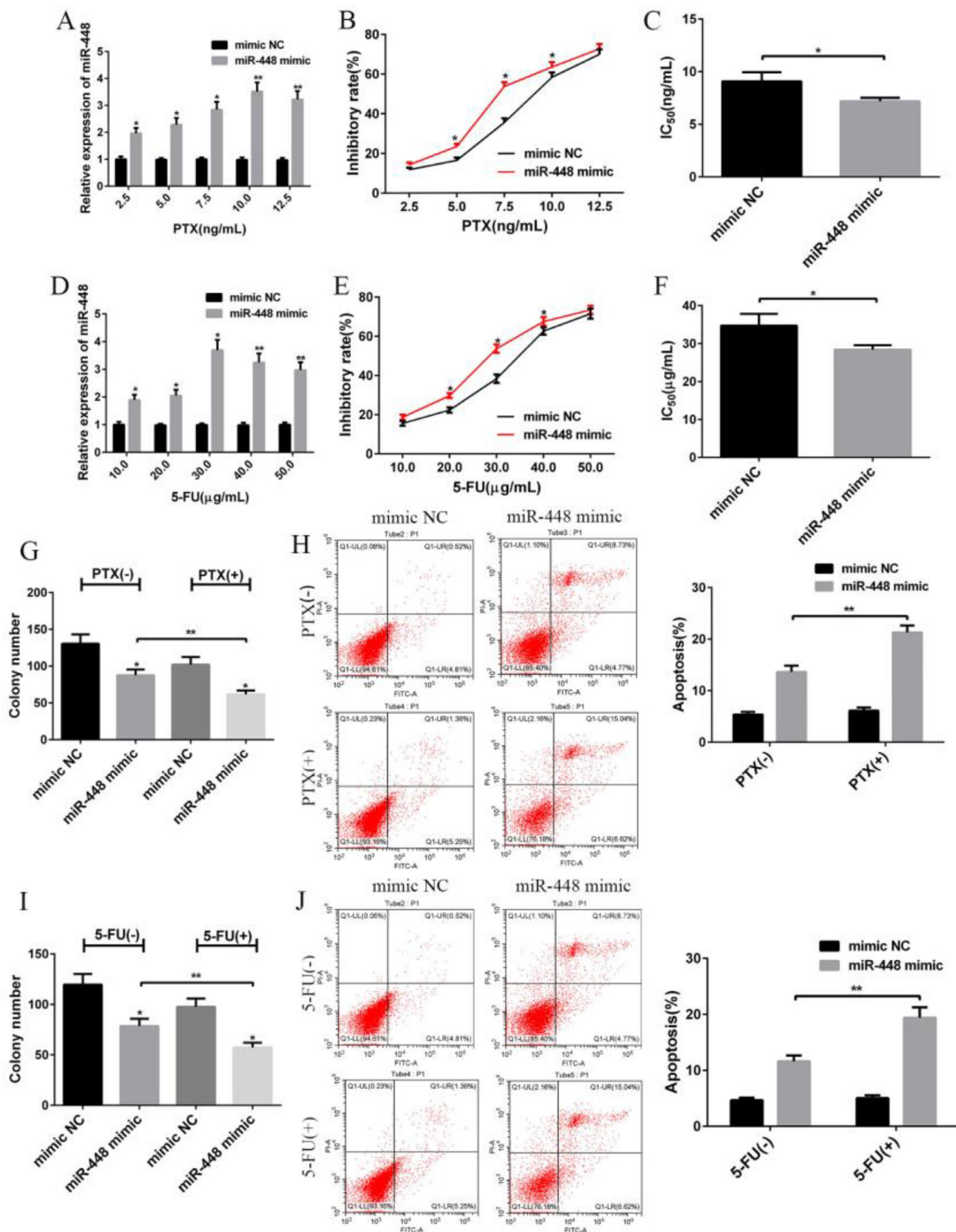


Fig. 5. Influence of miR-448 on the sensitivity of BC cells to paclitaxel and 5-fluorouracil. (A) Comparison of transfection efficiency of miR-448 with different concentrations of paclitaxel; (B) Growth inhibition rates of cells following treatment with different concentrations of paclitaxel; (C) IC50 value of paclitaxel in each group of cells; (D) Comparison of transfection efficiency of miR-448 with different concentrations of 5-fluorouracil; (E) Growth inhibition rates of cells following treatment with different concentrations of 5-fluorouracil; (F) IC50 value of 5-fluorouracil in cells of each group; (G) miR-448 enhances paclitaxel-induced inhibition of MCF-7 cell proliferation; (H) miR-448 enhances the apoptosis-promoting effects of paclitaxel on MCF-7 cells; (I) miR-448 enhances 5-fluorouracil-induced inhibition of MCF-7 cell proliferation; (J) miR-448 enhances MCF-7 cell apoptosis promoted by 5-fluorouracil; * Compared with the control group, $p < 0.05$; ** $p < 0.01$.

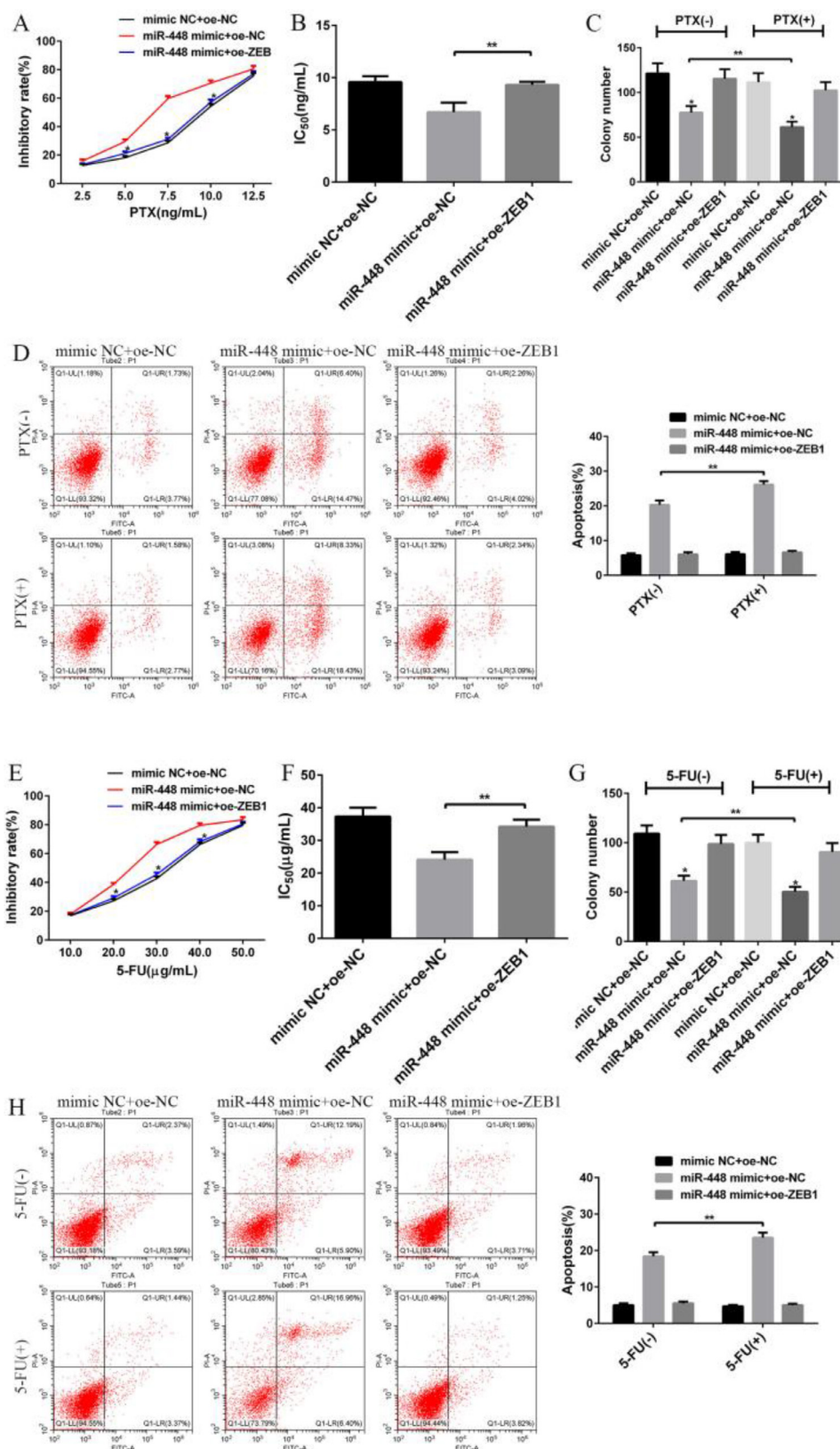


Fig. 6. Influence of miR-448 and ZEB1 on the sensitivity of BC cells to paclitaxel and 5-fluorouracil. (A) Growth inhibition rates of cells from each group following treatment with different concentrations of paclitaxel; (B) IC₅₀ value of paclitaxel in cells of each group; (C) Effects of miR-448 and ZEB1 combined with paclitaxel on the proliferation of MCF-7 cells; (D) Effects of miR-448 and ZEB1 combined with paclitaxel on the apoptosis of MCF-7 cells; (E) Growth inhibition rates of cells from each group following treatment with different concentrations of 5-fluorouracil; (F) IC₅₀ value of 5-fluorouracil in cells of each group; (G) Effects of miR-448 and ZEB1 combined with 5-fluorouracil on the proliferation of MCF-7 cells; (H) Effects of miR-448 and ZEB1 combined with 5-fluorouracil on MCF-7 cell apoptosis; * Compared with the control group, $p < 0.05$; ** $p < 0.01$.

to verify the present research findings in vitro. Third, more clinical data needs to be collected to confirm the importance of micro ribonucleic acid-448.

In summary, miR-448 expression decreased in BC tissues. Upregulation of miR-448 expression inhibits cancer cell proliferation and invasion, promotes apoptosis by targeting ZEB1, and increases the sensitivity of the cells to PTX or 5-FU. This indicates that upregulation of miR-448 expression is an effective method to enhance the efficacy of BC treatment.

Authors' contributions

Yizhou Zhou, Li Sun, Yangmei Zhang and Kai Chen conceived the project, designed, and performed the experiments, analyzed the data. Yizhou Zhou and Kai Chen wrote and revised the manuscript.

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Declaration of Competing Interest

The authors declare no conflicts of interest.

Acknowledgments

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