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Abstract. Doxorubicin (Dox) is widely used in the treatment of triple-negative breast cancer cells (TNBCs), however resistance limits its effectiveness. Cancer stem cells (CSCs) are associated with Dox resistance in MCF-7 estrogen receptor positive breast cancer cells. Signal transducer and activator of transcription 3 (Stat3) may functionally shift non-CSCs towards CSCs. However, whether Stat3 drives the formation of CSCs during the development of resistance in TNBC, and whether a Stat3 inhibitor reverses CSC-mediated Dox resistance, remains to be elucidated. In the present study, human MDA-MB-468 and murine 4T1 mammary carcinoma cell lines with the typical characteristics of TNBCs, were compared with estrogen receptor-positive MCF-7 cells as a model system. The MTT assay was used to detect cytotoxicity of Dox. In addition, the expression levels of CSC-specific markers and transcriptional factors were measured by western blotting, immunofluorescence staining and flow cytometry. The mammosphere formation assay was used to detect stem cell activity. Under long-term continuous treatment with Dox at a low concentration, TNBC cultures not only exhibited a drug-resistant phenotype, but also showed CSC properties. These Dox-resistant TNBC cells showed activation of Stat3 and high expression levels of pluripotency transcription factors Octamer-binding transcription factor-4 (Oct-4) and c-Myc, which was different from the high expression of superoxide dismutase 2 (Sox2) in Dox-resistant MCF-7 cells. WP1066 inhibited the phosphorylation of Stat3, and decreased the expression of Oct-4 and c-Myc, leading to a reduction in the CD44-positive cell population, and restoring the sensitivity of the cells to Dox. Taken together, a novel signal circuit of Stat3/Oct-4/c-Myc was identified for regulating stemness-mediated Dox resistance in TNBC. The Stat3 inhibitor WP1066 was able to overcome the resistance to Dox through decreasing the enrichment of CSCs, highlighting the therapeutic potential of WP1066 as a novel sensitizer of Dox-resistant TNBC.

Introduction

Triple-negative breast cancer (TNBC), which is negative for estrogen receptor (ER), progesterone receptor and human epidermal growth factor 2, is an aggressive subtype of breast cancer that has significant morbidity and mortality rates (1-5). Doxorubicin (Dox), a topoisomerase II inhibitor, is usually applied for the treatment of TNBC (1,2,6,7). However, the effectiveness is restricted by the development of resistance (7). Drug resistance was originally attributed to drug efflux mediated by adenosine triphosphate-binding cassette transporter (8,9). Increasing evidence has indicated the existence of cancer stem cells (CSCs) in breast cancer, and these CSCs are involved in the process of cancer initiation, metastasis, recurrence and therapeutic resistance (3,6,10-22). CSCs have cancer-initiating capability and metastatic potential (4,23-27). CD44 is a cell surface receptor for the extracellular matrix protein hyaluronan. CSCs expressing CD44 are considered to be responsible for adhesion, invasion, metastasis and the acquisition of resistance to apoptosis (6,12,25,28-33). ATP-binding cassette subfamily G member 2 (ABCG2) is overexpressed in several types of tumor and tumor cell lines (5,10,17,34-38). ABCG2 is not only associated with MDR, but may also function as a breast cancer stem cell marker (20,34,35,37,38). CD133 was initially described as a specific marker of human hematopoietic stem cells (13,14,17,34,39,40), CD133 has been used as the primary marker of putative CSCs and has been reported to be the most reproducible marker of breast CSCs (BCSCs) (14,24,27,39,41). Therefore, cells expressing high levels of cell surface markers, including CD44, ABCG2 and CD133, have stem-like activities (10,17,24). As the majority of information on CSCs in breast cancer cells has

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come from studies on the MCF-7/ADR cell line, originated from ER-positive MCF-7 cells (2,14,16), whether resistance of TNBCs to Dox is associated with the development of CSCs and, if so, the molecular characteristic phenotype of TNBC-derived CSCs remain to be elucidated.

The development and enrichment of CSCs may rely on the orchestration of multiple critical transcription factors. Pluripotent transcription factors contribute to the reprogramming of somatic cells into an embryonic stem cell (ESC)-like state (3,18). CSCs express members of the ESC core pluripotency complex, including octamer-binding transcription factor-4 (Oct4), superoxide dismutase 2 (Sod2) and Nanog, which are positively correlated with the development of CSCs (3,18,24,42-44). Furthermore, the Myc oncogene also forms a complex on its own (Myc-complex) to mediate CSCs (45-47). In addition to the overexpression of the above specific pluripotency transcriptional factors, signal transducer and activator of transcription 3 (Stat3) is important for mammosphere formation and the CSC process in breast cancer (3,15,46,48,49). The activation of Stat3 shifts non-CSC dynamics towards CSCs and regulates its downstream genes (3,18,50), in addition to other signaling molecules, which are important in tumorigenesis. It has been reported that the inhibition of Stat3 increased Dox sensitivity in the MDA-MB-231 human breast cancer cell line. Therefore, it was hypothesized that Stat3 inhibition may decrease the enrichment of CSCs during the induction of Dox resistance in TNBC cells.

In the present study, human MDA-MB-468 and murine 4T1 mammary carcinoma cell lines with the typical characteristics of TNBC were used as a model system to investigate the effects of the Stat3 signaling pathway and its inhibitor, WP1066, on the enrichment of CSCs and development of Dox resistance in breast cancer cells. TNBC cultures under prolonged selection with Dox at a low concentration exhibited a drug-resistant phenotype and had CSC properties with high expression levels of CD44, CD133 and ABCG2; this was regulated by Stat3-driven stemness through upregulating the expression of Oct-4 and c-Myc. This indicated a Stat3/Oct-4/c-Myc signaling circuit for the modulation of CSC-mediated Dox resistance in TNBC. The data provided direct evidence that WP1066 inhibited the Stat3-mediated enrichment of CSCs, in turn overcoming the resistance to Dox.

Materials and methods

Reagents. Dox (Solarbio, Beijing, China) was dissolved in sterile water in a 10 mM stock solution. WP1066 (Selleckchem, Shanghai, China) and 3-(4,5-dimethyl-2-thiazolyl)-2,5-di-phenyl-2-H-tetrazolium bromide (MTT; Solarbio) were lysed in DMSO (Beijing Solarbio Science & Technology Co., Ltd.). All cultures were maintained at 37°C in a humidified 5% CO₂ incubator. The MCF-7/Epr human epirubicin-resistant breast cancer cell line was obtained from Shanghai Bogoo Biotechnology (Shanghai, China), which was originally selected with 1,000 ng/ml epirubicin.

Mammosphere formation assay. Tumorsphere culture was performed in low attachment dishes (Corning Incorporated, Corning, NY, USA). The cells were seeded at a density of 8,000 cells per well into a 6-well plate supplemented with 1X B27 and 2X N-2 (both from Gibco; Thermo Fisher Scientific, Inc., Waltham, MA, USA), 20 ng/ml IGF, 20 ng/ml EGF, 10 ng/ml FGF-basic (all from PeproTech, Inc., Rocky Hill, NJ, USA) and 5 µg/ml heparin. The cells were maintained at 37°C in a humidified incubator containing 5% CO₂. Following 7 days of culture, the wells were examined under an inverted microscope at x400 magnification.

Immunofluorescent staining. The Dox-resistant tumor cells and parental tumor cells were harvested by cytopsin, respectively, and fixed in 4% paraformaldehyde for 10 min. The slides were washed three times in 1X PBS and, using goat sera (Wuhan Boster Biological Technology, Ltd., Wuhan, China), the cells were blocked for 1 h. The cells were incubated with CD133 (1:500 dilution, cat. no. ab130244; Abcam, Cambridge, MA, USA) primary antibodies at 4°C. Following a wash in PBS, the cells were incubated with immunoglobulin G/tetramethylrhodamine goat anti-rabbit antibody (1:200 dilution, cat. no. ZF0316; OriGene Technologies, Inc., Beijing, China) for 1 h at 4°C. The cells were then washed with PBS, stained with DAPI for 10 min, washed, and covered with mounting medium. Immunofluorescent staining was observed and images were captured.

Western blot analysis. Total protein was extracted from the Dox-resistant TNBC cells and parental cells, respectively. Briefly, the cells were lysed with RIPA lysis buffer combined with PMSF (both from Beijing Solarbio Science & Technology Co., Ltd.) and then centrifuged at 12,000 x g for 15 min at 4°C. The supernatants were collected for western blot analysis and protein concentration was determined using the BCA method (CW/Bio, Beijing, China). Total protein (25 µg) was separated by 10% sodium dodecyl sulfate-polyacrylamide gel electrophoresis and transferred onto a polyvinylidene fluoride membrane (EMD Millipore, Billerica, MA, USA). The membranes were blocked with 5% non-fat dry milk in TBST for 1 h and then blotted overnight at 4°C with the following primary antibodies: Mouse monoclonal β-actin (1:3,500 dilution, cat. no. 66009-1-lg; PeproTech, Inc.); mouse monoclonal Stat3 (1:1,000 dilution, cat. no. 9139s); mouse monoclonal phosphorylated (p)-Stat3 (1:1,000 dilution, cat. no. 4113s) (both from Cell Signaling Technology, Inc., Danvers, MA, USA); rabbit monoclonal Oct-4 (1:1,000 dilution, cat. no. ab200834); rabbit monoclonal c-Myc (1:5,000 dilution, cat. no. ab32072); rabbit monoclonal Sox2 (1:1,000 dilution, cat. no. ab92494); mouse monoclonal ABCG2 (1:1,000 dilution, cat. no. ab130244) (all from Abcam) and rabbit polyclonal CD133 (1:1,000 dilution, cat. no. 18470-1-AP; PeproTech, Inc.)
antibodies. The membranes were then blotted for 1 h at room temperature with anti-mouse (1:5,000 dilution, cat. no. A0216) and anti-rabbit (1:3,500 dilution, cat. no. A0208) secondary antibodies (both from Beyotime Institute of Biotechnology, Shanghai, China). The signal was detected using enhanced chemiluminescence (Immobilon Western Chemiluminescent Horseradish Peroxidase Substrate, cat. no. WBKLS0100; EMD Millipore) and recorded on X-ray film.

Flow cytometry. The dissociated MDA-MB-468-Dox cells, 4T1-Dox cells and their parental cells were counted and transferred into a tube, respectively. The cells were washed and resuspended in PBS at a concentration of 10^6/ml. The cell suspensions were incubated with PE-rat anti-mouse CD44 antibody, which also cross-reacts with human cells (1:100 dilution, cat. no. 553134; BD Biosciences, Franklin Lakes, NJ, USA) for 30 min at room temperature. Following washing with PBS, the samples were analyzed using a FACS Calibur. Isotype-matched PE-rat IgG2b (1:100 dilution, cat. no. 553989; BD Biosciences) served as negative control.

**MTT assay.** The MDA-MB-468-Dox cells, 4T1-Dox cells, MCF-7-Dox cells and their parental cells were seeded into 96-well plates at a density of 1x10^3 cells/well in RPMI-1640 medium supplemented with 10% FBS. The cells were treated with Dox at various concentrations (0, 0.05, 0.1, 0.5, 1, 5 and 10 µM) for 48 h at 37°C. MTT solution (5 mg/ml) was added to the plates (20 µl per well). The plates were incubated at 37°C for an additional 4 h. The supernatant was discarded, and 150 µl DMSO was added to each well. The plates were thoroughly mixed and read at 570 nm on a microplate reader.

**Statistical analysis.** Each experiment was performed at least three times. Data are presented as the mean ± standard deviation. A one-way analysis of variance was performed for comparisons between two subgroups, and the Fisher’s least significant difference post hoc test was used for determination of the significance among the means of multiple groups using SPSS version 17.0 software (SPSS, Inc., Chicago, IL, USA). P<0.05 was considered to indicate a statistically significant difference.

**Results**

**Establishment of Dox-resistant TNBC cells by long-term exposure to Dox at a low concentration.** Dox has been documented to induce the resistance of ER-positive MCF-7 breast cancer cells by exposure to a high dose of Dox. To establish Dox-resistant TNBC cell lines, human MDA-MB-468 and murine 4T1 cells were treated with a long-term low dose of Dox. Based on the dose-dependent curve of Dox, the concentration of 0.05 µM Dox was used to stimulate MDA-MB-468 cells and 0.1 µM of Dox was used to stimulate the 4T1 cells. Following continuous treatment with Dox for 4 weeks, morphological differences were observed between the parental cells and the Dox-resistant TNBC cells (Fig. 1A and B).
Dox-treated cells were termed MDA-MB-468-Dox cells and 4T1-Dox cells, and were banked for further investigations.

The cytotoxicity of Dox in the Dox-resistant cell lines and their parental cells was examined using an MTT assay. The dose-dependent effects of Dox in the above two pairs of cell lines are shown in Fig. 1C and D. Compared with the parental cancer cells, the cytotoxicity of Dox towards the Dox-resistant TNBC cells was significantly decreased. The EC_{50} of Dox in the parental MDA-MB-468 cells was 0.47 µM, whereas the EC_{50} of the MDA-MB-468-Dox was increased to 0.92 µM (P<0.05; Fig. 1C). Similarly, the EC_{50} of the 4T1-Dox cells was significantly increased from 0.98 to 5.37 µM (P<0.05; Fig. 1D). These data suggested that the sensitivity of TNBC cells against Dox was significantly decreased by long-term exposure with Dox at a low concentration.

To compare Dox resistance in TNBC cells to ER-positive cells, a human epirubicin-resistant breast cancer cell line MCF-7/Epr was induced into a Dox-resistant cell line, termed MCF-7-Dox, by prolonged exposure to 0.9 µM Dox for 4 weeks (Fig. 1E). The MTT assay demonstrated that the EC_{50} of the parental MDA-MB-468 cells was 0.47 µM, whereas the EC_{50} of the MDA-MB-468-Dox cells was increased to 0.92 µM (P<0.05; Fig. 1C). Similarly, the EC_{50} of the 4T1-Dox cells was significantly increased from 0.98 to 5.37 µM (P<0.05; Fig. 1D). These data indicated that the sensitivity of TNBC cells against Dox was significantly decreased by long-term exposure with Dox at a low concentration.

Expression levels of Oct-4 and c-Myc are increased in Dox-resistant cells. The high expression levels of pluripotent transcription factors, including Oct-4, Sox2 and c-Myc, has been reported in CSCs, which may promote stem cell self-renewal and differentiation (3,18,24,26). To clarify the roles of pluripotent transcription factors in regulating the enrichment of CSCs induced by Dox, the present study examined the expression of these genes in BCSCs induced by Dox. As shown in Fig. 3A, compared with their parental cancer cells, the expression levels of Oct-4 and c-Myc were increased in the Dox-resistant TNBC cells, whereas the expression of Sox2 was not altered significantly. However, the expression of Sox2 in MCF-7-Dox cells was increased. These results indicated that the upregulated expression of Sox2 may be a major factor in the CSC activities of MCF-7-Dox cells. Notably, Dox induced the expression of Oct-4 and c-Myc, but did not affect the expression of Sox2 to promote the CSC enrichment in the TNBC cells.

Activation of Stat3 is involved in CSC enrichment induced by Dox. In addition to specific pluripotency transcriptional factors, Stat3 is already known to be a key mediator for mammosphere formation and the CSC process in breast cancer (3,15,18,49,50). To investigate whether Stat3 is involved in CSC enrichment induced by Dox, the present study examined the expression and activation of Stat3. As shown in Fig. 3B, the phosphorylation of Stat3 was elevated in Dox-resistant breast cancer cells, whereas there was no significant change in the total expression of Stat3. This finding suggested that the activation of Stat3 was also involved in the regulation of CSC formation induced by Dox in breast cancer cells.

Inactivation of Stat3 by WP1066 reverses Dox resistance of TNBC cells by downregulating the expression of c-Myc and Oct-4 and reducing CSC enrichment. From the above data, it was observed that the pluripotency transcriptional factors Oct-4 and c-Myc, and the activation of Stat3 were involved in the Dox-induced CSC enrichment in TNBC. The CSC transcriptional factor c-Myc is a recognized transcriptional target of Stat3 in embryonic stem cells and hematopoietic stem cells (18,47). Oct-4 also acts as a downstream factor of Stat3 during mammosphere culture (3). Therefore, the Stat3 inhibitor WP1066 (51-56) was used to examine whether the inactivation of Stat3 affects the interaction between Stat3 and Oct-4/c-Myc in the process of Dox-induced CSC enrichment. Treatment with WP1066 (1.25 µM) not only decreased the phosphorylation of Stat3 (Fig. 4A), but also markedly downregulated the expression of Oct-4 and c-Myc (Fig. 4A). These data suggested that continuous Dox stimulation promoted the enrichment of CSCs through the activation of Stat3, which increased the expression of pluripotency transcriptional factors Oct-4 and c-Myc.

An MTT assay was performed to evaluate whether WP1066 restored the sensitivity of MDA-MB-468-Dox and 4T1-Dox cells to Dox. Following incubation with WP1066 (1.25 µM), the EC_{50} of MDA-MB-468-Dox cells was decreased from 0.32 to 0.14 µM (P<0.05; Fig. 4B). Similarly, the EC_{50} of 4T1-Dox cells was decreased from 0.98 to 0.47 µM (P<0.05; Fig. 4B). These data suggested that WP1066 reversed Dox resistance in TNBC cells.

The present study also performed mammosphere formation assays to evaluate the sphere-forming ability of Dox-resistant 4T1 cells. As shown in Fig. 2F, the volume of the sphere formed by Dox-resistant 4T1 cells was larger than that of the parental 4T1 cells. The numbers of mammospheres of the Dox-resistant 4T1 cells were also higher compared with those of the control 4T1 cells. These data obtained from the expression of surface molecular markers and mammosphere formation capacity indicated that continuous stimulation with Dox at a low dose induced the enrichment of CSCs in TNBC cells.
of 4T1-Dox was decreased from 5.37 to 2.24 µM (P<0.05; Fig. 4C). There was no significant difference in EC50 values between the WP1066 treatment group and parental MDA-MB-468 or 4T1 cells (P>0.05). These data suggested that WP1066 treatment sensitized the MDA-MB-468-Dox and 4T1-Dox cells to Dox, as demonstrated by EC50 values in WP1066-treated cells, which were similar or close to those in parental cells. In addition, following treatment with WP1066 at the concentration of 1.25 µM, the maximal response to Dox in the 4T1-Dox cells was lower than that in the parental 4T1 cells, indicating that this concentration of WP1066 did not completely restore the sensitivity of 4T1-Dox cells, which may be improved by increasing the dose and incubation time of WP1066.

The present study also examined whether WP1066 affects the expression of CSC markers on Dox-resistant cells. As shown in Fig. 5A, the increased expression of CD133 and ABCG2 on the MDA-MB-468-Dox cells was reduced by WP1066 treatment (Fig. 5A). The effect of WP1066 treatment on the CD44-positive cell population was also analyzed by flow cytometry. WP1066 treatment reduced the CD44-positive cell population in the MDA-MB-468-Dox cells (Fig. 5B) and 4T1-Dox cells (Fig. 5C). Analysis of the peak channel of the flow cytometry assay showed that, under treatment with WP1066, the mean peak value of 4T1-Dox was shifted left from 731 to 533. In addition, the mean peak value of MDA-MB-468-Dox was shifted left from 896 to 771. These results indicated that the inhibition of Stat3 downregulated the Dox-induced enrichment of the CSC population of TNBC cells, and led to reversion of TNBC cell resistance to Dox.
Discussion

Dox has a broad antitumor spectrum and is widely used in the treatment of various types of cancer, including breast cancer (1,6,7,54,55). Although Dox is effective in the majority of cases of primary breast cancer and almost half of the metastases at the early period of chemotherapy (5,54,56), the repeated application of Dox may lead to the development of chemotherapeutic resistance (5,7,9). ER-positive MCF-7/ADR cells selected by high dose of Dox have been found to overexpress ABCB1 protein and are widely used for investigations of the multidrug-resistant phenotype (2,5,35). In the present study, it was found that, following long-term continuous exposure to Dox at a low concentration, human and mouse TNBC cell lines exhibited a drug-resistant phenotype.

The enrichment of the CSC population may cause drug resistance, leading to recurrence and metastasis of tumors with poor prognosis (4,10,11,12,15,21-23,25,28,36,39). In the present study, MDA-MB-468-Dox and 4T1-Dox cells were used to investigate whether Dox induced CSC-enrichment. Using a sphere-forming assay, it was found that the low concentration of Dox (0.1 µM) stimulation resulted in increased mammosphere formation capacity in 4T1-Dox cells. In addition, increased expression levels of CSC biomarkers, including CD44, ABCG2 and CD133, were observed in the MDA-MB-468-Dox and 4T1-Dox cells, compared with those in their control parental cells. These data confirmed that exposure of TNBC cells to low dose of Dox resulted in the enrichment of BCSCs. These findings were consistent with previous observations on lung cancer cells (7). The CSCs may evade or efficiently repair DNA damage from radiation and chemotherapy, and these stem cell properties are likely to be responsible for the difficulties existing in the treatment of breast cancer (8,16,31,32). Therefore, the enrichment of CSCs may contribute to the development of TNBC resistance to Dox.

Dox-induced chemoresistance in breast cancer cells may occur through the activation of Stat3 (49,54). Gariboldi et al observed that the inactivation of Stat3 by AG490 increased the sensitivity of MDA-MB-231 cells to Dox treatment (49). However, how the activated Stat3 regulates the resistance of breast cancer cells to Dox remains to be elucidated. Pluripotent transcription factors, including Oct-4, c-Myc and Sox2, are important in the induction of CSCs through promoting CSC self-renewal (4,18,24,26). The cooperation between the CSC transcriptional factors c-Myc, Oct-4, and the activation of Stat3 has been considered as a mechanism in sustaining and
Figure 4. Inactivation of Stat3 by WP1066 restores the sensitivity of MDA-MB-468-Dox and 4T1-Dox cells to Dox. (A) Western blot analysis of Stat3, p-Stat3, c-Myc and Oct-4 in MDA-MB-468-Dox and 4T1-Dox cells treated with or without WP1066. (B) Effect of WP1066 on the sensitivity of MDA-MB-468-Dox to Dox was examined by MTT assay. (C) Effect of WP1066 on the sensitivity of 4T1-Dox to Dox was examined by MTT assay. The results are representative of three independent experiments. Dox, doxorubicin; MTT, 3-(4,5-dimethyl-2-thiazolyl)-2,5-diphenyl-2-H-tetrazolium bromide; ctrl, control.

Figure 5. Inactivation of Stat3 by WP1066 reduces cancer stem cells enriched by Dox. (A) Changes in the expression of CD133 and ABCG2 in MDA-MB-468-Dox cells with the use of WP1066 were examined by western blot analysis. (B) Changes in the expression of CD44 with the use of WP1066 in MDA-MB-468-Dox cells were detected by flow cytometry. (C) Changes in the expression of CD44 with the use of WP1066 in 4T1-Dox cells was detected by flow cytometry. The results are representative of three independent experiments. Dox, doxorubicin; ABCG2, ATP-binding cassette subfamily G member 2; ctrl, control.
amplifying pluripotent stem cells (44,57-59), however, the functional interaction has proven obscure. In the present study, it was found that the pluripotent transcription factors Oct-4 and c-Myc were expressed at high levels, and the phosphorylation of Stat3 was elevated in the Dox-resistant TNBC cells. Based on the above data and previous studies, a hypothetic model was suggested for the different pathways involved in the regulation of Dox-induced CSC enrichment during the development of Dox resistance between TNBCs and ER-positive breast cancer cells (Fig. 6). The activation of Stat3 promotes the expression of Oct-4 and c-Myc, and further induces the enrichment of CSCs in TNBC cells. However, in the case of ER-positive MCF-7 cells, a high dose of Dox induces the resistance to Dox through the formation of CSCs mediated by activation of the Stat3/Sox2 pathway, which differs from the regulatory pattern of Stat3 during the development of resistance of TNBCs to Dox.

The Stat3 inhibitor WP1066 has been shown to be a potent antitumor agent in the treatment of malignant glioma and acute myelogenous leukemia cells (53,54). However, whether WP1066 reverses the resistance to Dox in breast cancer cells remains to be fully elucidated. In the present study, it was shown that inhibiting the activation of Stat3 with WP1066 markedly decreased the expression of pluripotent transcription factors Oct-4 and c-Myc. The data from the flow cytometry assay (Fig. 4B) and peak channel analysis further demonstrated that the inhibition of Stat3 affected the enrichment of CSC populations in Dox-resistant TNBC cells. Under treatment with WP1066, the proportion of CD44-positive cells was markedly reduced in 4T1-Dox and MDA-MB-468-Dox cells. These data revealed that the inhibition of Stat3 led to a decrease in CSC populations and restored the sensitivity of TNBC cells to Dox. As CSCs exhibit increased drug resistance, tumorigenesis and metastasis, compared with non-CSC cancer cells, it may be worthwhile to investigate the possible differential effects of the inhibition of Stat3 on sorted CSC and non-CSC population in future investigations.

In conclusion, the data obtained in the present study provided direct evidence that a low concentration of Dox induced the enrichment of BCSCs in TNBC cells. The activated Stat3-driven stemness mediated the resistance of TNBC cells to Dox. As CSCs exhibit increased drug resistance, tumorigenesis and metastasis, compared with non-CSC cancer cells, it may be worthwhile to investigate the possible differential effects of the inhibition of Stat3 on sorted CSC and non-CSC population in future investigations.

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Availability of data and materials

The datasets used and/or analyzed during the current study are available from the corresponding author on reasonable request.

Authors' contributions

CNC and LHS contributed to study design, experimental investigation, data analysis and manuscript drafting; XJW, SXW and XQW participated in stem cell culture and characterization; SRL, YFW and CNC conducted flow cytometry assays and data analysis; ZL and CNC performed experiments analyzing the phosphorylation and expression of transcription factors; LHS, LHW and YD contributed to the conception, analysis and interpretation of data; ZL, LHW and YD wrote, reviewed and edited the manuscript. All authors have read and approved the final manuscript.

Ethics approval and consent to participate

Not applicable.

Consent for publication

Not applicable.

Competing interests

The authors declare that they have no competing interests.

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