Role of Bcl-2 on drug resistance in breast cancer polyploidy-induced spindle poisons

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Abstract. Spindle poisons are chemotherapeutic drugs used in the treatment of malignant tumors; however, numerous patients develop resistance following chemotherapy. The present study aimed to induce polyploidy in breast cancer cells using the spindle poison nocodazole to investigate the mechanism of polyploid-induced tumor resistance. It was revealed that the spindle poison nocodazole induced apoptosis in HCC1806 cells but also induced polyploidy in MDA-MB-231 cells. The drug sensitivities of the polyploid MDA-MB-231 cells to paclitaxel, docetaxel, epirubicin, 5-fluorouracil and oxaliplatin were lower than those of the original tumor cells; however, the polyploid MDA-MB-231 cells were more sensitive to etoposide than the original tumor cells. The expression of F-box and WD repeat domain containing 7 (FBW7) was decreased, while the expression of MCL1 apoptosis regulator BCL2 family member (MCL-1) and Bcl-2 was increased, and caspase -3/9 and Bax were not expressed in MDA-MB-231 cells. The resistance to docetaxel and etoposide was reversed, but the sensitivity of paclitaxel was not changed following Bcl-2 silencing. The formation of polyploidy in tumors may be one of the molecular mechanisms underlying tumor resistance to spindle poisons. Expression of the Bcl-2 family members, for example FBW7 and MCL-1, plays a key role in apoptosis and the cell escape process that forms polyploid cells. However, Bcl-2 silencing has different reversal effects on different anti-tumor drugs, which requires further investigation.

Introduction

Tumor chemotherapy is an important part of the comprehensive treatment for solid tumors (1). Paclitaxel, vincristine and other spindle poisons are widely used as first-line chemotherapy for various types of malignant tumor, such as breast cancer, cervical cancer and melanoma (2,3). One of the important methods in treating cancer is inducing tumor cell death. In the clinical setting, it was observed that certain types of tumor are sensitive to this type of drug, and in such cases, the tumor tissue disappears following several cycles of chemotherapy (4,5). However, more commonly, the tumor tissue is sensitive to the drug at the beginning of chemotherapy, but drug resistance eventually occurs, or the tumor tissue has no obvious response to this class of drugs (6). The latter two phenomena are the root cause of uncontrollable tumors and recurrence. Previous studies demonstrated that spindle poisons play a role in the induction of apoptosis in some cells (2,7). However, certain cells do not undergo apoptosis when exposed to such drugs, and instead form polyploids. Polyploid tumors are extremely insensitive to common radiotherapy and chemotherapy options, and patients who have tumors with polyploid subclones have a worse prognosis (8,9). Therefore, it was speculated that the formation of polyploids may be associated with tumor resistance.

For a number of years, research has focused on elucidating the molecular mechanism underlying tumor resistance, and numerous theories and potential mechanisms have been proposed; however, a resolution to this clinical issue has not yet been identified (10,11). With continuous scientific development, it has become apparent that the causes of tumor resistance are multi-faceted. In addition to the current multi-drug resistance genes that are known, it was speculated that the formation of polyploids is likely to lead to resistance to spindle poisons and is an important cause of tumor drug resistance after use (10-12); however, a mechanism for this possible drug resistance has not yet been proposed. The phenomenon of tumor-forming polyploidy induced by spindle poison was used
as a basis for the design of the present study. MDA-MB-231 cells were treated with the spindle poison nocodazole in order to establish a stable polyploid cell model. Subsequently, the sensitivity of chemotherapeutic drugs on polyploid and original tumors was investigated with the aim of elucidating the association between polyploid formation and tumor resistance, and to attempt to determine the molecular mechanism underlying polyploidy occurrence through in-depth analyses. A potential intervention point/method to address such resistance was identified, and a new strategy was proposed in the present study to prevent the occurrence of tumor resistance by inhibiting the formation of polyploid subclones.

Materials and methods

Cell lines and cell culture. For routine culturing, human breast cancer HCC1806 and MDA-MB-231 cells were obtained from China Infrastructure of Cell Line Resources, Institute of Basic Medical Sciences, Chinese Academy of Medical Sciences, and MDA-MB-231 polyploid cells (T-MDA-MB-231) were established in the laboratory. These cells were cultured in RPMI-1640 medium (Corning Inc.) containing 10% newborn calf serum (Corning, Inc.) at 37°C and 5% CO₂. Cells in the logarithmic growth phase were used in the subsequent experiments.

Nocodazole drug induction assay. HCC1806 and MDA-MB-231 cells (1x10⁶ cells/ml) were inoculated into 10-cm culture dishes for 24 h and then the antibiotic-free RPMI 1640 medium was replaced fresh antibiotic-free RPMI-1640 medium with 10% fetal bovine serum (Corning, Inc.) containing nocodazole (Sigma-Aldrich; Merk KGaA). The base volume was 10 ml, and the final concentration of nocodazole was 100 µg/l; the cells were treated for 0-72 h at 37°C, and 3 duplicate wells were examined for each group. The morphological changes in the HCC1806 and MDA-MB-231 cells prior to and following nocodazole treatment were observed under a light microscope at x200 magnification and images were captured at randomly selected fields.

Cell cycle analysis. MDA-MB-231 and HCC1806 cells were treated with 100 µg/l nocodazole for 0, 6, 12, 24, 48 or 72 h, and the cells were trypsinized by 0.25% trypsin, collected, fixed in 70% cold ethanol and incubated overnight at 4°C. After centrifuging for 5 min at 300 x g and at room temperature, the ethanol was discarded, and the pellets were washed with PBS and incubated with 100 ng/l RNAse for 37 min at 37°C. A total of 500 µl propidium iodide (PI; 2 µg/ml) stain solution was added, the cells were incubated at room temperature for 30 min in the dark prior to being counted (1x10⁶ cells/ml). The cell cycle (G₀/S or G₂/M) was detected via flow cytometry (FACSCanto II; BD Biosciences), and the percentages of diploid, tetraploid and octoploid cells in the total cell population were determined by the FACSDiva software (version 8.0.1; BD Biosciences).

Flow cytometry sorting of polyps. According to the results of the cell cycle assay, 100 µg/l nocodazole-treated MDA-MB231 cells were harvested at the point of maximal polyploid formation (72 h), and the cells were resuspended in RPMI 1640 medium at 4°C at a concentration of 1x10⁶ cells/ml. Subsequently, Hoechst 33342 (5 µg/ml) was added, and the cells were incubated at 37°C for 90 min. After centrifuging at 300 x g for 10 min at 4°C, the supernatant was discarded, and the cells were resuspended in pre-chilled Hanks' Balanced Salt Solution (Beyotime Institute of Biotechnology). A total of 500 µl PI (2 µg/ml) was then added to the solution prior to sieving using a 200-mesh nylon cell sieve. Flow cytometry (FACSCanto II) was used to obtain drug-treated polyploid cells (T-MDA-MB-231) using the FACSDiva software (version 8.0.1). After routine stable growth, polyploid cells in the logarithmic growth phase were subjected to secondary sorting using the same method. Subcloning of polyploid cells was performed by limiting dilution.

Apoptosis experiment. The present study aimed to compare the sensitivity of polyploid cells and their diploid parental cells to commonly used chemotherapeutic drugs. Therefore, representative drugs from different chemotherapeutic drug classes, paclitaxel and docetaxel (Jiangsu Hengrui Medicine Co., Ltd.), were used for the apoptosis experiments. MDA-MB-231 and T-MDA-MB-231 cells in the logarithmic growth phase (5x10⁶ cells/ml) were plated into 6-well cell culture plates and cultured for 24 h at 37°C and 5% CO₂ using the 30% inhibitory concentration (IC₃₀) and the 60% inhibitory concentration (IC₆₀) of 0.66 and 2.50 µM for paclitaxel and 3.48 and 12.10 µM for docetaxel, respectively, to treat the MDA-MB-231 and T-MDA-MB-231 cells. After incubating for 72 h at 37°C, the cells were centrifuged at 300 x g for 4 min at 4°C, and 5 µl Annexin V and 10 µl PI were added, followed by incubation for 15 min at room temperature in the dark. Apoptosis was subsequently detected by flow cytometry (FACSCanto II; excitation wavelength, 488 nm) with the FACSDiva software (version 8.0.1).

MTT assay. T-MDA-MB-231 cells were seeded at 5x10⁵ cells/well in 96-well plates using 2 and 4 µM of the multi-kinase inhibitor sorafenib (Sigma-Aldrich; Merck KGaA) alone or in combination with 100 µg/l nocodazole or 100 nmol/l paclitaxel for 48 and 72 h, with 5 replicate wells per group; 4 µl of a 5 g/l MTT solution were added for 4 h. At termination, the medium was removed from each well, and 200 µl of dimethyl sulfoxide was added per well, followed by shaking until the crystals dissolved. The same method was applied to compare differences in the cell cytotoxicity of MDA-MB-231 and T-MDA-MB-231 cells treated with spindle poisons, paclitaxel (0, 0.39, 0.78, 1.56, 3.125, 6.25, 12.5, 25 and 50 µM), docetaxel (0, 0.39, 0.78, 1.56, 3.125, 6.25, 12.5, 25 and 50 µM), epirubicin (0, 0.39, 0.78, 1.56, 3.125, 6.25, 12.5, 25 and 50 µM), 5-fluorouracil (FU) (0, 0.78, 1.56, 3.125, 6.25, 12.5, 25, 50 and 100 µM), oxaliplatin (0, 0.78, 1.56, 3.125, 6.25, 12.5, 25, 50, and 100 µM) or etoposide (0, 0.0195, 0.039, 0.078, 0.156, 0.3125, 0.625, 1.25 and 2.50 µM). Epirubicin was purchased from Pfizer Inc. 5-FU, oxaliplatin or etoposide were purchased from Jiangsu Hengrui Medicine Co., Ltd. Differences in the cytotoxicity of paclitaxel, docetaxel, epirubicin, 5-FU, oxaliplatin or etoposide on T-MDA-MB-231 cells and MDA-MB-231 cells were also examined by MTT assay. The absorbance value at 490 nm of each well was measured using a microplate reader with the tecan i-control 2.0 software, and the cytotoxicity rate was calculated as 1-ΔΔA⁶₀⁰ drug group/ΔΔA⁶₀⁰ control group x100%.
Western blotting. MDA-MB-231 and T-MDA-MB-231 cells (3x10⁵) were treated with 100 µg/l nocodazole for 48 h and then digested and centrifuged at 300 x g for 5 min at 4°C and collected. Subsequently, the cells were lyed using radioimmunoprecipitation assay lysis buffer (Beyotime Institute of Biotechnology), and the cell lysates (total protein) were collected by centrifugation at 16,000 x g for 10 min at 4°C. Protein concentration was determined using a BCA kit (Beyotime Institute of Biotechnology). For each sample, 100 µg of total cell protein per lane was separated via SDS-PAGE (12% gel) and then transferred to a polyvinylidene fluoride membrane. Following blocking with 5% skimmed milk at 4°C for 1 h, the membrane was incubated with primary monoclonal antibodies (all purchased from Abcam) against FBW7 (1:1,500; monoclonal antibodies (all purchased from Abcam) against β-actin (1:500; cat. no. ab179467) overnight at 4°C. The membrane was washed three times to remove the primary antibody, and then incubated with goat anti-rabbit-horse-radish peroxidase-labeled secondary antibody (1:1,000; cat. no. ab6721; Abcam) for 1 h at room temperature. After washing three times, Immobilon Western chemiluminescent horseradish peroxidase substrate (EMD Millipore) was used to develop the membranes. The experiment was performed three times and visualized by Image J (National Institutes of Health; version 1.8.0).

Reverse transcription-quantitative PCR (RT-qPCR). MDA-MB-231 and T-MDA-MB-231 cells were digested, centrifuged at 300 x g for 5 min at 4°C and collected. Total RNA was extracted using TRZol® (Invitrogen; Thermo Fisher Scientific, Inc.), and cDNA was obtained via the BeyoRT™ RT kit (Beyotime Institute of Biotechnology), according to the manufacturer's protocol. qPCR was subsequently performed using BeyoFast™ SYBR Green qPCR Mix kit (Beyotime Institute of Biotechnology) according to the manufacturer's protocol. The following primer pairs were used: Bcl-2 forward, 5'-ATG TGT GTG GAG AGC GTC AA-3' (140 bp) and reverse, 5'-GCC GTA GTG GAG AGC GTC AA-3' (140 bp); and GAPDH forward, 5'-GAAGATGGTAAGGTCGGAGTC-3' (225 bp) and reverse, 5'-GGCCGTA CAGTTCCACAAAGG-3' (140 bp); and GAPDH forward, 5'-GAAGATGGTAAGGTCGGAGTC-3' (225 bp) and reverse, 5'-GGCCGTA CAGTTCCACAAAGG-3' (140 bp); and a final extension was performed at 72°C for 10 min. GAPDH was used as an internal reference to calculate the relative expression of the Bcl-2 gene via the 2−ΔΔCq method (13).

RNA interference. T-MDA-MB-231 cells were transfected using Lipofectamine™ 2000 (Invitrogen; Thermo Fisher Scientific, Inc.) according to the manufacturer's protocol, and MDA-MB231 cells were used as controls. Logarithmic growth phase T-MDA-MB231 cells at 1.5x10⁵ cells per well were added to 6-well plates and incubated for 24 h, after which Lipofectamine™ 2000 diluted in Opti-MEMI (Corning; Thermo Fisher Scientific, Inc.) was added at 50 µl/well and mixed at room temperature for 5 min. After 5 min, 100 µl of liposomes (Invitrogen; Thermo Fisher Scientific Inc.) and Bcl-2 small interfering (si) RNA (forward, 5'-UGUAGGAUGA CUGAUAUCCUGA-3'; reverse, 5'-UCAGGUAUCAGUCA UCCACAGG-3') or control siRNA (AATTCTCGAACGTG TCACGTCTCGTCTC) were purchased from Ambion (Thermo Fisher Scientific, Inc.) and incubated at room temperature for 20 min. The liposome-siRNA complexes were added to the cells and then incubated at 37°C for 6 h, at which time the media was replaced with fresh RPMI-1640 medium with 10% fetal bovine serum to continue the culture. The mRNA expression of Bcl-2 in T-MDA-MB-231 cells was detected by RT-qPCR as aforementioned, 72 h following siRNA transfection.

Statistical analysis. The statistical analyses were performed using SPSS software (version 17.0; SPSS, Inc.). Data are expressed as the mean ± standard deviation from at least three separate experiments. Experimental results were assessed using one-way analysis of variance and SNK-q post hoc tests. P<0.05 was considered to indicate a statistically significant difference. GraphPad Prism software (version 5.0; GraphPad Software, Inc.) was used to calculate the inhibitory concentrations.

Results

Nocodazole induces apoptosis in HCC1806 cells and polyploidy in MDA-MB-231 cells. A final nocodazole concentration of 100 ng/ml was used in the present study as it has been demonstrated that 100 ng/ml could induce polyploid (14,15) After nocodazole was applied to MDA-MB-231 and HCC1806 cells for 24 h, the cells began to appear rounded, and after 48 h, the cells were all rounded. However, with prolonged exposure for 72 h to nocodazole, the HCC1806 cells exhibited typical apoptotic morphological changes; the cell volume gradually decreased, the cells became deformed and floated, their nuclei shrank, and apoptotic bodies formed. Fragmentation was gradually increased, and after 72 h, apoptotic bodies were widespread. However, the MDA-MB-231 cells exhibited increased cell volume over time with drug exposure, and typical polyploid morphological changes were apparent, such as gradually increasing nuclei. After 72 h, the polyploid appearance was prevalent (Fig. 1A).

In order to further verify the responses of MDA-MB-231 and HCC1806 breast cancer cells to nocodazole in the present study, samples of the two cell lines were analyzed by flow cytometry in the present study. After HCC1806 cells were treated with nocodazole for 6 h, the percentage of G2/M phase cells increased, and the proportion of sub-G0 phase cells gradually increased. For MDA-MB-231 cells treated with nocodazole for 6, 12 and 24 h, the percentages of cells in the sub-G1 and G2/M phases were the same as those of HCC1806 cells, but the percentage of cells in the G1/S/G2/M phase increased after 36 h, and body (tetraploid and octoploid) cell formation increased significantly (Fig. 1B). These experiments indicated that nocodazole can induce apoptosis in HCC1806 cells but instead induces polyploidy in MDA-MB-231 cells.

The western blot analysis revealed that, compared with the control group, HCC1806 cells expressed caspase-9 and caspase-3 after 24 h of incubation, indicating that nocodazole induced apoptosis in HCC1806 cells. The
expression of caspase-9, caspase-3 and Bax was not detected in MDA-MB-231 cells. However, the Bax protein was continuously expressed in HCC1806 cells. In MDA-MB-231 cells, the expression of Bcl-2 increased significantly until 36 h, and its expression levels were significantly different. Bcl-2 was also expressed in HCC1806 cells, and its expression increased after 36 h (Fig. 2). These experiments indicated that an increasing Bcl-2 in HCC1806 cells promoted apoptosis following treatment with nocodazole, while the high expression of Bcl-2 in MDA-MB-231 cells inhibited apoptosis and formed polyploid cells. High expression levels of Bcl-2 inhibit nocodazole-induced apoptosis in breast cancer cells and may be one of the primary molecular mechanisms underlying polyploid formation.

Effect of polyploid MDA-MB-231 on chemosensitivity. The appearance and polyploid certification of MDA-MB-231 parent cells and T-MDA-MB-231 polyploid cells is presented in Fig. 3A and B, viewed under a light microscope at x200 magnification and flow cytometry. Meanwhile, there was a slower rate of cytotoxicity in T-MDA-MB-231 cells, demonstrated by the MTT assay (Fig. 3C). The MTT assay results also revealed that T-MDA-MB-231 cells were less sensitive to paclitaxel, docetaxel, vincristine, oxaliplatin, 5-FU and epirubicin than MDA-MB-231 cells (all P<0.05). However, T-MDA-MB-231 polyploid cells were more sensitive to etoposide (Fig. 3D). The flow cytometry results revealed that following treatment with paclitaxel and docetaxel at IC_{30}, IC_{50} and IC_{70} for 72 h, the survival rates of the polyploid T-MDA-MB-231 cells were higher than those of the
parental MDA-MB-231 cells. The proportion of cells in early apoptosis was also lower in polyploid cells (Fig. 3E), indicating that polyploid tumors are more resistant to these anticancer drugs. 

Role of the Bcl-2 protein family in polyploidy-induced tumor resistance. The RT-qPCR results revealed that the relative expression of Bcl-2 mRNA in T-MDA-MB-231 polyploid cells was significantly higher than that of the MDA-MB-231 parental cells. After silencing the expression of Bcl-2 mRNA in T-MDA-MB-231 cells using siRNA technology, the relative expression decreased significantly (P<0.05; Fig. 4A) but was not significantly different to the expression levels
Figure 4. Role of the Bcl-2 protein family in polyploidy-induced tumor resistance in MDA-MB-231. (A) The relative mRNA expression of Bcl-2 of T-MDA-MB-231 cells, MDA-MB-231 parental cells and T-MDA-MB-231 cells following Bcl-2 silencing. (B) The half maximal inhibitory concentration of docetaxel, paclitaxel and etoposide for T-MDA-MB-231 cells with or without Bcl-2 silencing. (C) The protein expression of FBW7 and MCL-1 in MDA-MB-231 cells at different time points. (D) The protein expression of MCL-1 protein in MDA-MB-231 cells following treatment with nocodazole and sorafenib. β-actin was used as an internal control. 1, control; 2, 2 µM sorafenib + 100 µg/l nocodazole; 3, 100 µg/l nocodazole; 4, 2 µM sorafenib. (E) The protein expression of MCL-1 protein in MDA-MB-231 cells following treatment with paclitaxel and sorafenib. β-actin was used as an internal control. 1, control; 2, 2 µM sorafenib + 100 nM paclitaxel; 3, 100 nM paclitaxel; 4, 2 µM sorafenib. (F and G) The effect of ploidy on treatment with nocodazole, sorafenib, paclitaxel and sorafenib in the MDA-MB-231 cells. (H) The effect of sorafenib on the cytotoxicity of MDA-MB-231 at 48 and 72 h. *P<0.05 vs. 100 µg/l Nocodazole group; ΔP<0.05 vs. 100 µg/l Nocodazole + 2 µM Sorafenib group; **P<0.05 vs. 100 nM Paclitaxel group; #P<0.05 vs. 100 nM Paclitaxel + 2 µM Sorafenib group.
of the MDA-MB-231 parental cells. Following the silencing of Bcl-2 expression, mRNA was downregulated, the IC_{50} of docetaxel for T-MDA-MB-231 polyploid cells was significantly decreased, and the IC_{50} of etoposide was significantly increased; however, Bcl-2 silencing had no significant effect on the paclitaxel IC_{50} (Fig. 4B). These results indicated that the higher expression of Bcl-2 in polyploidy cells plays a key role in the process by which cells escape apoptosis and further form polyploid cells, but has different effects on anti-tumor drugs with different mechanisms.

MCL-1 is an important anti-apoptotic protein in the Bcl-2 family, and it may play an important role in polyploidy resistance (16). With the prolonged nocodazole treatment time, expression of the FBW7 protein was decreased in MDA-MB-231 cells, while expression of MCL-1 was increased (Fig. 4C). The present study observed that after 48 h of treatment, the expression of MCL-1 was decreased and demonstrated by comparing the nocodazole+sorafenib group with the nocodazole group, and the paclitaxel+sorafenib group with the paclitaxel group in MDA-MB-231. At 48 h, with sorafenib, expression of the MCL-1 protein was decreased, the number of diploid cells was increased, and the number of polyploid cells was decreased in MDA-MB-231 cells (Fig. 4D and E). There was an increase in diploid and decrease in polyploid cells in the nocodazole+sorafenib and sorafenib groups compared with the nocodazole group. There was a similar effect of ploidy in the nocodazole+sorafenib, paclitaxel+sorafenib and sorafenib groups, which polyploid was downregulated after sorafenib treating cells for 48 and 72 h (Fig. 4F and G), indicating that the combination of sorafenib and single-purpose spindle poisons can increase drug sensitivity and be used to reverse polyploidy resistance, the occurrence of which may be associated with the regulation of MCL-1 protein expression and the cell cycle. The present study tested different concentrations of nocodazole and sorafenib and it was concluded that 100 μg/l nocodazole+4 μM sorafenib may be the best option by increasing cytotoxicity (Fig. 4H).

Discussion

At present, paclitaxel, vincristine and other spindle poisons are the first-line chemotherapy options for the majority of solid tumors, such as breast and ovarian cancer (17,18). These drugs affect the structure and function of spindles during the process of mitosis in tumor cells, destroying the kinetochore and affecting the adhesion of the tube and the integrity of the mitotic apparatus to prevent normal splitting of the chromosome centromere, which is originally placed in the center of the equatorial plate (19,20). Consequently, the chromosome cohort cannot be normally separated from the two poles of the spindle, and the spindle assembly checkpoint is activated. Under normal conditions, if the cell damage cannot be repaired, then the cells are unable to undergo mitosis, thereby initiating the apoptotic pathway and inducing apoptosis to achieve the ultimate goal of tumor cell death (21-24). However, certain cells ignore the spindle structure that is formed during mitosis, and in the case of functional disruption, the nucleus does not divide prior to entering the next cell cycle, forming polyploid tumor cells, which enhances resistance to chemotherapeutic drugs (25-27).

Nocodazole is responsible for inducing cell cycle arrest in the G2/M phase (28-30). Nocodazole interferes with the cell microtubules to form a spindle, which blocks cell division (7). The results of the present study demonstrated that nocodazole had a significant effect on the cell cycle, and G2/M phase arrest occurred following induction in MDA-MB-231 cells; however, the cells did not completely exit mitosis or activate apoptosis and instead entered the next cell cycle. The human breast cancer MDA-MB-231 cells were polyploidized, with typical polyploid morphological changes such as cell volume increase and nuclear enlargement observed. At present, the occurrence of drug resistance following the polyploidization of tumor cells is controversial. Castedo et al (31) reported that polyploid tumors exhibit significant resistance to cisplatin and camptothecin. Havas et al (32) demonstrated that cells were more sensitive to chemotherapeutic drugs following polyploidization. The present study revealed that polyploid tumor cells induced by spindle poisons were less sensitive to paclitaxel, docetaxel, vincristine, oxaliplatin, 5-FU and epirubicin than the original tumor cells. In addition, the results indicated that induced polyploid cells are relatively resistant to the majority of commonly used chemotherapeutic drugs and relatively sensitive to the topoisomerase II inhibitor etoposide, which not only validates the genetic instability of polyploid cells but also suggests that induced polyploid cells are likely to be a key problem in drug resistance during tumor treatment. It was hypothesized that there may be an especial mechanism underlying topoisomerase II inhibitors, for example, etoposide for polyploid breast cancer, which is different with other drugs such as paclitaxel and docetaxel. Therefore, the present study further investigated the specific molecular mechanism underlying polyploid cell resistance.

Nocodazole did not produce polyploidy in HCC1806 cells. MDA-MB-231 and HCC1806 cells have distinct p53 mutations that occur in response to spindle poisons, and both have complete spindle assembly checkpoints (33). MDA-MB-231 cells are characterized by polyploid formation, and HCC1806 cells are characterized by apoptosis, meaning that their differential responses to nocodazole may be associated with inhibition of the apoptotic pathway. In order to further investigate whether the apoptotic pathway is involved in the formation of polyploid tumor cells induced by nocodazole, expression of the apoptotic pathway proteins Bcl-2 and Bax in nocodazole-induced polyploid tumors cells was investigated using two strains treated with the drug. Human breast cancer cells with different responses were studied: MDA-MB-231 cells were characterized by the formation of polyploid cells, and HCC1806 cells were characterized by apoptosis. Whether key proteins of the apoptosis pathway, Bcl-2 and Bax, were involved in the mechanism of polyploid tumor formation was preliminarily investigated.

The flow cytometry analysis revealed that the percentage of HCC1806 cells in the G2/M phase increased following nocodazole treatment for 6 h, and the subdiploid peak (sub-G0 phase), also called the ‘apoptosis peak’, was increased at 36 and 48 h compared with 0 h. Changes in the percentages of MDA-MB-231 cells in the G2/M phase and sub-G0 phase following nocodazole treatment for 6, 12 and 24 h were the same as those of the HCC1806 cells. However, as the cells were incubated with the drug for a prolonged period of time, the two human breast cancer cells exhibited different outcomes:
mRNA expression was downregulated in MDA-MB-231 polyploid cells. The IC50 of docetaxel for T-MDA-MB-231 polyploid cells was significantly decreased, suggesting they may affect the formation of polyploidy in MDA-MB-231 cells. FBW7 degrades the downstream oncogenic protein MCL-1 by recognizing phosphorylation groups, promoting apoptosis and exerting anticancer effects. Therefore, in the present study, both mdm2 and p53 were detected in MDA-MB-231 parental cells and T-MDA-MB-231 polyploid cells. The western blot analysis revealed that the overexpression of mdm2 with no expression of p53 in MDA-MB-231 cells, which leads to polyploid cell tumor formation. Furthermore, mdm2 expression and tumor resistance are multifactorial, with numerous important molecular mechanisms. The present study investigated the molecular mechanisms underlying polyploidy and the association between polyploidy and tumor resistance in an attempt to ascertain potential interventional methods, which may lead to an improved understanding of polyploid tumor cell formation.

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Availability of data and materials
The datasets used and/or analyzed during the present study are available from the corresponding author on reasonable request.

Authors' contributions
BY and YZ contributed to the study conception and design. YZ wrote and revised the manuscript. JH, QZ and YW were
Ethics approval and consent to participate

Not applicable.

Patient consent for publication

Not applicable.

Competing interests

The authors declare that they have no competing interests.

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