Lapatinib sensitizes quiescent MDA-MB-231 breast cancer cells to doxorubicin by inhibiting the expression of multidrug resistance-associated protein-1

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Abstract. The quiescent state plays an important role in tumor recurrence because it protects cancer cells from chemotherapy. Previously, we optimized tumorsphere cultures for in vitro screening methods for targeting quiescent cell population since the majority of cells in tumorspheres are quiescent. In this study, we analyzed efficacies of current chemotherapeutics in tumorsphere assays to seek better strategies for eradicating quiescent cell population. Tumorspheres generated from MDA-MB-231 cells exhibited accumulations of cells in the G0/G1 phase as compared with cells in monolayer culture, suggesting that sphere formation contributes to an increase of quiescent cells. As a result of a decreased doxorubicin uptake, MDA-MB-231 tumorspheres exhibited chemoresistance to both doxorubicin and paclitaxel. Since we found that the enhanced EGFR signaling is characteristics of MDA-MB-231 tumorspheres, the combination effects of chemotherapy with lapatinib, a dual ErbB1/ErbB2 inhibitor, were accessed in tumorsphere assays. Western blot analysis revealed that lapatinib inhibited the phosphorylation of EGFR, AKT and p38 in doxorubicin-treated tumorspheres. The inhibition of EGFR signaling by the treatment with lapatinib suppressed the expression of multidrug resistance-associated protein-1 (MRP-1), leading to increased cytotoxicity of doxorubicin to tumorspheres. Furthermore, blockade of the PI3K/AKT and p38 MAPK signaling pathways resulted in a remarkable decrease in the expression of MRP-1 in doxorubicin-treated tumorspheres. These results demonstrate that lapatinib sensitizes quiescent MDA-MB-231 breast cancer cells to doxorubicin by inhibiting doxorubicin-induced MRP-1 expression via PI3K/AKT and p38 MAPK signaling pathways. Thus, this study suggests that treatment with lapatinib in combination with anti-mitotic drugs maybe a useful approach to improve clinical response by eradicating the quiescent cancer cell population.

Introduction

Chemoresistance is a major cause of cancer treatment failure. The mechanisms involved in the development of chemoresistance are complex and not fully understood. Since conventional chemotherapies target proliferating cells and require active cycling for induction of apoptosis, it has been proposed that cells in the quiescent state within tumors are associated with resistance and cell survival in chemotherapies (1). Moreover, the quiescent nature of cells is known to be characteristic of cancer stem cells, which have the ability to self-renew and differentiate to produce heterogeneous tumor cell lineages (1-3). Increasing evidence indicates that cancer stem cells are responsible for treatment failure and tumor recurrence because their quiescent nature is likely to contribute to the survival in response to chemotherapy (4). Thus, selectively targeting quiescent cell population including cancer stem cells offers possible way forward to overcome chemoresistance and improve the clinical outcomes of cancer patients.

The sphere culture has been proposed to propagate cells with stem cell properties and has been widely adopted to study stem cell biology (5-8). It is a relatively easy, rapid and non-animal-depending model to assess stem cell activity, but the application of sphere-forming assays for high-throughput screening is limited due to formation of variable sizes of spheres and lack of a simple and well-established analytical...
tool. Moreover, cell aggregation in spheres can cause misinterpretation. Previously, we showed that sphere cultures exhibit higher proportions of quiescent cells and we optimized tumorosphere cultures for the in vitro screening of chemotherapeutics against the quiescent cell population (9).

In this study, we utilized tumorsphere cultures to identify better ways of eradicating quiescent tumor cell population in MDA-MB-231 human breast cancer cells. MDA-MB-231 cells are representative of triple-negative breast tumors, which are characterized by the absence of estrogen receptor (ER), progesterone receptor (PR) and human epidermal growth factor receptor 2 (HER2) (10). This triple negative tumor subtype is mainly correlated to poor outcomes, showing the worst overall and disease-free survival rates due to a lack of effective targeted therapies (11,12). Thus, cytotoxic chemotherapeutics such as doxorubicin/paclitaxel remain the mainstay of treatment for triple-negative breast cancer, but resistance is common and can develop rapidly (13). To seek better strategies to overcome chemoresistance, we analyzed the efficacies of chemotherapeutics, doxorubicin and paclitaxel, on MDA-MB-231 tumorspheres. Since we found that the enhanced epidermal growth factor receptor (EGFR) signaling pathway is characteristic of MDA-MB-231 tumorspheres, we assessed the combination effects of doxorubicin and lapatinib (a dual ErbB1/ErbB2 inhibitor) in tumorsphere assays.

**Materials and methods**

**Adherent cell culture.** The MDA-MB-231 human breast cancer cell line was purchased from the Korean Cell Line Bank (Seoul, Korea) and routinely maintained in DMEM (Welgene, Daegu, Korea) supplemented with 10% fetal bovine serum (Invitrogen, Carlsbad, CA, USA) and 1% antibiotic-antimycotic solution (Welgene).

**Suspension sphere cultures.** The protocol used for tumorsphere culture was as previously described (5,6,8,14). Briefly, cells were suspended in serum-free DMEM/F12 (Welgene) supplemented with 1:50 B27 (Gibco BRL, Grand Island, NY, USA), 10 µg/ml insulin (Welgene), 20 ng/ml recombinant human epidermal growth factor (EGF; R&D Systems, Minneapolis, MN, USA), 10 ng/ml recombinant human fibroblast growth factor (FGF; R&D Systems), and 1% antibiotic-antimycotic solution (Welgene) and cultured in non-adherent plates.

**Cell kinetic assay.** To examine cell proliferation rates, MDA-MB-231 cells were plated at different concentrations (3,000-20,000 cells/well) into 96-well plates under non-adherent (see above), or monolayer culture conditions. After 4 days, premixed cell proliferation reagent WST-8 (Dojindo Laboratories, Kumamoto, Japan) was added to each well and the absorbance of the water-soluble formazan produced by viable cells was measured at 450 nm according to the manufacturer’s instructions.

**Cytotoxicity assay.** To compare the chemosensitivity of cells in the TS and 2D culture systems, several chemotherapeutics including doxorubicin (Sigma, St. Louis, MO, USA), paclitaxel (Sigma), lapatinib (a dual ErbB1/ErbB2 inhibitor; LC Laboratories, Woburn, MA, USA), U0126 (a MEK inhibitor; LC Laboratories), or LY294002 (a PI3K/AKT inhibitor; LC Laboratories) were added into cells grown in either adherent or non-adherent 96-well plates and cell viabilities were measured 3 days later. To examine the combinational effects of doxorubicin and lapatinib, cells were treated with doxorubicin in a range of 0.2-1 µM in the presence of 5 µM lapatinib.

**Flow cytometry analysis: cell cycle analysis and doxorubicin uptake.** For cell cycle analysis, cells grown in 2D or TS culture for 4 days were trypsinized after washing with PBS, centrifuged at 1,000 rpm for 3 min and fixed in cold 70% ethanol. After centrifugation, the cells were washed with PBS containing 2% FBS and stained in the dark with 20 µg/ml propidium iodide (Sigma) and 200 µg/ml RNase A (Sigma) for 30 min at room temperature. The cells were analyzed by FACS Calibur II flow cytometry (Becton Dickinson Biosciences, San Jose, CA, USA). To measure intracellular doxorubicin accumulations in cells grown in 2D or tumorsphere cultures, cells were seeded into adherent or non-adherent 6-well plates for 3 days. Cells were treated with 0.5 µM doxorubicin for 30 min, trypsinized, and washed twice with PBS containing 2% FBS. At 24 h after resuspending in PBS containing 2% FBS, the cells were analyzed by FACSCalibur II flow cytometry (Becton Dickinson Biosciences).

**Western blotting.** Cells grown in 2D or TS conditions were lysed with RIPA buffer (50 mM NaCl, 1% Triton X-100, 1% sodium deoxycholate, 0.1% SDS, 50 mM Tris-HCl, pH 7.5 and 2 mM EDTA). Phosphatase and protease inhibitor cocktails were added, and then proteins were separated by SDS-PAGE and transferred to a nitrocellulose membrane. Membranes were incubated with primary antibodies against drug resistance-associated protein-1 (MRP-1) and GAPDH and washed extensively with PBST (phosphate buffered saline with 0.1% Tween-20) prior to incubation with horseradish peroxidase-conjugated secondary antibodies and detection with a chemiluminescence detection system (ECL; GE Healthcare). A chemiluminescence image was obtained using a ChemiDoc Imaging System (Bio-Rad Laboratories). The intensity of the bands was quantified using the ImageJ software (National Institutes of Health, Bethesda, MD, USA).

**RNA extraction, RT-PCR and quantitative real-time PCR.** Cells cultured under 2D and TS conditions were treated with 0.3 µM doxorubicin, 5 µM LY294002 or 10 µM SB203580 (a p38 inhibitor; LC Laboratories) for 3 days, and harvested for RNA isolation. Total RNA was extracted using the easy-Blue™ Total RNA Extraction kit (iNtRON Biotechnology Inc., Sungnam, Korea) and cDNA was synthesized with reverse transcriptase (TaKaRa, Shiga, Japan). RT-PCR for cyclin D1, MDR-1, and GAPDH were conducted as previously described (15). Densitometric analysis was performed using Scion Image software (Scion Co., Frederick, MD, USA). The utilized primer sequences for the RT-PCR reactions were as follows: CyclinD1 (forward) 5'-AGCTCTGTGCTGCGAAGTG ACAAAC-3' and Cyclin D1 (reverse) 5'-AGTGTTCAAT GAAATCGTGCGGGG-3'; MDR-1 (forward) 5'-GCC TGGCAGCTGGAAGACAAATACACAAATT-3' and MDR-1 (reverse) 5'-CAGACAGCACGCTGACAGTCACA TAAGACACAGACGACT-3'; GAPDH (forward) 5'-ATCCCATCAC CATCTTCAG-3' and GAPDH (reverse) 5'-TTCTAGACGC AGTGATTGTAG-3'. The real-time PCR reactions for multi-drug resistance-associated protein-1 (MRP-1) and GAPDH were performed using QuantiMix SYBR green kit (Philekorea, Daejeon, Korea) in Eco Real-time PCR (Illumina, San Diego, CA, USA). mRNA expression level of MRP-1 was calculated by the 2^(-ΔΔCt) method. The expression levels were normalized to GAPDH. The utilized primer sequences for the real-time PCR reactions were as follows: MRP-1 (forward) 5'-GGCAAGGTGTCCTCCCTCAAAAG-3' and MPR-1 (reverse) 5'-TCTCCTCAGGTTGATGCTTGC-3'; GAPDH (forward) 5'-ATCCCATCAC CATCTTCAG-3' and GAPDH (reverse) 5'-TTCTAGACGC AGTGATTGTAG-3'. Western blotting. Cells grown in 2D or TS conditions were lysed with RIPA buffer (50 mM NaCl, 1% Triton X-100, 1% sodium deoxycholate, 0.1% SDS, 50 mM Tris-HCl, pH 7.5 and 2 mM EDTA). Phosphatase and protease inhibitor cocktails were added, and then proteins were separated by SDS-PAGE and transferred to a nitrocellulose membrane. Membranes were incubated with primary antibodies against drug resistance-associated protein-1 (MRP-1) and GAPDH and washed extensively with PBST (phosphate buffered saline with 0.1% Tween-20) prior to incubation with horseradish peroxidase-conjugated secondary antibodies and detection with a chemiluminescence detection system (ECL; GE Healthcare). A chemiluminescence image was obtained using a ChemiDoc Imaging System (Bio-Rad Laboratories). The intensity of the bands was quantified using the ImageJ software (National Institutes of Health, Bethesda, MD, USA).
(GenDepot, Barker, TX, USA) were added immediately before use. Lysates were cleared of debris at 13,000 rpm for 10 min, and protein concentrations were determined using bicinchoninic acid reagent (Sigma). Equal amounts of protein were separated by SDS-PAGE and transferred to polyvinylidene fluoride (PVDF) membranes, which were then blocked with 5% non-fat skim milk in 1X TBS-0.1% Tween-20 (TTBS) for 2 h and incubated with a primary antibody (EGFR, p-EGFR, AKT, p-AKT, ERK 1/2, p-ERK 1/2, p38, p-p38 or GAPDH; Cell Signaling, Beverly, MA, USA) overnight. hRP-conjugated secondary antibody (Santa Cruz Biotechnology, Santa Cruz, CA, USA) diluted 1:5,000 was incubated with blots for 1 h at room temperature. Blots were developed using Luminescent Image Analyzer LAS-4000 (Fujifilm, Tokyo, Japan).

Statistical analysis. Statistical significance was determined using the Student’s t-test. All experiments were conducted in triplicates, and results are presented as mean ± SD. P-values of <0.05 were considered significant.

Results

Quiescence in tumorspheres generated from MDA-MB-231 cells. First, we cultured MDA-MB-231 breast cancer cells in non-adherent culture condition for 4 days to test their ability to form tumorspheres. As previously reported (14), tumorspheres generated from MDA-MB-231 cells exhibited much looser structures (Fig. 1A) than those derived from MCF-7 cells (data not shown). To evaluate the cell growth rates of tumorspheres and monolayer cultures, cells were plated at different concentrations (3,000-20,000 cells/well) into 96-well plates, in either non-adherent plates, or regular tissue culture plates. After 4 days of 2D or TS culture, cell viabilities were assayed by measuring WST-8 absorbance. The overall WST-8 readings of suspension cultures were at least three times less than those of 2D cultures, seeded with the same cell numbers (Fig. 1B). The cell cycle analysis revealed that this slow cell growth rate in tumorspheres correlated with the accumulation of cells at the G0/G1 phase, showing that 79.81±9.4% cell population in tumorspheres was in the G0/G1 phase, whereas 68.76±5.5% of the 2D cultured cell population was in the G0/G1 phase (Fig. 1C). Moreover, the mRNA expression of cyclin D1 in cells grown as 2D and TS.

Chemoresistance of tumorspheres to doxorubicin and paclitaxel. To investigate whether culturing cells as spheres affects chemo-sensitivity, MDA-MB-231 culture under TS or 2D conditions were exposed to different concentrations of doxorubicin (0.2-1 µM) or paclitaxel (5-60 nM) for 3 days and then assessed for cell viability. It was found that tumorspheres were resistant to both doxorubicin and paclitaxel (Fig. 2A and B). In fact, the IC50 of doxorubicin for tumorspheres was at least three fold higher than its IC50 for 2D cultured cells (Fig. 2A). Similarly, tumorspheres exhibited significant resistance to paclitaxel with an IC50 value ten times that of 2D cultured cells (Fig. 2B).

We further analyzed intracellular doxorubicin accumulation using flow cytometry (16). Cells grown from either monolayer or tumorsphere cultures were treated with 0.5 µM doxorubicin for 30 min and then doxorubicin fluorescence was analyzed by flow cytometry. The treatment of 2D cultured cells with doxorubicin for 30 min caused a right-shift of fluorescence intensity of doxorubicin as compared with untreated MDA-MB-231 cells, confirming the intracellular accumulation of doxorubicin within cells. On the other hand, the fluorescence intensity of tumorspheres was lower than that of monolayer cultured cells (Fig. 2C), implying that the decreased accumulation of intracellular doxorubicin may contribute to chemoresistance of tumorspheres.

It has been well established that the intracellular accumulation of doxorubicin is associated with the expression of the ABC family of drug transporters (17), and therefore, we examined the mRNAs expression levels of MDR-1 and MRP-1. It was observed...
that the expression of MDR-1 was similarly induced by doxorubicin in TS and 2D cultured cells (Fig. 2D). However, mRNA expression of MRP-1 was induced in tumorspheres by doxorubicin but not in monolayer cultures (Fig. 2E), suggesting that MRP-1, and not MDR-1, was involved in the chemoresistance demonstrated by tumorspheres.

**Enhanced EGFR signaling in tumorspheres.** To elucidate the molecular mechanisms responsible for the chemoresistance of tumorspheres, we first analyzed which signaling pathways are upregulated in the tumorspheres. Western blot analysis showed that the EGFR signaling pathway was more activated in TS than in 2D cultured cells (Fig. 3A). Furthermore, the phosphorylation of EGFR in tumorspheres was associated with concomitant increases in the phosphorylations of ERK1/2 and AKT.

We next explored the role of the EGFR signaling pathway in the formation of tumorspheres. Cells were treated with lapatinib, U0126, or LY294002 to block the EGFR, MAPK, and PI3K/AKT signaling pathways, respectively, and then cell viabilities were assessed. Of note, responses to lapatinib were similar for TS and 2D cultured cells (Fig. 3B), whereas TS cultured cells were less sensitive to U0126 and to LY29400 (Fig. 3C and D). These results suggested that the EGFR signaling pathway plays an important role in mediating the survival of cells in the quiescent state rather than pathways downstream of EGFR, such as, the MAPK and PI3K/AKT pathways.

**Lapatinib sensitized tumorspheres to doxorubicin by inhibiting the expression of MRP-1.** We next examined whether the blockade of EGFR signaling by lapatinib enhanced cytotoxic effect of doxorubicin on quiescent MDA-MB-231 cells. Cells were treated with different concentrations of doxorubicin in the presence of 5 µM lapatinib for 3 days and the cell viability was measured. Noteworthy, the chemosensitivity of tumorspheres to doxorubicin was more enhanced by lapatinib than that of 2D cultured cells (Fig. 4A and B). As shown in Fig. 4A, treatment with lapatinib mildly increased cytotoxicity of doxorubicin in the monolayer culture but the chemosensitivity of tumorspheres was dramatically increased in the presence of lapatinib (Fig. 4B). In fact, the 60% cell viability after treatment with 1 µM of doxorubicin was significantly decreased to <5% when TS cultured cells were treated with 1 µM doxorubicin in the presence of 5 µM lapatinib (Fig. 4B).

Since we found that increased MRP-1 expression was responsible for the chemoresistance of tumorspheres (Fig. 2E), we tested whether treatment with lapatinib affected the expression of MRP-1 in doxorubicin-treated tumorspheres. As shown in Fig. 4C, doxorubicin-induced MRP-1 expression was significantly suppressed in the presence of lapatinib, suggesting that lapatinib sensitizes tumorspheres to doxorubicin by inhibiting the expression of MRP-1.

**Lapatinib inhibits doxorubicin-induced MRP-1 expression by inhibiting PI3K/AKT and p38 MAPK signaling pathways.** To obtain more insight into the mechanism underlying the inhibitory effects of lapatinib against doxorubicin-induced MRP-1 expression in tumorspheres, we first analyzed the effects of lapatinib on EGFR and its downstream signaling pathways. Treatment with lapatinib was found to inhibit the phosphorylation of EGFR and that of AKT and p38 in doxorubicin-treated
Figure 3. Enhanced epidermal growth factor receptor (EGFR) signaling in tumorspheres. (A) Whole cell lysates of tumorsphere (TS) or monolayer (2D) cultured cells were analyzed by western blotting for EGFR, AKT and ERK 1/2. Chemotherapeutic responses of 2D and TS cultured cells to (B) lapatinib (an inhibitor of ErbB1 and ErbB2), (C) LY294002 (a PI3K/AKT inhibitor), or (D) U0126 (a MEK1/2 inhibitor). After treatment with each inhibitor for 3 days, cell viability was assessed using WST-8 reagent. Results represent three independent experiments performed in triplicate. Error bars indicate the SD of means.

Figure 4. Lapatinib sensitizes tumorspheres to doxorubicin (Dox) by inhibiting the expression of MRP-1. Synergistic cytotoxic effects of Dox and lapatinib (Lap) on (A) monolayer (2D) cultured cells and on (B) tumorsphere (TS) cultured cells. MDA-MB-231 cells were treated with various concentrations of Dox and 5 µM lapatinib (Dox+Lap) for 3 days and viabilities were assessed using WST-8 reagent. (C) Inhibitory effect of lapatinib on Dox-induced MRP-1 expression in tumorspheres. Tumorspheres were treated with 0.3 µM Dox and/or 5 µM Lap for 3 days and the mRNA expression of MRP-1 was analyzed by quantitative real-time PCR. Results represent three independent experiments performed in triplicate. Error bars represent SD of means. *P<0.05, **P<0.01.

Figure 5. Lapatinib inhibits MRP-1 expression via PI3K/AKT and p38 MAPK pathways in tumorspheres. (A) Cell lysates from tumorspheres (TS) treated with 0.3 µM doxorubicin (Dox) and/or 5 µM lapatinib for 3 days were analyzed by western blotting for EGFR, AKT, and p38. Downregulation of Dox-induced MRP-1 expression by (B) LY294002 and by (C) SB203580 (a p38 inhibitor) in tumorspheres. TS were treated with 0.3 µM Dox and/or 5 µM LY294002 or 10 µM SB203580 for 3 days and the mRNA expression of MRP-1 was analyzed by quantitative real-time PCR. Results represent three independent experiments performed in triplicate. Error bars on the graph indicate the SD of means. *P<0.05, **P<0.01, ***P<0.001.
ongoing arguments about the enrichment of cancer stem cells such as cell density and culture duration (14,18,19). Despite as a suitable in vitro cancer stem cell enrichment and disagreed on considering it that the formation of tumorspheres does not always predict in stem cell biology (5,6,8). However, several studies reported adapted to detect and propagate human breast cancer stem cells breast cancer cells. Tumorsphere culture has been widely tion of quiescent cell population in MDA-MB-231 human strategies to overcome chemoresistance based on the eradica -tion of quiescent cell population in MDA-MB-231 human breast cancer cells. Tumorsphere culture has been widely adapted to detect and propagate human breast cancer stem cells in stem cell biology (5,6,8). However, several studies reported that the formation of tumorspheres does not always predict cancer stem cell enrichment and disagreed on considering it as a suitable in vitro culturing method for cancer stem cells because the formation of tumorspheres is influenced by factors such as cell density and culture duration (14,18,19). Despite ongoing arguments about the enrichment of cancer stem cells in tumorspheres, the generation of tumorspheres confers interesting and unique features such as quiescence (20,21). Previously, we showed that most cells in tumorsphere cultures are quiescent, whereas cells in monolayer culture have a high mitotic index (9). Similar to this study, the sphere-forming population of hepatoma cells contained a higher proportion of cells in the G0/G1 phase than the same cells cultured as monolayers (22). Since the quiescence is one of the traits in understanding the contribution of cancer stem cells to chemoresistance, we previously optimized tumorsphere cultures for in vitro screening methods for evaluating chemotherapeutics against quiescent cell population (9).

In the present study, tumorspheres generated from MDA-MB-231 cells exhibited chemoresistance to both doxorubicin and paclitaxel. However, we found that the epidermal growth factor receptor (EGFR) signaling pathway was more activated in TS than in 2D cultured cells and this enhanced activation of EGFR signaling in tumorspheres mediated survival of cells in quiescent state. EGFR is a member of the ErbB family of receptors and its activation by specific ligand binding triggers several signal transduction cascades, principally the PI3K/AKT and MAPK pathways, leading to cell proliferation, adhesion, and migration (23,24). In human tumors, EGFR and the other three members of the EGFR family, HER2, HER3, and HER4, are often overexpressed or dysregulated, which promotes tumor growth and/or progression (23,24). EGFR is frequently overexpressed in triple-negative breast cancer and is emerging as a therapeutic target (25). Although the use of single-agent of tyrosine kinase inhibitors targeting EGFR in triple-negative breast cancer patients have produced the disappointing results (26,27), several studies have reported that cytotoxic chemotherapy in combination with EGFR inhibition has shown promising results in treatment of breast cancer patients (28,29). Lapatinib is an orally active small molecule, which inhibits the tyrosine kinases of EGFR and HER2, and has been approved by the FDA in combination with other anticancer agents for the treatment of HER2-positive breast cancers (30). More recent studies reported that lapatinib enhanced the cytotoxic effect of chemotherapeutics including paclitaxel, vincristine, and topotecan by inhibiting the drug efflux function of ABC transporters, such as P-glycoprotein (P-gp), MRP's, or ABCG2 (BCRP) transporters (31-33).

ABC transporters have been linked to the development of resistance to anticancer drugs as they are involved in the ATP-dependent efflux of xenobiotics or chemotherapeutics from cells and tissues (34,35). Consistent with other studies, our results also show that blockade of the EGFR signaling pathway by lapatinib significantly increased the anticancer activities of doxorubicin on quiescent MDA-MB-231 cells by inhibiting the expression of MRP-1. However, the inhibition of MRP-1 expression may be insufficient to increase the anticancer activity of doxorubicin, because we observed that, unlike lapatinib, the blockade of the PI3K/AKT or p38 MAPK signaling pathways with LY294002 or SB203580, respectively, did not increase the cytotoxic effect of doxorubicin on tumorspheres although they were able to suppress the expression of MRP-1 in tumorspheres. These observations suggest that the synergistic effects of lapatinib and doxorubicin may arise from the inhibition of MRP-1 expression and the inhibition of EGFR-mediated survival signaling pathways.

In summary, we propose that although EGFR inhibition alone does not represent an effective therapeutic approach to triple-negative breast cancer, treatment with lapatinib in combination with cytotoxic chemotherapy may provide a useful approach to improve clinical responses by eradicating the quiescent cell population.

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