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Inhibiting autophagy does not decrease drug resistance in breast cancer stem-like cells under hypoxic conditions

Bui Dinh Khan^{1,2,*}, Tran Ngo The Nhan^{1,2}, Phuc Van Pham^{2,3}

ABSTRACT

Introduction: Breast cancer has been one of the most frequently diagnosed cancers in women over the past decade. The presence of a breast cancer stem cell (BCSC) population within breast tumors significantly contributes to malignancy and drug resistance. Under the hypoxic conditions of breast tumors, BCSCs exhibit increased resistance to chemotherapy, complicating treatment and reducing its efficacy. Consequently, targeting BCSCs is considered a crucial strategy for breast cancer treatment. We hypothesize that the chemotherapy resistance of BCSCs is dependent on the upregulation of autophagy triggered by hypoxia. This study aims to investigate the relationship between autophagy and drug resistance of BCSCs in hypoxic conditions. Methods: BCSCs were isolated from Vietnamese breast cancer cell line 1 (VNBRCA1) based on expression of CD44 and CD24. Cobalt (II) chloride (CoCl₂) was used to simulate a hypoxic environment in monolayer cell cultures. The hypoxic status of BCSCs was confirmed by the upregulation of the HIF-1 α gene. Autophagy activation was assessed through the expression of autophagy-related genes (LC3B, Beclin-1, AMPK, PI3K, AKT). Drug resistance was evaluated by measuring the expression of ABCC10 and assessing cell proliferation under cisplatin treatment. **Results**: The study revealed that CoCl₂ treatment led to high expression of HIF-1 α and significant upregulation of LC3B in BCSCs, together with enhanced expression of ABCC10. Consequently, BCSCs exhibited marked resistance to cisplatin in the hypoxic environment. However, inhibition of autophagy using the autophagy inhibitor 3-methyladenine (3-MA) did not reduce cisplatin resistance in hypoxic conditions. Conclusion: The findings indicate that in hypoxic conditions, BCSCs enhance both the autophagy process and drug resistance. However, the drug resistance of BCSCs in hypoxic conditions is not directly associated with the upregulation of autophagy. Therefore, targeting autophagy inhibition may not be an effective strategy to reduce drug resistance in hypoxia.

Key words: breast cancer stem-like cells, hypoxia, autophagy, drug resistance, cobalt chloride

INTRODUCTION

According to the 2022 GLOBOCAN statistics, breast cancer is the second most common cancer, making up 11.5% of the 19,976,499 new cases, and ranks fourth in cancer-related deaths, accounting for 6.8% of the 9,743,832 fatalities globally. Furthermore, this data indicates that in Viet Nam, breast cancer represented the highest proportion of new cancer diagnoses in 2022, with the majority of patients being women. Primary prevention measures like reducing excess body weight, lowering alcohol consumption, increasing physical activity, and promoting breastfeeding can help reduce breast cancer incidence. In 2021, WHO launched the Global Breast Cancer Initiative to reduce mortality by 2.5% annually, aiming to save 2.5 million lives in 20 years through health promotion, timely diagnosis, and comprehensive management¹.

Cancer stem cells (CSCs) play an important role in contributing to the heterogeneity of many kinds of

cancer². About 20 years ago, Al-Hajj et al. determined that a small breast cancer cell population strongly expressed CD44 and weakly expressed CD24, which was able to develop tumors in NOD/SCID mouse models³. Later, ALDH1 was also detected as a novel marker for breast cancer stem cell isolation, which is correlated with poor clinical outcomes in breast cancer patients⁴. Breast cancer stem cells are significant due to their role in metastasis and therapy resistance. Efforts are focused on characterizing them, understanding their pathways, and developing targeted therapies to eradicate them. New therapeutic approaches, including drug delivery, gene targeting, and immunotherapy, hold promise for effectively targeting breast cancer stem cells in cancer treatment⁵. Cancer stem cells are usually located at the core of solid tumors, distant from the blood vessels, creating a hypoxic microenvironment (hypoxia). In comparison, while the level of oxygen pressure in normal tissue reaches over 40 mmHg, concentration in cancer tumors is frequently maintained between 0 and

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20 mmHg. Transcription factors, which are responsible for adapting to hypoxic conditions, are known as hypoxia-induced factors (HIFs)⁶. We have reviewed the important effects of hypoxia as well as the roles of HIFs during tumor progression⁷. Particularly, *HIF-1* α and *HIF-2* α regulate cancer stem cells' adaptation to hypoxia and resistance to therapies⁸.

The term "autophagy" comes from a Greek word meaning "eating itself" and was first introduced in 1963 by a Belgian physiologist Christian de Duve. Mechanisms related to this biological process were studied and published by Yoshinori Ohsumi and his colleagues from the 1960s to the 1990s^{9,10}. This multi-step process includes autophagy induction, recognition, autophagosome formation, fusion with lysosome, and autophagolysosome destruction¹¹. Each step involves the regulation of many genes called autophagy-related genes (ATGs). At least 40 ATGs have been discovered and 20 of them play key roles in the autophagy of eukaryotic cells¹². In cancer cells, autophagy acts not only as a "friend" since it stimulates cancer cells to proliferate, but also as a "foe" due to its inhibitory impacts on cancer cell survival¹³. In the case of tumor suppression, autophagy protects cancer cells by preventing the accumulation of DNA-damaging reactive oxygen species (ROS) and by removing damaged proteins and organelles to maintain homeostasis^{14,15}. Conversely, autophagy activation also aids tumor cell survival by regulating various mediators^{16,17}. Overall, the process of autophagy in cancer cells is highly complex and not yet fully understood.

Hypoxia has been shown to induce autophagy in some types of cancer stem cells^{18,19}, which has led to studies on signaling pathways related to autophagy in hypoxic cancer stem cells. In this study, we use cobalt (II) chloride (CoCl₂) to mimic hypoxic conditions in breast cancer stem-like cell populations to investigate changes in the expression of genes related to autophagy induction and regulation. Additionally, we examine changes in drug resistance gene expression *ABCC10* and cisplatin resistance in breast cancer stem-like cells in a hypoxia-mimic model. Based on these results, we draw preliminary conclusions about the molecular relationship between autophagy and drug resistance in hypoxic breast cancer stem-like cells.

METHODS

Cell Culture

VNBRCA1 stands for the Vietnamese breast cancer cell line no. 1, which was isolated and characterized

from Vietnamese breast cancer patients and was described and used by our laboratory in previous studies²⁰. Cancer cells were cultured in DMEM (Lonza, Switzerland) supplemented with 10% FBS (Thermo Fisher Scientific, USA) and 1% antimycotic-antibiotic (HyClone, USA) for enrichment in a T-25 flask (SPL Life Science, Korea Republic). All flasks were incubated at 37°C and 5% CO₂, and the medium was freshly changed after 48 to 72 hours. After reaching 80-90% confluency, cells were harvested by Deattachment reagent (Regenmedlab, Viet Nam) to perform subculture.

Flow cytometry assay for CD44^{high}CD24^{low} population

Cells were harvested by Deattachment reagent as described above. Approximately 1×107 VNBRCA1 cells were stained with 10 µL of anti-CD44 (BD Biosciences, USA) and 10 µL of anti-CD24 (Sigma-Aldrich, USA) in phosphate-buffered saline (supplemented with 2% BSA) at 4°C in the dark for 20 minutes. Then, all cells were sorted based on fluorescentbound antigen signals using the BD FACSMelody Cell Sorter (BD Biosciences, USA). The cell population that is strongly positive for CD44 and negatively or dimly expressed CD24 is considered breast cancer stem cells (BCSCs). They are marked as BC-SCs hereafter in this article. BCSCs were cultured in M171 medium (Thermo Fisher Scientific, USA) supplemented with 1% MEGS (Thermo Fisher Scientific, USA) at 37° C and 5% CO₂.

Cobalt (II) Chloride Cytotoxicity Assay

Cytotoxicity was determined using the AlamarBlue assay²¹. BCSCs were seeded at a density of 3,000 cells per well in a 96-well plate (SPL Life Science, Korea Republic). Each well was filled with 100 μ L of M171. After 24 hours of incubation, the medium was withdrawn and replaced with new medium containing CoCl₂ at different concentrations. Medium without CoCl₂ was used as a negative control. The percentage of cell viability was calculated based on the following formula:

Rate of viable cells (%) = $\frac{OD \text{ value of treated group} - OD \text{ value of blank}}{OD \text{ value of untreated group} - OD \text{ value of blank}} x 100\%$

Cisplatin Cytotoxicity Assay

Breast cancer stem-like cells were seeded at a density of 3,000 cells per well in a 96-well plate. They were then pretreated with different concentrations of CoCl₂ for various periods of time. Subsequently, the M171 medium was withdrawn and replaced with new medium containing cisplatin (Sigma-Aldrich, USA) at concentrations of 50, 100, 150, 200, and 250 μ M, respectively. BCSCs were incubated at 37°C for the next 24 hours, and cell viability was assessed via the AlamarBlue assay as described above.

Gene Expression Analysis

RNA isolation was performed using the easy-BLUETM Total RNA Extraction Kit (iNtRON Biotechnology, South Korea). The expression levels of target genes were determined by real-time quantitative RT-PCR (reverse transcriptase polymerase chain reaction) with specific primers (listed in Table 1). All primers were purchased from PHUSA Biochem Co., Viet Nam. RNA samples were divided into qPCR tubes (Thermo Fisher Scientific, USA) with appropriate amounts and supplemented with the necessary components for a standard PCR reaction, following the instructions in the Luna Universal One-Step RT-qPCR Kit (New England Biolabs, USA). Thermal cycling was performed using the Replex system (Eppendorf, USA). Gene expression levels were semi-quantitated using the Livak formula, with β -actin used for normalization.

Statistical Analysis

All assays were performed in triplicate. Raw data were collected using Microsoft Excel 365 and analyzed with GraphPad Prism 9.0 software. All data are displayed as mean \pm standard error of the mean (SEM). Differences between treatments were examined using the one-way ANOVA method, with a p-value < 0.05 considered statistically significant.

RESULTS

Establishing model of hypoxic BCSCs using CoCl₂

After sorting, the BCSC population predominantly exhibits a $CD44^{high}CD24^{low}$ phenotype (over 96%) and shows an approximately 1.5-fold increase in ALDH1 gene expression compared to the pre-sorting population (**Figure 1A**). However, this increase in ALDH1 expression is not statistically significant (p-value = 0.20) (**Figure 1B**). This population was used for further analysis.

In the first assay to evaluate the cytotoxicity of CoCl₂, CoCl₂ displayed high toxicity in BCSCs when used in high concentrations. Indeed, when compared to cells not treated with CoCl₂, the survival rate of BC-SCs markedly decreases in a concentration- and timedependent manner when exposed to CoCl₂ concentrations ranging from 100 to 800 μ M for 24 to 72 Table 1: Sequences of primer pairs used for qRT-PCR

| • | | • |
|------------------|-------------------------|------------------------|
| Genes | Primer sequence (5'-3') | Referrence |
| β -actin F | CTGGAACGGTGAAGGTGA | CA 23 |
| β -actin R | AAGGGACTTCCTGTAACAA | TGCA |
| HIF-1α F | TGGTGACATGATTTACATT | гстд <mark>а́</mark> 4 |
| HIF-1α R | AAGGCCATTTCTGTGTGTA | AGC |
| LC3B F | TGTCCGACTTATTCGAGAG | CAGC ²⁵ |
| LC3B R | TTCACCAACAGGAAGAAG | GCCTGA |
| Beclin-1 F | ATGCAGGTGAGCTTCGTGT | G ²⁶ |
| Beclin-1 R | CTGGGCTGTGGTAAGTAAT | GGA |
| ABCC10 F | CGGGTTAAGCTTGTGACAC | GAGC 27 |
| ABCC10 R | AACACCTTGGTGGCAGTG | AGCT |
| AMPK F | CAGGCATATGGTGGTCCAT | AGAG ²⁸ |
| AMPK R | TCATGGGATCCACCTGCAG | C |
| PI3K F | TGTGTGGGGACTTATTGAGG | TG 22 |
| PI3K R | ACACAGTATCCAGCACATG | AAC |
| AKT F | TCTATGGCGCTGAGATTGT | G 29 |
| AKT R | CTTAATGTGCCCGTCCTTG | Т |
| | | |

Abbreviations: F: Forward, R: Reverse

hours (**Figure 1C**). In contrast, BCSCs exposed to $CoCl_2$ concentrations below 100 μ M for 24 hours show a generally reduced survival rate, but it is not significantly different from the negative control (**Figure 1D**).

At low concentrations of CoCl₂ (25 to 100 μ M), BC-SCs demonstrated a significant rise in *HIF-1* α gene expression. Notably, exposure to 100 μ M CoCl₂ results in the most significant increase, with a 4.54-fold elevation in *HIF-1* α gene expression compared to cells not treated with CoCl₂ (Figure 1E).

Autophagy is stimulated under hypoxic condition

The hypoxic environment caused by adding $CoCl_2$ strongly induced autophagy in BCSCs. In fact, in the presence of $CoCl_2$, *LC3B*, a central marker of autophagy induction, exhibits increased gene expression in a concentration-dependent manner. As $CoCl_2$ concentration rises, *LC3B* gene expression increases from a 1.45 to a 4.09-fold change compared to untreated cells. The expression of the *Beclin-1* gene, another autophagy marker, also increases at $CoCl_2$ concentrations of 50, 75, and 100 μ M (1.49, 2.28,













Abbreviations: 3-MA: 3-Methyladenine, BCSCs: Breast cancer stem cells, *p-value < 0.05

and 1.64-fold change, respectively), but it shows no significant change at 25 μ M (p > 0.05). Overall, both genes display increased expression with higher CoCl₂ concentrations, although these changes are not statistically significant (p > 0.05) (**Figure 2A**). In general, genes involved in autophagy regulation exhibit elevated expression levels when exposed to various CoCl₂ concentrations (**Figure 2B**). The AMPK gene shows the greatest expression increase at 50 μ M CoCl₂ (3.14-fold change), while PI3K and AKT genes reach their peak expression at 75 μ M CoCl₂ (2.69 and 2.14-fold change, respectively). After reaching peak expression, the levels of these genes gradually decline

with further increases in CoCl₂ concentration.

The inhibition of autophagy did not reduce the drug resistance of BCSCs

The potential for drug resistance in BCSCs was assessed using a cytotoxicity assay with the anti-tumor drug cisplatin. The drug resistance of hypoxic BCSCs to cisplatin was evaluated either in the presence or absence of the autophagy inhibitor 3-MA in the culture medium. In this assay, BCSCs were first induced to become hypoxic by culturing them in a medium containing CoCl₂ (ranging from 25 to 100 μ M) for 24 hours. Subsequently, these hypoxic BCSCs were cultured in a fresh medium containing 60 $\mu\mathrm{M}$ cisplatin for the next 24 hours.

The AlamarBlue assay results indicated that the viability of hypoxic BCSCs increased with the concentration of cobalt (II) chloride, although the increase was not significant. Concurrent treatment with 60 μ M cisplatin and 5 mM autophagy inhibitor 3-MA also enhanced the viability of the BCSCs population in relation to the concentration of CoCl₂.

The results showed that, whether treated with or without 3-MA, the cell viability of BCSCs remained similar and peaked at 75 μ M CoCl₂ (Figure 3).



Figure 4: The expression of the drug resistance gene *ABCC10* increased when breast cancer cell populations were treated with gradually increasing concentrations of CoCl₂ compared to BCSCs not treated with CoCl₂. *p-value < 0.05

Hypoxia enhanced the expression of ABCC10

Treatment with CoCl₂ results in an upregulation of the drug resistance gene *ABCC10*. The most significant increase in expression, observed as a 9.17-fold change, occurs at a concentration of 75 μ M CoCl₂ compared to breast cancer cells not exposed to CoCl₂ (see **Figure 4**).

DISCUSSION

The Cluster of Differentiation 44 (CD44) is a complex transmembrane adhesion glycoprotein involved in numerous signaling pathways significant in carcinogenesis and tumor progression³⁰. Meanwhile, Cluster of Differentiation 24 (CD24) serves as a crucial biomarker, indicating poor prognoses like increased invasion, metastasis, and reduced survival in

cancer patients³¹. The enzyme Aldehyde dehydrogenase (ALDH1) catalyzes the conversion of intracellular aldehydes to carboxylic acids and is expressed highly in a small subset of breast tumor cells capable of tumor generation in implanted mice³². Breast cancer cells displaying the CD44⁺CD24^{-/low}ALDH1^{high} phenotype are considered cancer stem cells due to their association with poorer outcomes in breast cancer patients³³. In this study, breast cancer cells isolated from Vietnamese patients were cultured in DMEM with 10% FBS to expand their population. Flow cytometry sorting was then used to purify cells with high CD44 expression and low CD24 expression. As such, CD44^{high}CD24^{low} breast cancer cells exhibiting elevated ALDH1 and EpCAM can be classified as breast cancer stem-like cells (BCSCs).

Cobalt (II) chloride (CoCl₂) has been used previously to modulate HIF expressions in various cancer³⁴⁻³⁶ and non-cancer cell lines^{37,38}. Past studies commonly used CoCl₂ concentrations over 100 μ M; however, our research demonstrated that BCSC proliferation was significantly inhibited at concentrations above 100 μ M for extended durations (24 to 72 hours). Consequently, we opted for lower concentrations of CoCl₂, under 100 µM, administered for shorter times (4 to 24 hours). The AlamarBlue assay revealed reduced BCSC proliferation, with no significant difference compared to cells cultured without CoCl₂. Similarly, a Thai study reported comparable outcomes using CoCl₂ concentrations of 50 or 100 μ M³⁹. To prevent extensive cell death affecting other evaluations, we employed CoCl2 concentrations ranging from 25 to $100 \,\mu$ M in subsequent experiments.

Autophagy under controlled hypoxia promotes cancer cell survival; however, under severe hypoxic stress, it can lead to cell death⁴⁰. In normoxic conditions, hypoxia-inducing factors (HIFs) are deactivated as prolyl hydroxylase enzymes hydroxylate them using oxygen. In hypoxia, HIFs stabilize and translocate to the nucleus, activating oncogenes⁴¹. Among HIF isoforms, HIF-1 α is the most studied due to its mRNA levels increasing with the severity of hypoxia in many cancers ^{42–44}. Thus, HIF-1 α can serve as a biomarker for evaluating tumor hypoxia levels. Using RT-qPCR, we found that CD44^{high}CD24^{low} breast cancer cells exhibited increased HIF-1 α expression correlating with rising CoCl₂ concentrations. Various mechanisms involving cobalt ions (Co²⁺) enhancing HIF- 1α expression have been suggested, generally affecting cofactors, inhibiting PHDs, and stabilizing HIF- $1\alpha^{45}$. Higher CoCl₂ concentrations in experimental models reflect a more pronounced hypoxic state in cancer tumors.

Beclin-1 is essential for signaling pathways activating autophagy and autophagosome formation, requiring interaction with hVsp34, while LC3B, a lipidated form of microtubule-associated protein lightchain 3, is a reliable autophagy marker, increasing in response to stressors⁴⁶. Our study found that Beclin-1 and LC3B gene expressions increased with rising CoCl₂ concentrations, indicating strong autophagy induction by this agent, similar to results in other cell lines⁴⁷. The observed increase in HIF-1 α expression suggests a link between autophagy and hypoxic response in BCSCs. According to Chen et al., $CoCl_2$ exposure increased HIF-1 α and its downstream protein BNIP3, involved in autophagy induction through interaction with Beclin-1 and LC3⁴⁷. The AMPK/PI3K/AKT pathway is implicated in autophagy across mammalian cell lines, typically with AMPK activation and increased expression, while PI3K and AKT expressions decrease under autophagy-inducing agents⁴⁸⁻⁵⁰. However, our results showed all three genes increased expression in BCSCs exposed to 25-75 µM CoCl₂, decreasing at 100 µM. Prior research also indicated mTOR, downstream of PI3K and AKT, plays a key role in autophagy regulation, with reduced PI3K and AKT expressions potentially inhibiting mTOR and initiating autophagy⁵¹. AMPK activation relates to numerous autophagy process steps, given its involvement in several autophagy factors⁵². Variances in gene expressions within BCSCs suggest autophagy regulation through alternate signaling pathways, necessitating further investigation into AMPK/PI3K/AKT and autophagy relationships.

Autophagy links to drug resistance in different cancer cell lines. Experiments with cisplatin on CD44^{high}CD24^{low} BCSCs, treated with CoCl₂ to simulate hypoxia and stimulate autophagy, showed increased cell survival correlating with CoCl₂ concentrations. This survival increase aligned with changes in autophagy-related gene expressions and ABCC10, a gene associated with drug resistance in breast cancer stem cells²⁷. We earlier posited that autophagy acts as a "double-edged sword" in drug resistance among cancer stem cells, potentially enhancing sensitivity or decreasing responsiveness to chemotherapy drugs based on the signaling pathway involved 53. Thus, targeting autophagy may offer a strategy to combat drug resistance in cancer cells. Numerous studies revealed that inhibiting autophagy through chemicals or genetic alterations enhances apoptosis in chemotherapy-resistant cell lines⁵⁴⁻⁵⁷. We hypothesize that autophagy suppression increases sensitivity to cisplatin in CD44^{high}CD24^{low} BCSCs. Using 5

mM 3-MA, an autophagy inhibitor, we observed significant *LC3B* expression reduction compared to untreated cells (data not shown). Nevertheless, the response to cisplatin remained unchanged under 3-MA influence, suggesting a weaker linkage between autophagy and cisplatin response in our cell line compared to others. Hence, more specific methods are required to evaluate signaling pathways linking these properties in Vietnamese breast cancer stem-like cells.

CONCLUSIONS

BCSCs exhibit increased expression of the HIF-1 α gene when exposed to higher concentrations of cobalt (II) chloride (CoCl₂), which also leads to the upregulation of genes involved in the induction (Beclin-1, LC3B) and regulation (AMPK, PI3K, AKT) of autophagy. Under hypoxia-mimicking conditions, BC-SCs show increased resistance to cisplatin. However, inhibiting autophagy with the inhibitor 3-MA does not reduce the drug resistance of hypoxic BCSCs. Furthermore, under these conditions, an increase in the expression of the drug resistance gene ABCC10 is observed in response to CoCl2 levels. These findings suggest that the drug resistance of BCSCs under hypoxia is related to both autophagy upregulation and ABCC10 expression. Therefore, addressing drug resistance in BCSCs should consider both autophagy and ABCC10 pathways.

ABBREVIATIONS

3-MA: 3-Methyladenine, *ABCC10*: ATP-Binding Cassette Subfamily C Member 10, **AKT**: Protein Kinase B, **AMPK**: AMP-Activated Protein Kinase, **ANOVA**: Analysis of Variance, **BCSC**: Breast Cancer Stem Cell, **CoCl**₂: Cobalt (II) Chloride, **CSC**: Cancer Stem Cell, **DMEM**: Dulbecco's Modified Eagle Medium, **FBS**: Fetal Bovine Serum, **HIF**: Hypoxia-Induced Factor, **HIF-1** α : Hypoxia-Induced Factor 1 alpha, *LC3B*: Microtubule-Associated Proteins 1A/1B Light Chain 3B, **mRNA**: Messenger Ribonucleic Acid, **PI3K**: Phosphoinositide 3-Kinase, **RT-qPCR**: Real-Time Quantitative Polymerase Chain Reaction, **SEM**: Standard Error of the Mean, **VNBRCA1**: Vietnamese Breast Cancer Cell Line No. 1

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AUTHOR'S CONTRIBUTIONS

Bui Dinh Khan and Tran Ngo The Nhan equally contributed to developing the research methodology, conducting the experimental work, and collecting the data. Bui Dinh Khan took primary responsibility for data analysis and manuscript content. Pham Van Phuc suggested the layout, discussed the results, revised entire manuscript, and supervised the study. All authors contributed significantly to this work and have read and approved the final version of the manuscript.

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AVAILABILITY OF DATA AND MATERIALS

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

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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