Autophagy/ Apoptosis Induced by Geraniol Through HIF-1α/BNIP3/Beclin-1 Signaling Pathway in A549 CoCl₂ Treated Cells

Dina M. Abo El-Ella *

*Department of Pharmacology and Toxicology, Faculty of Pharmacy, October 6 University, 6 October City, 12566, Giza, Egypt

*Corresponding author. O6U, October 6 city, 12566, Giza, Egypt.

E-mail: Dinamoustafa@o6u.edu.eg

Telphone: +20-012-22760673

Running title:

Autophagy and Geraniol on A549 CoCl₂ Treated Cells

Abstract:

Background:

During cancer growth, hypoxia occurs along with autophagy as an adaptive response to overcome cellular stress. Geraniol is a natural isoprenoid known for its wide anticancer activity and autophagy induction in the cancer cell.

Purpose:

To investigate the antihypoxic potential of geraniol with the incidence of autophagy and apoptotic cell death in A549 CoCl₂ treated cells.

Methods:

A549 cells were incubated for 24 h with GE and CoCl₂ either alone or in combination. We examined the cytotoxicity and cell viability of GE either alone or in combination therapy using MTT and trypan blue assay. GE modulating effect was determined on lipid peroxidation, antioxidant capacity markers, gene expression levels of HIF-1α, NF-kB, VEGF, autophagy factors in different groups, besides apoptotic bodies using AO/EB.

Results:

GE and CoCl₂ combination therapy downregulated the expression of HIF-1α that suppressed A549 cell growth through downregulation of BNIP3 and Beclin-1 gene expression. This resulted in autophagy and
apoptotic cell death, in addition to the downregulation of NF-kB and VEGF expression. Also, GE treatment significantly reduced the oxidative stress markers and restored the antioxidant capacity.

Conclusion:

GE possesses an antihypoxic effect on A549 CoCl$_2$ treated cells and induces cell death via autophagy along with apoptosis through HIF-1α/BNIP3/Beclin-1 signaling pathway.

Key words:

Hypoxia, Geraniol, HIF-1α, Autophagy, BNIP3, Beclin-1, VEGF

Abbreviation:

GE, Geraniol; MDA, malondialdehyde; TAC, total antioxidant capacity; NF-kB, nuclear factor kappa-light-chain-enhancer of activated B cells; CoCl$_2$, Cobalt chloride; HIF-1α, hypoxia-inducible factor-1 alpha; BNIP3,Bcl-2 19KD interacting protein; Beclin-1, mammalian orthologue of yeast Atg6; VEGF, Vascular Endothelial Growth Factor.

1. Introduction

Lung cancer is one of the leading causes of death worldwide. This is due to the formation of wide areas of hypoxic tissue that reduce the response to radiotherapy and chemotherapy. Therefore, developing anti-hypoxic drugs is the main aim of lung cancer therapy by targeting hypoxia, concurrent sequences of angiogenesis, invasiveness, and tumor growth.

Hypoxia and necrotic areas are developed from insufficient oxygen supply during solid tumor growth. Hypoxia inducible factor-1 (HIF-1) is the main cellular regulator during hypoxia, composed of oxygen regulated HIF-1 α subunit as well as the constitutively expressed HIF-1 β subunit. HIF-1 α role during hypoxia is to maintain blood, nutrients as well as energy resumption through the expression of regulatory genes controlling glycolysis, angiogenesis, and erythropoiesis. Recent studies reported a bi-directional relation between HIF-1 α and NF-κB because HIF regulates the expression of NF-κB mRNA and vice versa.

Autophagy is a self-eating and dynamic phenomenon that depends on lysosomal degradation of the damaged mitochondria, cytoplasm, lipids, and misfolded proteins back to their basic components with cellular recycling to preserve cellular homeostasis. Autophagy is present at a low level under physiological conditions but increased under cellular stress ailments including hypoxia, and nutrient deficiency and ROS production. Autophagy and apoptosis share many regulators and interactions like Beclin-1/Bcl-2, and caspase/Beclin-1 cleavage. During moderate hypoxic conditions, HIF-1α induces activation of autophagy in a dependent manner and supplies cancer cells with cellular energy requirement for solid tumor proliferation. However, severe hypoxia induces autophagy in HIF-1α independent manner through involvement of the AMPK-mTOR and unfolded protein response pathways. BNIP3 and Beclin-1(Bcl-2 gene family) are cell death proteins that play an important role in hypoxia associated apoptosis, necrosis, and autophagy. Thus, BNIP3 and Beclin-1 are markers for appraising autophagy.

Geraniol is a natural isoprenoid monoterpenoid derived from the essential oils of various aromatic plants including lemon, ginger, rose, and orange. Previous in vitro and in vivo studies showed that GE possesses anti-microbial, anti-inflammatory, and anti-oxidant properties.
Moreover, GE exerts an apoptotic, antitumor, and anti-angiogenic effect on various cancer cells,\textsuperscript{24-26} besides being a potent inducer of autophagy.\textsuperscript{27}

Taking these previous pharmacological effects into account, this study was designed to determine the protective effect of GE against CoCl\textsubscript{2} induced hypoxia associated autophagy on the A549 cell line and focused on the relation between HIF-1\textalpha, autophagy, and apoptotic cell death.

2. Materials and methods
2.1. Media and reagents:
All chemicals and reagents were obtained from Sigma Aldrich Chemical Co. (St. Louis, MO, USA) unless otherwise stated during this study. This study was achieved at the Cell Culture Research Unit on October 6 University (Cairo, Egypt).

2.2. Cell line and cell culture:
Human lung adenocarcinoma epithelial A549 cells (ATCC, USA) were cultured according to the American Type Culture Collection protocol.\textsuperscript{28} All experiments were repeated three times independently, and the data were represented as the mean ± SD.

2.3. MTT assay:
A549 cells were grown to 90% confluence then undergo trypsinization followed by counting with a hemocytometer. Afterward, A549 cells were seeded in 96 well tissue culture plates at 10x 10\textsuperscript{3}/well in triplicates for 24 hours for cell attachment before further additions. The A549 cells were treated with various concentrations of GE (0.161 to 2.59 mM) and CoCl\textsubscript{2} (0.193 to 7.75mM) either alone or in combination then incubated for 24 hours.\textsuperscript{29,30} After treatment, the standard 3,4,5-dimethylthiazol-2,5- diphenyl tetrazolium bromide (MTT) method was used to assess the cytotoxicity of GE alone or its combination with CoCl\textsubscript{2}.\textsuperscript{31} The IC\textsubscript{50} concentration of GE and CoCl\textsubscript{2} either alone or in combination on the A549 cell line after 24 h were determined and used to assess other bioassays. The mean percentages of cell viability were detailed as mean ± SD.

2.4. Cell viability by the trypan blue dye exclusion assay:
Cell viability of A549 cells in treated and untreated groups was determined by the trypan blue assay that depends on quantify live cells by labeling dead cells exclusively. The A549 cells were stained with 10μL trypan blue dye (0.4% solution) and to count unstained live and blue stained dead cells under a phase contrast microscope with a hemocytometer.\textsuperscript{32} The mean percentages of live and dead cells per experiment were expressed as mean ± SD of three autonomous experiments.

2.5. Apoptosis and necrosis Staining:
Acridine orange/ethidium bromide staining (AO/EB) used to detect the mode of cell death.\textsuperscript{33} The mean percentages of live, apoptotic, and necrotic cells per experiment were determined according to Zakaria and her colleagues and were expressed as mean ± SD.\textsuperscript{34}

2.6. Cell lysate preparation:
The preparation of cell lysate was done according to Zakaria and her colleagues to be used in the analysis of oxidative stress and inflammatory markers.\textsuperscript{34} Bicinchoninic acid used to determine the protein content of total cell lysate using bovine serum albumin as a standard, data not cited.\textsuperscript{35}

2.7. Estimation of oxidative stress markers (MDA and TAC):
Lipid peroxidation of cell lysate of treated and untreated groups was determined using malondialdehyde (MDA) as a representative of its final product using a commercially supplied kit (Biodiagnostic).\textsuperscript{36} Also, the antioxidant capacity (TAC) of the cell lysate of different groups was assayed colorimetrically according to the protocol of the Biodiagnostic supplied kit.\textsuperscript{37}

2.8. Gene expression analysis using quantitative real-time polymerase chain reaction (qRT-PCR):
qRT-PCR was used to quantify the changes in HIF-1α, NF-κB, VEGF, BNIP3, and Beclin-1 mRNA expression in different groups. In ΔΔCT method, GAPDH mRNA was used as the internal control to detect the variation in the genes expression of HIF-1α, NF-κB, VEGF, BNIP3, and Beclin-1 in all groups. All primer sequences used in amplifying the HIF-1α, NF-κB, VEGF, BNIP3, and Beclin-1 genes are listed in table 1.

Table 1:
Primer sequences used in qRT-PCR analysis

<table>
<thead>
<tr>
<th>Target Gene</th>
<th>Primer Sequences</th>
</tr>
</thead>
<tbody>
<tr>
<td>GAPDH</td>
<td>F: CTCTGATTGTTGTCGTATTTG&lt;br&gt;R: TGGAAAGATGGTAGTGGATTTG</td>
</tr>
<tr>
<td>NF-κB</td>
<td>F: TGTTGCGCTACTGCTAACT&lt;br&gt;R: GGATGCACTTCAGCTTCTGT</td>
</tr>
<tr>
<td>HIF-1α</td>
<td>F: ATCCATGTGACCATGAGGAATG&lt;br&gt;R: TCGGCTAGTTAGGGTACACTTC</td>
</tr>
<tr>
<td>VEGF</td>
<td>F: AGGGCAGAATCATCACGAAGT&lt;br&gt;R: AGGGTCTCGATTGGATGGCA</td>
</tr>
<tr>
<td>BNIP3</td>
<td>F: CAGGCTCTGGTTAGAAGAC&lt;br&gt;R: CTACTCCGTCGATAGACATTC</td>
</tr>
<tr>
<td>Beclin-1</td>
<td>F: GCGTGAAGACTGGATAGG&lt;br&gt;R: CTGGGTCTGGGCATAACG</td>
</tr>
</tbody>
</table>

2.9 Statistical analysis:
Experimental data were analyzed by Graph- Pad Prism (ISI®, USA) software (version 5) and expressed as mean ± standard deviation (SD). One-way ANOVA followed by Tukey Kramer as a post hoc test was used to detect the variation between groups. A P-value of less than 0.05 was considered a statistically significant difference. (*) is significant compared with the A549 cells, and (#) is significant compared with the CoCl₂ treated group.

3. Results:
3.1 Cytotoxicity of GE and CoCl₂ either alone or in combination
A significant decrease in cell viability in the CoCl₂ (hypoxic) and GE groups was observed compared to untreated cells and the calculated IC₅₀ was 6.2 mM and 2.59 mM subsequently, as shown in Fig. 1a-b. The effect of combined therapy was assessed on the viability of A549 using the IC₅₀ of CoCl₂ added to various concentrations of GE for 24 h. As shown in Fig.1c, the combination therapy of CoCl₂ + 2.59 mM GE showed a significant reduction in cell viability (26 % ± 9) compared to untreated cells. Contrary to this, CoCl₂ + 1.29 mM of GE showed the least reduction in cell viability. Thus, the IC₅₀ value of GE (2.59 mM) and its diluted concentration ½ IC₅₀ 1.29 mM will be used in the combination therapy with IC₅₀ of CoCl₂ for 24 h in the following bioassays.

3.2 Trypan blue dye exclusion assay
As shown in Fig. 2, A549 cells treated with GE (2.59 and 1.29 mM) showed a significant decrease in the number of viable cells 48% ± 2.6, and 46% ± 3.2 respectively regarding that noticed in untreated cells (98% ± 0.6). Also, CoCl₂ A549 treated cells (6.2 mM) either alone or in combination, showed a major reduction in the number of viable cells (60% ± 2.6, 25% ± 3.8, and 32% ± 2.1 respectively) compared to that observed in untreated cells.
3.3. Apoptosis and necrosis staining

Under a fluorescence microscope, dual DNA staining by AO/EB dyes was used to determine the mode of cell death in untreated and treated A549 cell line. The untreated group showed no sign of cell death and most of the cells are viable with uniform green fluorescence color (Fig.3a-b). CoCl₂ treated group showed a high number of yellow to orange stained cells with a mean percentage of apoptotic cell death of 44%± 1.5 compared to that noticed in untreated cells (93% ± 1) (Fig.3a-c). Groups treated with GE (2.59 and 1.29 mM) and CoCl₂ showed a significant percentage of apoptotic cell death (85% ± 2 and 75% ± 2.5 respectively) compared to that detected in CoCl₂ treated group as well as few necrotic cells that display orange to red nuclei (Fig.3d). Moreover, GE treated groups showed a meaningful percentage of apoptotic cells (50%± 1 and 40%± 1.5 respectively) with bright green to yellow color compared to untreated groups (Fig.3e).

3.4. Oxidative stress marker and antioxidant activity

Combined therapy of GE (2.59 and 1.29 mM) and CoCl₂ (6.2 mM) on A549 for 24 h showed a significant reduction in lipid peroxidation in terms of MDA level (1.9 ± 0.1 and 2.8 ± 0.6 respectively) compared to that observed in CoCl₂ treated cells (34.4 ± 0.75); MDA is an eminent detector for oxidative stress (Fig. 4a). On the other hand, total antioxidant capacity in untreated and treated groups was determined to detect the effect of ROS generation on antioxidant system. Combination therapy of GE (2.59 and 1.29 mM) and CoCl₂ as well as GE treated groups showed a significant elevation in the TAC levels (0.161 ± 0.01, 0.173 ± 0.007, 0.16 ± 0.003, and 0.169 ± 0.003 respectively) compared to that noticed in CoCl₂ treated cells (0.11 ± 0.01) as presented in Fig. 4b.

3.5. Relative expression of HIF-1α, NF-κB, VEGF, BNIP3, and Beclin-1 mRNA in different groups:

As shown in Figs. 5a-e, CoCl₂ treated group showed a significant upregulation in the mRNA expression levels of HIF-1α (9.4 fold), NF-κB (11.74 folds), VEGF (8.46 folds), BNIP3 (5.3 folds), and Beclin-1 (2.7 folds) compared to untreated groups, indicating a significant incidence of hypoxia, autophagy as well as angiogenesis. On the other hand, Combined therapy of GE (2.59 and 1.29 mM) and CoCl₂ showed a significant downregulation in the mRNA expression levels of HIF-1α (6.4 and 7.15 folds respectively), NF-κB (5.1 and 7.4 folds respectively), VEGF (6.1 and 6.9 folds respectively), and BNIP3 (3.1 and 2.5 folds respectively) as well as a modest downregulation in the mRNA of Beclin-1 expression (2 and 1.9 folds respectively) compared to CoCl₂ treated group. It is worthwhile to mention that GE treated groups showed a significant upregulation in the mRNA expression levels of BNIP3 (1.95 and 2.05 folds respectively), and Beclin-1 (4.54 and 2.01 folds respectively) compared to that noticed in untreated groups.

4. Discussion

Exploration of antihypoxic agents from plant origin has drawn great attention these years in cancer research. From MTT results, GE induced cytotoxicity and loss of A549 cell viability at a concentration of 2.59 mM (IC₅₀ value). Yet, Galle and his colleagues reported that the IC₅₀ value of GE was 797.2 μM. Also, CoCl₂ is a hypoxic imitator agent that induces structural modification in the heme protein O₂ sensor that results in the generation of ROS, which provokes oxidative stress resulting in hypoxia-induced cytotoxicity. In this study, the IC₅₀ value of CoCl₂ was 6.2 mM. But Mahey and her colleagues reported that 29.81 mg/l (231 mM) is the IC₅₀ value of CoCl₂.

ROS formation in the mitochondria during hypoxia aggravates the oxidative stress, induces excessive autophagy, and disrupts the oxidant/antioxidant balance within the cells. This study showed that the IC₅₀ value of CoCl₂ (6.2 mM) disrupted the oxidant/antioxidant balance in A549 cells, which results in the incidence of lipid peroxidation with a subsequent reduction in the TAC level. This finding further corroborates earlier observations of Tripathi et al as well as Mohamed et al. GE (2.59 and 1.59 mM) possesses antioxidant capacity when combined with CoCl₂ by reducing the level
of MDA parallels with the elevation in the TAC level. This finding is similar to that of Khan and his colleagues. Abbreviated orthography.

The antioxidant activity of GE results from increasing endogenous antioxidant defense system through the elevation in glutathione peroxidase and superoxide dismutase activity via prompting their transcriptional up-regulation.

HIF-1α is released during hypoxia to maintain oxygen homeostasis besides the upregulation of transcriptional factor NF-κB; both are factors of stress response to tumor-associated hypoxia. Recent studies reported a bi-directional relation between HIF-1α and NF-κB because HIF regulates the expression of NF-κB mRNA and vice versa. Also, HIF and NF-κB activate the expression of several genes controlling angiogenesis (VEGF), cytokines (TNF-α), chemokines (IL-8), and cell death related proteins (Noxa and BNIP3). Under stress condition such as hypoxia, NF-κB induces autophagy and inflammation through activation of mTOR, which in turn stimulates HIF-1α and NF-κB transcriptional activation results in a sequence of hypoxic and inflammatory cascades that ends in autophagy and inflammation. Together, these findings further support the results of our study where CoCl2 treated cells showed a rapid upregulation in mRNA of HIF-1α, NF-κB, VEGF, BNIP3, and Beclin-1 expression. Accordingly, these results indicated the rapid induction of hypoxia with angiogenesis that is concurrent with autophagy, and this agrees with other researchers using a similar approach. The most consistent effect of CoCl2 to accumulate HIF-1α is through the direct binding of cobalt with oxygen-dependent degradation domain of HIF-α, followed by the suppression of hydroxylated HIF-1α to be produced, as well as its interaction with von Hippel-Lindau protein. Also, transcriptional activation of HIF-1α induced by CoCl2 results in the direct activation of BNIP3 (hypoxia response element), induced programmed cell death as well as regulates autophagy through hindering the binding between Bcl-2 and Beclin-1.

GE is a natural product that blocks the modification of IkB which results in the inhibition of NF-κB activation as well as its transcriptional activation in TPA treated mouse skin. Also, GE possesses a strong antiangiogenic, anti-inflammatory, antiproliferative, and apoptosis effect through the downregulation of VEGF, NF-κB, cyclin D1, PCNA, c-fos, p53 and Bcl-2 in buccal pouch carcinogenesis. Moreover, GE suppresses the AKT signal with the activation of the AMPK pathway, followed by the suppression of mTOR phosphorylation that results in autophagy in PC-3 cells. Furthermore, GE elevated the BNIP3 and BAX (Bcl-2 family members, apoptotic markers) as well as downregulated the cell cycle regulators (cyclins and CDKs) in PC-3 cells. Inhibition of HIF-1α can suppress autophagy by reducing the amount of LC3-II and LC3-I in oral squamous cell carcinoma. Together, these findings further support the results of this study. Combination therapy of GE (2.59 and 1.59 mM) with CoCl2 treated groups showed a significant upregulation in BNIP3 and Beclin -1 expression. This study showed that CoCl2 treated group induced a significant increase in the percentage of apoptotic cells compared to that of necrotic cells as well as upregulation in the expression levels of BNIP3 and Beclin-1. Cobalt ion possesses a cytotoxic activity and induces hypoxia followed by the upregulation and downregulation of apoptotic and anti-apoptotic.
proteins subsequently, which stimulates apoptotic cell death. Also, Chen et al showed that CoCl$_2$ upregulates the HIF-1$\alpha$ as well as autophagic proteins (LC3, BNIP3, and Beclin-1) in the C2C12-hypoxic model. These reports and our current results confirmed the link between CoCl$_2$, hypoxia, autophagy signaling, and apoptotic cell death.

Moreover, GE in the combination therapy with CoCl$_2$ showed a significantly higher percentage of apoptotic cells and a down regulation in the BNIP3/Beclin-1 level. Also, GE alone showed a significant percentage of apoptotic cells, an upregulation in the BNIP3/Beclin-1 level. Our results corroborate the observations of Kim and his colleagues who reported the GE induced autophagy and apoptotic cell death in PC-3 cell. BNIP3 (Bcl-2 19KD interacting protein) is a mitochondrial outer membrane protein that is activated under hypoxia by the action of HIF-1 to clear the impaired mitochondria; therefore driving autophagy by acting as a mitophagy receptor. Furthermore, BNIP3 can avert the connection between Beclin-1 and Bcl-2, freeing Beclin-1 to hasten the autophagic process through autophagosome formation. This in turn induces apoptosis by increasing mitochondrial membrane permeability. Also, Beclin-1 overexpression by an anticancer drug resulted in apoptotic cell death in cervical cancer cells.

5. Conclusion
In this study, we assessed the antihypoxic role of GE (2.59 and 1.29 mM) against CoCl$_2$ induced hypoxia in the A549 cell line which relies on restoring pro-oxidant / antioxidant equilibrium. Also, GE modulates the inflammatory and angiogenic markers through downregulation of the expression levels of NF-$\kappa$B and VEGF that results from the upregulation of HIF-1$\alpha$ during hypoxia. Additionally, GE reduces autophagy in combination therapy by downregulating the expression levels of BNIP3 and Beclin-1, associated with an elevation in apoptotic cell death through HIF-1$\alpha$ signaling pathways. Further experiments will be needed to explore the mechanism of GE in controlling hypoxia-associated autophagy at the molecular and cellular levels in different cell types.

6. Study highlights:

Current knowledge:
GE exerts an apoptotic, antitumor, and anti-angiogenic effect on various cancer cells, besides being a potent inducer of autophagy. No previous study mentioned the relation between GE and HIF-1$\alpha$ in hypoxia and how it connected to autophagy through the BNIP3/Beclin-1 signaling pathway.

New findings:
- GE reduces the oxidative stress induced by CoCl$_2$ in A549 cells by reducing MDA and increasing TAC content.
- GE and CoCl$_2$ combination therapy downregulate the expression of HIF-1$\alpha$ that in turn suppresses the A549 cell growth through downregulation of BNIP3 and Beclin-1 gene expression, results in autophagy and apoptotic cell death.
- GE and CoCl$_2$ combination therapy reduce the expression levels of inflammatory (NF-$\kappa$B) and angiogenic factors (VEGF).

7. Declarations:
Funding: This research received no specific grant from any funding agency in the public, commercial, or not-for-profit sectors.

Conflict of interest: The authors declare no potential conflicts of interest.

Author contributions: Dina M. Abo El-Ella designed experiments, performed the experiments, wrote the paper, revised and finalized the manuscript.
Ethical Issues: Not applicable.

8. References:


32. Tripathi VK, Subramaniam SA, Hwang I. Molecular and Cellular Response of Co-cultured Cells toward Cobalt Chloride (CoCl₂)-Induced Hypoxia. *ACS Omega* 2019;4(25):20882-20893. doi:10.1021/acsomega.9b01474


The manuscript will undergo copyediting, typesetting, and review of the resulting proof before it is published in its final form.
Fig 3

a. Bar graph showing the cell population (%): Necrotic cells, Apoptotic cells, and Vital cells across different conditions:
- A549 cells
- 6 mM CoCl2
- CoCl2 + 2.59 mM GE
- CoCl2 + 1.29 mM GE
- 2.59 mM GE
- 1.29 mM GE

b. Untreated A549 Cells
c. 6.2 mM CoCl2
d. Combination therapy
e. 2.59 & 1.29 mM GE
The manuscript will undergo copyediting, typesetting, and review of the resulting proof before it is published in its final form.

**Fig 5**

(a) HIF-1α folds of expression

(b) NF-kB folds of expression

(c) VEGF folds of expression

(d) BMP3 folds of expression

Legend:
- *: Statistically significant difference
- #: Statistically significant difference between groups