Synthesis of an anthraquinone derivative (DHAQC) and its effect on induction of G2/M arrest and apoptosis in breast cancer MCF-7 cell line

Abstract: Anthraquinones are an important class of naturally occurring biologically active compounds. In this study, anthraquinone derivative 1,3-dihydroxy-9,10-anthraquinone-2-carboxylic acid (DHAQC) (2) was synthesized with 32% yield through the Friedel–Crafts condensation reaction. The mechanisms of cytotoxicity of DHAQC (2) in human breast cancer MCF-7 cells were further investigated. Results from the 3-(4,5-dimethylthiazol-2-yl)-2,5-diphenyltetrazolium bromide (MTT) assay showed that DHAQC (2) exhibited potential cytotoxicity and selectivity in the MCF-7 cell line, comparable with the naturally occurring anthraquinone damnacanthal. DHAQC (2) showed a slightly higher IC50 (inhibitory concentration with 50% cell viability) value in the MCF-7 cell line compared to damnacanthal, but it is more selective in terms of the ratio of IC50 on MCF-7 cells and normal MCF-10A cells. (selective index for DHAQC (2) was 2.3 and 1.7 for damnacanthal). The flow cytometry cell cycle analysis on the MCF-7 cell line treated with the IC50 dose of DHAQC (2) for 48 hours showed that DHAQC (2) arrested MCF-7 cell line at the G2/M phase in association with an inhibited expression of PLK1 genes. Western blot analysis also indicated that the DHAQC (2) increased BAX, p53, and cytochrome c levels in MCF-7 cells, which subsequently activated apoptosis as observed in annexin V/propidium iodide and cell cycle analyses. These results indicate that DHAQC (2) is a synthetic, cytotoxic, and selective anthraquinone, which is less toxic than the natural product damnacanthal, and which demonstrates potential in the induction of apoptosis in the breast cancer MCF-7 cell line.

Keywords: cytotoxic, selective index, cell cycle

Introduction

Cancer is one of the most major health problems, according to any frame of reference, and, unfortunately, is not likely to lose that qualification any time soon. The breast is one of the four major cancer sites, and, of all cancers, breast cancer has the highest incidence rate and is the second highest cause of death in females.1 Thus, research in medicinal chemistry has been focused on the discovery of novel anticancer compounds, including natural compounds that can specifically target cancer cells and minimize the possible side effects on normal cells.2 Anthraquinones are potential natural anticancer compounds that can be isolated from Damnacanthus and Morinda spp. Among the natural anthraquinones, damnacanthal has been shown to possess potential cytotoxic, immunomodulatory,1 and anticancer activities.3 These cytotoxic and apoptotic activities against breast cancer cell lines were found to be regulated by damnacanthal via activation of p53 and p21 genes.3 Other naturally occurring anthraquinones have also been shown to exhibit similar anti-inflammatory, antibiotic, antiviral, and antineoplastic activities.4–6 However, due to the lack of accessibility, the time and cost of preparation, and the limited quantity of naturally occurring anthraquinones, chemically synthesized...
compounds have garnered a great deal of attention as a means of complementing the natural isolated compounds in anticancer drug discovery. Based on this connection, we synthesized the compound 1,3-dihydroxy-9,10-anthraquinone-2-carboxylic acid (DHAQC) (2) by Jones oxidation similar to the previously synthesized compound, 2-hydroxymethyl-1,3-dimethoxy-9,10-anthraquinone (1).

One of the major obstacles in anticancer compound discovery is that the candidate compounds could have nonspecific toxicity against both cancerous and normal cells. Thus, identification of a novel compound with high selectivity against cancerous cells rather than normal cells is one of the major objectives of this type of study. The current study aimed to evaluate and compare the selectivity of DHAQC (2) to damnacanthal on cancerous and normal cell lines. Moreover, the cell cycle arrest and apoptotic effects of DHAQC (2) on MCF-7 cells was also evaluated by quantitative polymerase chain reaction (PCR), Western blot, and flow cytometry analysis.

Materials and methods
Synthesis of DHAQC (2)
Damnacanthal was synthesized according to the methods of Akhtar et al. Compound (1) (2-hydroxymethyl-1,3-dimethoxy-9,10-anthraquinone) was synthesized as described in our previous publication. Briefly, the phthalic anhydride and 1,3-dihydroxy-2-methylbenzene were mixed in a molten mixture of AlCl₃/NaCl. Then, the product was acetylated with acetic anhydride and potassium carbonate. Next, the product underwent methylation with K₂CO₃/(CH₃)₂SO₄, which was further brominated with Wohl-Ziegler’s reaction. Finally, the product was hydrolyzed in acetic acid and water (8:2) to yield compound (1). For the synthesis of compound (2), DHAQC (2), compound (1) (500 mg, 1.7 mmol) was dissolved in 30 mL of acetone in a round bottle flask equipped with a CaCl₂ drying tube. The reaction mixture was stirred at 0°C–5°C in an ice bath. Jones reagent was prepared by mixing CrO₃ + H₂SO₄ at 0°C in acetone. After 2 hours of storage in a refrigerator, 5 mL of the Jones reagent was diluted with water and added slowly under nitrogen atmosphere until the yellow solution turned greenish. The product was treated with 5% solution of sodium bisulfite, extracted with ethyl acetate, and purified by column chromatography. The compound DHAQC (2) was analyzed by electron ionization mass spectrometry (EI-MS), infrared (IR), and nuclear magnetic resonance spectroscopic (NMR) techniques. Compound (2) was obtained as a yellow-green amorphous powder and purified by column chromatography. The yield was 32.0%. IR (cm⁻¹) in KBr disk: 3450-, 3220 (OH), 2973 (CH), 1646 (H-C=O), 1670 (non-chelated C=O), 1466 (C=C aromatic), 1244, 1262, 1230, 1132, 710. ¹H-NMR (500 MHz, acetone-d₆): δ 12.21 (s, OH), 8.22 (d, 1H, J=7.5 Hz, H-8), 8.09 (d, 1H, J=7.5 Hz, H-5), 7.75–7.72 (m, 2H, H-6, and H-7), 7.70 (s, 1H, H-4), MS m/z (relative intensity): 284.20 (M⁺, 242), 281 (45), 226 (56), 254 (79), 206 (7), 196 (24), 180 (57), 77, 65, 54.

Cell culture
Human erythromyeloblastoid leukemia K562 cells, estrogen-dependent breast cancer MCF-7 cells, estrogen-independent breast cancer MDA-MB-231 cells, and normal breast MCF-10A cells were purchased from American Type Culture Collection (ATCC) Manassas, VA, USA). K562 and MCF-7 cells were cultured in Roswell Park Memorial Institute (RPMI)-1640 (Sigma-Aldrich Co, St Louis, MO, USA) and supplemented with 10% fetal bovine serum (FBS) (GE Healthcare, Little Chalfont, UK), while MDA-MB-231 cells were cultured in Dulbecco’s Modified Eagle’s Medium (DMEM) (Sigma-Aldrich Co) supplemented with 10% FBS. MCF-10A cells were cultured in DMEM/F-12 (Sigma-Aldrich Co) supplemented with 10% FBS, hydrocortisone (0.5 μg/mL), insulin (10 μg/mL), and human epidermal growth factor (20 ng/mL) (Sigma-Aldrich Co). All cell lines were kept in a 37°C incubator with 5% CO₂.

3-(4,5-Dimethylthiazol-2-yl)-2,5-diphenyltetrazolium bromide (MTT) cell viability assay
MCF-7, MDA-MB-231, and MCF-10A cells were seeded in 96-well plates (0.8×10⁴ cells/mL) and cultured overnight. Then, starting at 30 μg/mL, serial 1:1 dilutions of the examined compounds were added to triplicate wells on the seeded plates. The nonadherent K562 cells were loaded (0.8×10⁵ cells/mL) on the 96-well plate, which contained the appropriate dilutions of compounds. All plates were incubated for 48 hours at 37°C, 5% CO₂. After the incubation period, 20 μL of MTT (5 mg/mL) solution (Sigma-Aldrich Co) was added to all wells and incubated in darkness for 3 hours. Then, 170 μL of solution was discarded and 20 μL of dimethyl sulfoxide (DMSO) was added to each well to solubilize the purple crystals. The plates were read using a microplate reader at 570 nm (BioTek Instruments, Inc., Winooski, VT, USA). IC₅₀ (inhibitory concentration with 50% cell viability) values, indicating the concentration of DHAQC (2) or damnacanthal that inhibited 50% of cell viability compared to untreated control, were obtained, and selective index was
calculated by dividing the IC$_{50}$ value of MCF-10A cells by the IC$_{50}$ value of MCF-7 or MDA-MB-231 cells.

**Cell preparation**

MCF-7 cells were seeded overnight and treated with the IC$_{50}$ value of DHAQC (2) for 24 hours for quantitative real-time PCR (qRT-PCR) and with IC$_{25}$ (inhibitory concentration that inhibits 25% of cell viability) (3.50 µg/mL) and IC$_{50}$ (7.20 µg/mL) of DHAQC (2) for 48 hours for flow cytometry, Western blot, and caspase-9 analysis. Untreated control cells were prepared simultaneously. After the treatment period, cells were harvested, washed, and subjected to the following assays.

**Flow cytometry cell cycle PI (propidium iodide) assay**

Treated and untreated MCF-7 cells were stained with BD CycleTest™ Plus DNA reagent kit (BD, Franklin Lakes, NJ, USA) according to the manufacturer’s protocol. The cell cycle phase was analyzed using BD CellQuest Pro software by FACSCalibur™ (BD, Franklin Lakes, NJ, USA).

**Flow cytometry annexin V/PI assay**

Treated and untreated MCF-7 cells were mixed with 100 µL of 1× binding buffer. Then, 5 µL of annexin V and PI, each from the Annexin V-FITC apoptosis detection kit (BD Biosciences, San Jose, CA, USA), were added to the cells and incubated for 15 minutes. Afterwards, stained cells were washed and mixed with 500 µL of 1× binding buffer, and the apoptosis of the cells was analyzed using BD CellQuest Pro software by FACSCalibur.

**qRT-PCR assay**

RNA from the treated and untreated MCF-7 cells was extracted using the RNeasy Plus Mini Kit (Qiagen NV, Venlo, the Netherlands) according to the manufacturer’s protocol. The RNA quality was quantified using a NanoDrop spectrophotometer (Thermo Fisher Scientific, Waltham, MA, USA). The RNA was converted to complementary (c)DNA using an iScript™ cDNA synthesis kit (Bio-Rad Laboratories Inc., Hercules, CA, USA). Expression of p21, PLK1, and FOXM1 were evaluated by qRT-PCR with an SYBR® Select Master Mix (Thermo Fisher Scientific, Waltham, MA, USA) on an iQ-5 5 Real Time PCR (Bio-Rad Laboratories Inc.) using the primers as stated in Table 1 and the following PCR conditions: one cycle of 50°C for 2 minutes for UDG (uracil-DNA glycosylase) activation; one cycle of 95°C for 2 minutes for DNA polymerase activation; 40 cycles of 95°C for 2 seconds for denature; and 60°C for 30 seconds for annealing and extension. Standard curves for housekeeping genes (β-actin, 18s rRNA, and GAPDH) and target genes (p21, PLK1, and FOXM1) were generated from the amplification of the serially diluted RNA samples isolated from the control MCF-7 cell line to optimize the assay. Each of the treated and untreated samples were run in triplicate and non-template control was included. Normalization of reference genes was done using geNorm.\(^\text{11}\)

**Western blot analysis**

Protein was extracted from IC$_{25}$ and IC$_{50}$ DHAQC (2)-treated and untreated control cells using radioimmunoprecipitation assay buffer, and the protein concentration was determined using Bradford protein assay (Thermo Fisher Scientific). The cell homogenates were electrophoresed by sodium dodecyl sulfate polyacrylamide gel electrophoresis (SDS-PAGE) (Bio-Rad Laboratories Inc., Hercules, CA, USA), transferred onto nitrocellulose membrane, blocked with Tris-buffered saline–Tween buffer containing 5% skimmed milk powder for 1 hour, and incubated with primary antibodies (anti-p53, anti-BAX, anti-Bcl-2, anti-cytochrome c, and anti-β-actin) at dilution 1:1000 (Abcam plc, Cambridge, UK) for another hour. Then, membranes were washed and incubated with the anti-rabbit secondary antibodies-conjugated with alkaline phosphatase (1:5000 dilution) (NEB [UK], Hitchin, UK) using a BioSpectrum® reagent (UVP, Upland, CA, USA). The size of the targeted proteins was determined by protein marker (PageRuler Prestained Protein Ladder; Thermo Fisher Scientific), and the density results obtained from this targeted protein were analyzed using VisionWorks®LS analysis software (Version 7.1 RC3.54; UVP) by normalizing against β-actin.

**Table 1** Quantitative real-time polymerase chain reaction primers designed for quantification of FOXM1, WEE1, p21, and PLK1 genes.

<table>
<thead>
<tr>
<th>Accession number</th>
<th>Gene</th>
<th>Sequence</th>
</tr>
</thead>
<tbody>
<tr>
<td>NM_001220778.1</td>
<td>p21</td>
<td>F: 5-TGTCGTCGTCAGACCCCATGC-3</td>
</tr>
<tr>
<td></td>
<td></td>
<td>R: 5-AAAACTGAAATTTCATGCTC-3</td>
</tr>
<tr>
<td>NM_0050303.3</td>
<td>PLK1</td>
<td>F: 5-CCTGCCCGGAACCCAGTTAT-3</td>
</tr>
<tr>
<td></td>
<td></td>
<td>R: 5-CCGCTATATTACGAGTTTTGTC-3</td>
</tr>
<tr>
<td>NM_001243089.1</td>
<td>FOXM1</td>
<td>F: 5-ATACGTTGATGAGGCACTACT-3</td>
</tr>
<tr>
<td></td>
<td></td>
<td>R: 5-TCCATGTCAGTGGACGGTTG-3</td>
</tr>
<tr>
<td>NM_001101.3</td>
<td>ACTB</td>
<td>F: 5-AGACACTAGCAGCTGGTCGAC-3’</td>
</tr>
<tr>
<td></td>
<td></td>
<td>R: 5-ACAAGTCTTGTGCGTACG-3’</td>
</tr>
<tr>
<td>HQ387008.1</td>
<td>18s rRNA</td>
<td>F: 5-GTAACCGTGCGACCCCATTT-3</td>
</tr>
<tr>
<td></td>
<td></td>
<td>R: 5-CCATCACAATCGCGGTAGCCG-3</td>
</tr>
<tr>
<td>NM_002046.4</td>
<td>GAPDH</td>
<td>F: 5-GATTGGTCGCTATTGGGCC-3</td>
</tr>
<tr>
<td></td>
<td></td>
<td>R: 5-TGGAAGATTGTGGTGAGGATT-3</td>
</tr>
</tbody>
</table>

**Note:** ACTB, 18s rRNA, and GAPDH were housekeeping genes used for normalization of the target genes.

**Abbreviations:** F, forward; R, reverse.
Caspase-9 fluorescence assay and viability in the presence of general caspase inhibitor
MCF-7 cells were pretreated with or without cell permeable general caspase inhibitor (Z-VAL-FMK) (Promega Corporation, Fitchburg, WI, USA) at 10 μM before treatment with IC_{25} and IC_{50} of DHAQC (2) for 48 hours. After the treatment period, viability of DHAQC (2)-treated MCF-7 cells with or without Z-VAL-FMK was quantified using MTT assay as described above. Then, protein was extracted from IC_{25} and IC_{50} DHAQC (2)-treated and untreated control cells with or without Z-VAL-FMK using radiolimunoprecipitation assay buffer, and the protein concentration was determined using Bradford protein assay. Levels of activated caspase-9 were detected using CaspGLOW Red Active Caspase-9 Staining Kit (BioVision, Inc., Milpitas, CA, USA) according to the manufacturer’s protocol. The fluorescence intensity in DHAQC (2)-treated and untreated MCF-7 cells was measured at excitation (Ex)/emission (Em) =540/570 nm using a microplate fluorometer (Arzoskan Flash; Thermo Fisher Scientific).

Statistical analysis
Data were presented as mean ± standard deviation from three experiments in three different biological replicates, and the significance (P<0.05) was evaluated by one-way analysis of variance (ANOVA) with Duncan test as post hoc analysis using SPSS software (v 14; SPSS Inc., Chicago, IL, USA).

Results and discussion
Chemical modifications in the side chain are one common approach in drug discovery for improving the therapeutic potential of a certain compound. Our main interest was to improve the yield of damnacanthal using different kinds of oxidizing agents, such as AgO/HNO₃, Vilsmeier reagent, and Lewis acid. None of the methods gave a reasonable yield and most gave oxidative demethylated products. However, the oxidation approach of a previously synthesized compound, 2-hydroxymethyl-1,3-dimethoxy-9,10-anthraquinone (1), with Jones reagents resulted in a 32% yield of DHAQC (2) (Figure 1).

Both damnacanthal and DHAQC (2) contain the anthraquinone planer basic skeleton, which is the major contributor to the cytotoxicity properties. The planar tricyclic structure of anthraquinone is essential for cytotoxic activities and has been widely investigated for cancer therapy. Anthraquinone derivatives such as doxorubicin, daunorubicin, and carminomycin are highly effective in the treatment of tumors. Cytotoxicity of DHAQC (2) on K562, MCF-7, and MCF-10A cell lines was evaluated using the MTT assay, and the IC_{50} value of these cell lines after 48 hours of treatment were obtained (Table 2). The selectivity of DHAQC (2) and damnacanthal on normal and cancerous cell lines was calculated by dividing the IC_{50} on normal MCF-10A cells by that of either estrogen-dependent MCF-7 or estrogen-independent MDA-MB-231 cells. Both DHAQC (2) and damnacanthal showed the highest cytotoxicity against MCF-7 compared to the other cell lines. Further, the IC_{50} values of DHAQC (2) and damnacanthal in MCF-7 and K562 were lower compared to the precursor, compound (1) with IC_{50} values of 12.80 and 14.00 μg/mL, respectively. Previously, we demonstrated the contribution of methoxy, CHO groups, and OH group to the cytotoxicity of anthraquinones against tested cancerous cell lines. DHAQC (2) possesses two OH groups and one carboxylic acid (COOH) group, which gives it a greater hydrophilic character compared to damnacanthal. The presence of hydroxyl and acidic groups in DHAQC (2) increases the solubility and might facilitate absorption or diffusion across the cellular membrane, as revealed by the theoretical membrane-interaction quantitative structure–activity relationship (MI-QSAR) studies of 188 drug-like compounds. The better selectivity of DHAQC (2) toward cancer cell lines is possibly a result of its hydrophilic character, which might fulfill Lipinski’s rule of five and have drug-like properties. Many drugs, for instance, mitoxantrone, possessing 1,4 hydroxyl groups have demonstrated potent antitumor activity and have been used widely in clinic since 1980. Moreover, the hydrophilic property of DHAQC (2)

![Figure 1](https://example.com/figure1.png)

**Figure 1** Synthesis of DHAQC (2) from 2-hydroxymethyl-1,3-dimethoxy-9,10-anthraquinone (1).

**Abbreviation:** DHAQC, 1,3-dihydroxy-9,10-anthraquinone-2-carboxylic acid.

**Table 2** The IC_{50} values of DHAQC (2) and damnacanthal in K562, MCF-7, MDA-MB-231, and MCF-10A cell lines after 48 hours of treatment

<table>
<thead>
<tr>
<th>Cell line</th>
<th>DHAQC (2) (μg/mL)</th>
<th>Damnacanthal (μg/mL)</th>
</tr>
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<tbody>
<tr>
<td>K562</td>
<td>9.10±1.16</td>
<td>6.20±1.32</td>
</tr>
<tr>
<td>MCF-7</td>
<td>7.20±1.00</td>
<td>4.30±0.67</td>
</tr>
<tr>
<td>MDA-MB-231</td>
<td>9.94±1.25</td>
<td>7.13±1.38</td>
</tr>
<tr>
<td>MCF-10A</td>
<td>24.15±1.13</td>
<td>6.51±1.65</td>
</tr>
<tr>
<td>Selective index of MCF-10A/MCF-7</td>
<td>3.35</td>
<td>1.51</td>
</tr>
<tr>
<td>Selective index of MCF-10A/MDA-MB-231</td>
<td>2.43</td>
<td>0.91</td>
</tr>
</tbody>
</table>

**Note:** Selective indexes on breast cancer cells were calculated based on IC_{50} of MCF-10A normal cells versus MCF-7 and MDA-MB-231 cells.

**Abbreviations:** DHAQC, 1,3-dihydroxy-9,10-anthraquinone-2-carboxylic acid; IC_{50}, inhibitory concentration with 50% cell viability.
may also be involved in its chelating with magnesium and zinc ions in DNA synthesis and the repair process. Added to this, the hydrophilicity or introduction of acidic groups increased the hydrophilicity of the rings and facilitated binding with receptor. It is well known that the polycyclic (aromatic) structure of anthraquinone can intercalate with DNA in between base pairs, either covalently or electrostatically. This DNA intercalation with anthraquinone compounds possibly inhibits the DNA replication in cancer cell lines. Previous research found that the position but not the total amount of hydroxyl in anthraquinone played an important role in determining its toxicity against normal and cancerous cell lines in vitro. Thus, modification of the hydroxyl group in DHAQC (2) might reduce its toxicity to normal cells while maintaining its apoptosis-inducing effect on cancer cells, thus leading to greater selectivity of DHAQC (2) compared to that of damnacanthal. As a result, the slight modification in natural compounds, as in DHAQC (2), could be useful for the development of new prototypes of anthraquinone-type cytotoxic agents.

Since DHAQC (2) showed the highest sensitivity and selectivity on hormone-dependent breast cancer MCF-7 cells, further evaluation on cell cycle regulation, apoptosis, gene expression, and apoptosis-related protein levels were evaluated. The cell cycle regulation and apoptotic effects were evaluated using flow cytometry after 48 hours of treatment. Many cytotoxic agents arrest the cell cycle

Figure 2 Cell cycle progression (A–C) and apoptosis (D–F) of DHAQC (2)-treated (IC_{50} values) and untreated MCF-7 cells after 48 hours of incubation. Notes: (A–C) indicate cell cycle phases of untreated control, IC_{25} of DHAQC (2)-treated MCF-7 cells, and IC_{50} of DHAQC (2)-treated MCF-7 cells, while (D–F) indicate apoptosis of untreated control, IC_{25} of DHAQC (2)-treated MCF-7 cells, and IC_{50} of DHAQC (2)-treated MCF-7 cells by annexin V/propidium iodide apoptosis assay, respectively.

Abbreviations: DHAQC, 1,3-dihydroxy-9,10-anthraquinone-2-carboxylic acid; FITC, fluorescein isothiocyanate; IC_{25}, inhibitory concentration with 25% cell viability; IC_{50}, inhibitory concentration with 50% cell viability.

Figure 3 Expression of p21, PLK1, and FOXM1 after 24 hours of IC_{50} DHAQC (2)-treated compared to untreated MCF-7 cells quantified by quantitative real-time polymerase chain reaction assay. Notes: All data are mean ± standard deviation of triplicate. *Indicates significant fold change (>twofold) compared to control, P-value <0.05.

Abbreviations: DHAQC, 1,3-dihydroxy-9,10-anthraquinone-2-carboxylic acid; IC_{50}, inhibitory concentration with 50% cell viability.
Figure 4 (A) Representative plots of Bcl-2, BAX, p53, cytochrome c, and β-actin (ACTB) of untreated, IC_{25} DHAQC (2)-treated, and IC_{50} DHAQC (2)-treated MCF-7 cells after 48 hours. (B) Fold change of IC_{25} DHAQC (2)-treated and IC_{50} DHAQC (2)-treated MCF-7 cells compared to untreated MCF-7 cells. (C) Ratio of BAX/Bcl-2 for untreated, IC_{25} DHAQC (2)-treated, and IC_{50} DHAQC (2)-treated MCF-7 cells after 48 hours. *Indicates significant fold change (P<0.05) compared to control.

**Abbreviations:** C, control; DHAQC, 1,3-dihydroxy-9,10-anthraquinone-2-carboxylic acid; M, protein marker; IC_{25}, inhibitory concentration with 25% cell viability; IC_{50}, inhibitory concentration with 50% cell viability.
of cancerous cells at the S and G2/M phases, followed by the induction of apoptosis. DHAQC (2) arrested MCF-7 at the S and G2/M phases (21.33%±1.74% and 37.66%±4.11%, respectively) as compared to the control (11.76%±1.31% and 4.48%±3.72%, respectively) after 48 hours of incubation. DHAQC (2) also induced 20.48%±2.91% of the MCF-7 cell population to enter the subG0 phase, which indicated the occurrence of DNA fragmentation in the DHAQC (2)-treated cells. This result was further supported by the annexin V/PI apoptosis study, wherein DHAQC (2) was able to induce 16.36%±3.72% of cells to undergo late apoptosis and 29.35%±3.81% to undergo early apoptosis (Figure 2). These cell cycle and apoptosis results were similar to those of our previous study, wherein damnacanthal was found to induce G2/M arrest, DNA fragmentation, and apoptosis toward MCF-7 cells.

Since cell cycle arrest was observed in DHAQC (2)-treated MCF-7 cells, expression of p21, PLK1, and FOXM1 genes involved in G2/M arrest was evaluated using qRT-PCR. PLK1 is a key regulator of G2/M phase progression, while forkhead box M1 (FOXM1) is a direct binding partner of PLK1, FOXM1 plays an important role in regulating the mitotic entry and expression of clusters of G2/M target genes. Downregulation of PLK1 and FOXM1 expression are always associated with G2/M arrest and apoptosis (Kim et al, 2012). On the other hand, upregulation of p21 was found to arrest the cell cycle

![Diagram](image.png)

Figure 5 Cell viability (A) and active caspase-9 fold change (B) of DHAQC-treated and IC50 DHAQC-treated MCF-7 cells with or without general caspase inhibitor Z-VAD-FMK.

Note: Different lowercase letters indicate significant fold change (P<0.05) of standard error of the mean ± standard deviation compared to control.

Abbreviations: DHAQC, 1,3-dihydroxy-9,10-anthraquinone-2-carboxylic acid; IC50, inhibitory concentration with 50% cell viability; IC25, inhibitory concentration with 25% cell viability;
in the G2/M phase as reported in damnacanthal-treated MCF-7 cells. Based on the qRT-PCR results, DHAQC (2) was able to inhibit the expression of PLK1 significantly. Further, expression of p21 and FOXM1 were up- and down-regulated, respectively, although it is not significant (Figure 3). Depletion of PLK1 gene was also reported to subsequently induce apoptosis in cancer cells (Liu and Erikson, 2003).27 Thus, significant PLK1 downregulation by DHAQC (2) contributed to the G2/M arrest and apoptosis in MCF-7 cells as observed in the flow cytometry analysis.

Besides regulation of G2/M arrest-related genes, effects of DHAQC (2) on apoptosis-related proteins, including Bcl-2, BAX, p53, and cytochrome c, were also evaluated by Western blot analysis. After 48 hours of treatment, both IC_{25} and IC_{50} of DHAQC (2) were able to enhance the levels of p53, BAX, and cytochrome c, but had no dose-dependent effect on Bcl-2 (Figure 4). The ability of DHAQC (2), as an anthraquinone, to bind and chelate the DNA may contribute to enhancement of p53 levels.28 Furthermore, this DNA binding ability may be further enhanced by the increased hydrophilic property of DHAQC (2). p53 is a tumor suppressor protein that plays an important role in regulating G2/M arrest and apoptosis via upregulation of p21, transcriptional target of p53, and multidomain proapoptotic protein BAX. BAX is essential for the onset of mitochondrial permeabilization and apoptosis via the intrinsic pathway. The balance between expression levels of antiapoptotic Bcl-2 and proapoptotic BAX is crucial for cell survival or death, wherein increase of the BAX/Bcl-2 ratio will activate the release of cytochrome c from mitochondria and subsequently activate the intrinsic apoptotic pathway.29 Induction of BAX increased the

![Proposed mechanisms of DHAQC (2)-induced G2/M phase arrest and apoptosis in MCF-7 cells.](image-url)

**Abbreviations:** Casp9, caspase-9; Cytoc c, cytochrome c; DHAQC, 1,3-dihydroxy-9,10-anthraquinone-2-carboxylic acid; FITC, fluorescein isothiocyanate.
BAX/Bcl-2 ratio, which subsequently enhanced the release of cytochrome c from the mitochondria (Figure 4) and increased the level of activated caspase-9 (Figure 5), which indicates the activation of the intrinsic apoptotic pathway. Similarly to another anthraquinone derivative, dantran, 

stimulation of activated caspase-9 and reduction of cell viability was suppressed in the presence of general caspase inhibitor Z-VAD-FMK. These results indicate that activation of caspase-9 is crucial for induction of apoptosis in MCF-7 by DHAQC (2). Furthermore, activation of p53 that induced G2/M arrest and the intrinsic apoptotic pathway was similar to the effects induced by damnamcanth. Additionally, regulation of these apoptosis-related proteins was in a dose-dependent manner, wherein upregulation of BAX, p53, cytochrome c, and caspase-9 were found to be higher in MCF-7 cells treated with IC50 of DHAQC (2) than in those treated with IC25 of DHAQC (2).

Conclusion

This study indicates that the production of DHAQC (2) warranted a better yield (32%) than the synthesis of damnamcanth. The synthesized DHAQC (2) also possessed better selectivity against estrogen-dependent breast cancer MCF-7 cells compared to damnamcanth. The better selectivity of DHAQC (2) compared with damnamcanth may have resulted from the increase of hydrophilicity or introduction of acidic groups. Induction of cell death by DHAQC (2) was contributed by G2/M cell cycle arrest and the downregulation of the PLK1 gene induces the intrinsic pathway of apoptosis via upregulation of p53, BAX/Bcl-2 ratio, and release of cytochrome c (Figure 6). Caspase inhibitor Z-VAD-FMK prevented the cell death induced by DHAQC (2) and inhibited the increase of active caspase-9 levels, suggesting that caspase-9 might be involved in DHAQC (2)-induced apoptosis of MCF-7 cells. Besides cell cycle arrest and apoptosis, damnamcanth demonstrated other bioactivities, including inhibition of cell migration and invasion via inhibition of kinases. Thus, further evaluation and comparison of the mechanisms of in vitro and in vivo antitumor and antimetastatic effects via inhibition of protein tyrosine kinase between DHAQC (2) and damnamcanth should be carried out.

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Author contributions

All authors made substantial contributions to conception and design, acquisition of data, or analysis and interpretation of data. All authors took part in either drafting the article or revising it critically for important intellectual content. All authors gave final approval of the version to be published.

Disclosure

The authors report no conflicts of interest in this work.

References


