The Chemopreventive Effect of *Tanacetum Polycephalum* Against LA7-Induced Breast Cancer in Rats and the Apoptotic Effect of a Cytotoxic Sesquiterpene Lactone in MCF7 Cells: A Bioassay-Guided Approach

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Key Words
*Tanacetum polycephalum* • Asteraceae • Sesquiterpene lactone • 8β- hydroxyl- 4β, 15-dihydrozaluzanin C • Breast cancer • Apoptosis • LA7 cells • MCF7 cells

Abstract
**Background:** *Tanacetum polycephalum* L. Schultz-Bip is a member of the Asteraceae family. This study evaluated the chemopreventive effect of a *T. polycephalum* hexane extract (TPHE) using in *in vivo* and *in vitro* models. **Methods and Results:** Five groups of rats: normal control, cancer control, TPHE low dose, TPHE high dose and positive control (tamoxifen) were used for the *in vivo* study. Histopathological examination showed that TPHE significantly suppressed the carcinogenic effect of LA7 tumour cells. The tumour sections from TPHE-treated rats demonstrated significantly reduced expression of Ki67 and PCNA compared to the cancer control group. Using a bioassay-guided approach, the cytotoxic compound of TPHE was identified as a tricyclic sesquiterpene lactone, namely, 8β- hydroxyl- 4β, 15-dihydrozaluzanin C (HDZC). Signs of early and late apoptosis were observed in MCF7 cells treated with HDZC and were attributed to the mitochondrial intrinsic pathway based on the up-regulation of Bax and the down-regulation of Bcl-2. HDZC induced cell cycle arrest in MCF7 cells and increased the expression of p21 and p27 at the mRNA and protein levels. **Conclusion:** This results of this study substantiate the anticancer effect of TPHE and highlight the involvement of HDZC as one of the contributing compounds that act by initiating mitochondrial-mediated apoptosis.

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Introduction

Breast cancer is the most prevalent cancer amongst females and presents an emerging major health problem, both socially and economically. In the United States, it is estimated that approximately 3 million women are living with a history of invasive breast cancer [1]. The metastasizing ability of breast cancer cells to various organs, mostly lymph nodes, lungs, the liver and bones, is accountable for the poor prognosis and high mortality rate in breast cancer patients [2]. Therefore, metastasis is still a high risk factor for medical practitioners involved in breast cancer treatment [3]. Furthermore, the development of resistance in breast cancer epithelial cells to currently used anticancer drugs is a growing challenge in the field of cancer therapy [4]. Although, theoretically, anticancer agents should target only tumour cells, there is a plethora of adverse side effects with the currently available treatments, including bleeding, immunosuppression, hair loss and diarrhoea [5].

Disruption of the native homeostatic mechanism or major perturbations in the cell lifecycle can affect the delicate maintenance of the entire population of a certain tissue. Consequently, the extreme increase of cell growth results in serious complications, such as invasion, metastasis and mechanical pressure [6]. These dramatic changes are mostly associated with a wider spread of the cancer, transformation from a benign to malignant tumour and/or resistant to conventional chemotherapies. A series of events can revolutionize the static state of the organ and initiate a drawback scenario in which unwanted diverse effects or an irreversible mutation may occur [7]. Nevertheless, it has been established that the evasion of apoptosis is one of the six critical changes in cell physiology that can lead to the development of malignant growth [8, 9]. In contrast, apoptosis (programmed cell death) is an important physiological hallmark that is responsible for regulating the generation or degeneration of newly born cells. Continuous insults and prolonged exposure to risk factors can alter the cross-talk between the cell and the surrounding tissue as well as the speed of the release of regulatory proteins and genes, including those related to apoptosis, i.e., Bax, Bcl-2, and caspases, among others [10]. In fact, new strategies for tumour eradication are often built upon the induction of apoptosis and the suppression of the cell cycle cascade at critical phases [10].

*Tanacetum polycephalum* (L.) Schultz-Bip, a member of the Asteraceae family, is an aromatic perennial plant that grows in many regions of the Northern Hemisphere [11, 12]. *T. polycephalum* has been reported to have antiallergic, anticancer, anti-irritant, antiseptic, analgesic and antihypertensive effects [13]. Previous studies have shown that the anti-allergic and anti-inflammatory effects of this aromatic plant are due to the presence of sesquiterpene lactones and volatile oils [14, 15]. In our previous report, we showed that the *T. polycephalum* hexane extract (TPHE) from leaves has selective cytotoxic effects against different cancer cell lines, including A549, CEMss, HepG2, HT29, MCF7, MDA-MB-231 and PC3. Further, we investigated the mitochondrial-mediated apoptotic effect of the extract on MCF7 breast cancer cells and detected a potent effect corresponding to an IC\textsubscript{50} value of 6.42 ± 0.35 μg/mL [16]. The present study was designed to examine the chemopreventive activity of TPHE on the development and growth of LA7-induced breast cancer in rats. In addition, TPHE was subjected to a bioassay-guided approach to identify the cytotoxic compound 8β-hydroxyl-4β, 15-dihydrozaluzanin C (HDZC, Fig. 6) from *T. polycephalum* and to investigate its apoptosis-inducing effects.

Materials and Methods

Plant materials and the preparation of the extract

The leaves of *T. polycephalum* were collected from Shahrekord, Chaharmahal and Bakhtiari, Iran, in May 2013, and a voucher specimen was deposited at the Herbarium, Biological Institute, Shahrekord Azad University. Air-dried leaves were reduced to powder using a grinder. The powder of leaves (4 kg) was used
for hexane extraction at room temperature, and the filtrates were concentrated under reduced pressure using a Buchi R110 Rotavapor (Buchi Labortechnik AG, Flawil, Switzerland) at 40 °C.

Cell culture maintenance and viability
LA7 (rat mammary tumour cells), MCF10A (human breast epithelial cells) and MCF7 (human breast cancer cells) cell lines were purchased from American Type Cell Collection (ATCC, Manassas, VA, USA) and maintained in DMEM, MEBM (Sigma, St. Louis, MO, USA) or RPMI-1640 (Sigma) supplemented with 10% foetal bovine serum (FBS, Pasching, Austria) in a 37 °C incubator with 5% CO₂ saturation. Medium containing 0.1% DMSO (Sigma) was used as a vehicle control for the in vitro assays.

Animals
Pathogen-free female Sprague Dawley rats aged 6-10 weeks were obtained from the Animal Facility of the Faculty of Medicine, University of Malaya, Kuala Lumpur, Malaysia. They were housed at 25 ± 3 °C with a relative humidity of 55-60 °C (a cycle of 12 h-light and 12 h-dark) and were provided with standard food pellets and tap water ad libitum. The animal studies were sanctioned and approved by the Institutional Animal Ethical Committee (FAR/26/07/2013HK).

Experimental design and animal treatment
A total of thirty female rats were randomly divided into five different groups (n=6). The animals were classified as group A-E. Group A consisted of the non-treatment LA7-induced non-treated tumour control animals, group B consisted of animals treated with a low dose (250 mg) of TPHE, group C consisted of animals treated with a high dose (500 mg) of TPHE, group D consisted of animals treated with tamoxifen, and group E consisted of animals that were the normal control group. For treatment, TPHE and tamoxifen were dissolved in Tween-20 and fed orally to the rats once a day for 6 weeks using gastric tubes two weeks before LA7 cell injection. The body weight of each rat was recorded weekly. At the end of the experiment, all of the rats were sacrificed via CO₂ asphyxiation. Mammary tissue samples were obtained, washed twice with ice-cold PBS and kept for histopathological analyses.

Induction of mammary gland tumours
After reaching 90% of confluence, LA7 cells grown in DMEM medium were removed from the medium and the monolayer was washed with PBS. The cells were then detached from the culture flask by adding trypsin (EDTA). The cells were placed into a falcon tube and centrifuged at 1800 rpm for 5 min at 4 °C and then washed with PBS. The cells were then counted using a haemocytometer. The cell suspensions were used within one h of harvesting. LA7 cell suspensions containing 5 × 10⁶ cells in 300 μl of PBS were injected subcutaneously into the left flank mammary fat pad of each rat using a tuberculin syringe and a 21-gauge needle.

Tumour growth
To monitor mammary tumour development, the diameters of the tumours were measured and recorded vertically and horizontally. Tumour volume (V) was calculated by the modified ellipsoidal formula: V = (ab²)/2, where ‘a’ and ‘b’ are the longest and shortest diameter of the tumour, respectively.

Histopathological examination
The breast tumours were inserted in blocks, fixed with 10% buffered formalin and embedded in paraffin. The blocks were sectioned into approximately 5 μm thick sections, which were then stained with Hematoxylin & Eosin (H & E). The stained sections were then evaluated and images were captured using light microscopy (Olympus BX51, Tokyo, Japan) by a histopathologist who was blinded to the treatment assignments.

TUNEL assay
The TUNEL assay was used to measure fragmented DNA in apoptotic cells by combining flourescein-12-dUTP (a) at 3′-OH DNA ends using the terminal recombinant deoxynucleotidyl transferase enzyme (rTdT). The mammary tumour tissues were subjected to the DeadEnd Fluorometric TUNEL system according to the manufacturer’s instructions (Promega Inc., Madison, WI, USA). Then, the sections were evaluated and
images were captured and analysed under a confocal microscope (Zeiss LSM 510, Zeiss, Thornwood, NY, USA) using standard fluorescein filters.

**Immunohistochemistry**

The paraffin sections were fixed with formaldehyde and were then subjected to graded alcohol dehydration and embedded in paraffin. The sections were incubated with primary antibodies against PCNA, Ki67, Bax, Bcl-2, Caspase 3, p21, p27 and p53 (Santa Cruz Biotechnology, Santa Cruz, CA, USA) and were incubated with a biotinylated secondary antibody. Colorimetric detection was performed using a DAB detection kit (Dako, Glostrup, Denmark). Images were captured using a light microscope (Olympus BX51).

**General experimental procedures**

Column Chromatography (CC) was run on silica gel 60 (40-63 μm), while TLC was performed on aluminium and glass plates pre-coated with silica gel 60 F<sub>254</sub> (Merck, Germany). UV spectra were recorded on a Shimadzu UV-160A spectrophotometer using MeOH as a solvent, and 1H NMR and 13C NMR spectra were collected (JEOL JNM-FX500). The separation step was performed using HPLC with a PDA detector and an ODS, C-18 column (Phenomenex). MS was obtained using an Agilent 6530 and FT-IR: Perkin-Elmer RX 1 (Fourier transform infra-red) spectrometer was utilized to analyse the IR at frequencies of 4000-400 cm<sup>-1</sup>.

**Cell viability analysis**

Cell viability analysis was carried out using a MTT (3-(4,5-dimethylthiazol-2-yl)-2,5-diphenyltetrazolium) assay as previously described [17]. Briefly, cells in the exponential growth phase were treated with HDZC at different concentrations (0.39 μg/ml to 50 μg/ml) in 96-well plates and were incubated for 12, 24 and 48 h. Cells treated with tamoxifen (Sigma) were used as the positive controls. After incubation, MTT dye (5 mg/ml, 50 μl, Sigma) was added to each well, and the resulting formazan crystals were dissolved using DMSO (200 μl). Absorbance was measured at 570 nm using a microplate reader (Asys UVM340, Eugendorf, Austria), and the anti-proliferative potential of HDZC was expressed as the IC<sub>50</sub> value.

**Bioassay-guided fractionation and isolation of the compounds**

Hexane crude extract (4.6 g) was chromatographed on a silica gel 60 column (40-63 μm particle size) and eluted sequentially with Hexane/CH<sub>2</sub>Cl<sub>2</sub> mixtures (70:30 → 0:100). Eluates were collected, and those displaying similar R<sub>f</sub> values on TLC were pooled to yield ten fractions (F<sub>1</sub>−F<sub>10</sub>). Fraction F<sub>9</sub> (0.53 g) was further purified by micro column chromatography using CH<sub>2</sub>Cl<sub>2</sub>/MeOH (80:20 → 0:100) and was separated into 6 and 4 sub fractions (F<sub>91</sub>−F<sub>96</sub>) and (F<sub>951</sub>−F<sub>954</sub>) (Fig. 7). The separation of fraction F<sub>953</sub> (0.23 g) by preparative HPLC with a ODS C-18 4.6 x250 mm, 5.0 μm, 70 A column and the mobile system of (50-100% ACN-H<sub>2</sub>O, detection at 210 nm, 7 mL/min) successively yielded a single chromatogram that was identified as 8β-hydroxy-4β,15-dihydrozaluzanin C<sub>1</sub> (1, 5 mg, 0.0025 %). This is the first time that 8β-hydroxy-4β,15-dihydrozaluzanin C<sub>1</sub> (Fig. 6) has been isolated from T. polycephalum. It was confirmed to be tricyclic sesquiterpene lactone by comparing NMR, mass and other physical properties with previously reported data [18].

**Characterization data**

The white amorphous solids were characterized as follows: UV (MeOH), m<sub>max</sub> 3401, 1704 cm<sup>-1</sup>; LCMS m/z 264.14 [M+H]]<sup>+</sup> (calcld for C<sub>15</sub>H<sub>20</sub>O<sub>3</sub>); <sup>1</sup>H NMR (CDCl<sub>3</sub>, 500 MHz), δ 6.24 (d, <i>J</i> = 5.8 Hz, 1H, H-3α), δ 5.53 (d, <i>J</i> = 3.1 Hz, 1H, H-3β), δ 5.36 (t, <i>J</i> = 2.46 Hz, 1H, H-3α), δ 3.99 (m, 1H, H-4), δ 1.53, 2.30 (m, 2H, H-5), δ 4.99 (m, 2H, H-6α and H-6β), δ 2.69 (m, 1H, H-7a), δ 1.97 (m, 2H, H-7), δ 4.28 (m, 1H, H-8), δ 1.56 (d, <i>J</i> = 3.2 Hz, 3H, H-9), δ 3.20 (m, 1H, H-10), δ 2.43 (m, 1H, H-10a), δ 4.30 (m, 1H, H-11); <sup>13</sup>C NMR (CDCl<sub>3</sub>, 125 MHz), δ, 169.90 (C-2), δ 156.09 (C-3), δ 152.90 (C-4), δ 80.57 (C-5), δ 37.7 (C-6), δ 112.78 (C-6 αβ), δ 5.242 (C-7a), δ 2.13 (C-13), δ 126.61 (C-13), δ 82.47 (C-9), δ 44.42 (C-10), δ 47.41 (C-10a), δ 83.08 (C-11), δ 139.11 (C-3), and δ 148.31 (C-6) (Table 2).

**Cell cycle analysis**

Cycle progression in the cells treated with HDZC was examined using flow cytometric analysis [19]. In brief, MCF7 cells (5 × 10<sup>4</sup> cells/mL), at the exponential phase of growth, were treated with HDZC at the
IC₅₀ concentration and incubated for 12, 24 and 48 h. After the incubation, the cells were harvested by trypanosinization and washed twice with PBS (Sigma) prior to fixation with 70% ethanol. The cells were then washed and stained with propidium iodide (PI, 1 mg/mL, 50 µL). RNase A (1 mg/mL, 20 µL) was added to limit the binding of PI to the DNA molecules. The cycle distribution of 10,000 cells per sample was examined using the BD FACSCanto II flow cytometer (BD Biosciences, San Jose, CA, USA), and the results were analysed using the Cell Fit Cell analysis program.

**Annexin V-FITC analysis**
Flow cytometric analysis of early and late apoptosis was investigated using Annexin-V-FITC/PI staining [20]. Briefly, after treating the cells with HDZC at IC₅₀ concentrations for 12, 24 and 48 h, MCF7 cells were harvested and washed twice with PBS. The cells were suspended in Annexin-V binding buffer (BD Biosciences) and were stained with Annexin V-FITC and PI (BD Biosciences) for 30 min in the dark. The fluorescent intensity was analysed via quadrant statistics for early and late apoptotic signs using a BD FACSCanto II flow cytometer.

**Caspase bioluminescent analysis**
The assay was carried out using Caspase-Glo 7 and Caspase-Glo 9 commercial kits (Promega, Madison, WI, USA) as previously described [21]. Briefly, MCF7 cells (1 × 10⁵ cells/mL) were seeded onto a white 96-well microplate and were treated with HDZC at an IC₅₀ concentration followed by incubation for 3, 6, 12, 24 and 48 h. The cells were then supplemented with caspase-Glo reagents according to the manufacturer’s instructions. The bioluminescent activity of the caspases was measured using the Infinite-200 Pro luminescence microplate reader (Tecan, Männedorf, Switzerland).

**Real-time Q-PCR analysis**
The time-dependent mRNA expression of apoptotic related genes in treated breast cancer cells was examined using quantitative PCR analysis [22]. In brief, the RNeasy plus commercial kit (Qiagen, Valencia, CA, USA) was used to extract total RNA from MCF7 cells after treatment with the IC₅₀ concentration of HDZC. cDNA synthesis from extracted RNA was performed using an iScript cDNA synthesis kit (Bio-Rad, Hercules, CA, USA) according to the manufacturer’s protocol. Q-PCR analysis was performed with SsoFast EvaGreen Supermix (Bio-Rad) on a StepOne PLUS real-time PCR machine (Applied Biosystems, Foster City, CA, USA). The mRNA expression of genes was expressed after normalization with the β-actin housekeeping gene. The IDs for Solaris Gene Expression (Thermo Fisher Scientific, Waltham, MA, USA) Assays used in this study are Bax: AX-003308-00-0100; Bcl-2: AX-003307-00-0100; Caspase-7: AX-004407-00-0100; Caspase-9: AX-003309-00-0100; p21: AX-003471-00-0100; p27: AX-003472-00-0100 and β-actin, AX-003451-00-0100.

**Western blot analysis**
A time-dependent western blot analysis was performed as previously described by Hajrezaie and colleagues with some modifications [23]. In brief, the total proteins were extracted from treated cells using Pierce cell lysis buffer (Rockford, IL, USA) according to the manufacturer’s instructions. Extracted proteins (40 µg/mL) were separated by 10% SDS-PAGE and transferred to PVDF membrane (Bio-Rad). Casein (Pierce) was used to block the PVDF membrane prior to washing with TBST. The membrane was incubated overnight with specific primary antibodies, namely, Bax: 1:10,000 (Cat: sc-493); Bcl-2: 1:10,000 (Cat: sc-492); Caspase 7: 1:10,000 (Cat: sc-33773); Caspase 9: 1:10,000 (Cat: sc-7885); p21: 1:10,000 (Cat: sc-397); p27: 1:10,000 (Cat: sc-528) and β-actin, 1:10,000 (Cat: sc-7210) (Santa Cruz Biotechnology, Inc., CA, USA). After incubation, appropriate peroxidase-coupled secondary antibodies were applied to the blot and the detection of bands was carried out using the Fusion FX7 system (Vilber Lourmat, Germany).

**Statistical analysis**
Data from the in vivo study were reported as the means ± SEM of n animals per group. The in vitro results for at least three independent experiments were expressed as the mean value ± SEM, and the differences were considered to be significant at the *p<0.05 level. Statistical analysis was carried out using GraphPad prism (version 4.0 Graphpad software Inc., San Diego, CA, USA). Analyses of variance were performed using the one-way ANOVA procedure followed by Tukey’s post hoc test.
Results and Discussion

Tumour development, body weight, tumour size and volume

Tumours appeared as early as 10 days after LA7 cell injections. The body weights, tumour volumes and tumour percentages (%) of treated and control animals are represented in Table 1. There was a significant body weight decrease in the cancer control group compared with the normal control animals, however, the low dose did not show any significant changes in weight compared to the controls.

Tumours in group A were enlarged up to 2,137 ± 324 mm³; however, a significant decrease appeared in the high TPHE doses (457 ± 132 mm³) and tamoxifen (396 ± 96 mm³) treated groups compared with the tumour control group. The low dose of TPHE did not show a significant decrease in tumour size compared to the tumour control and tamoxifen group (1,781 ± 248 mm³), but the high dose of TPHE was shown to significantly reduce the tumour size comparable to tamoxifen treatment. Based on our finding of the size of the tumour and the body weight of the animal, we can conclude the efficacy of TPHE in reducing tumour size, which is associated with increases in body weight (Table 1).

Table 1. Tumor size and body weight of rats. I. Normal control group (NC), II. Tumor control group (TC), III. Tumor treated with low dose of TPHE (TT + LD), IV. Tumor treated with high dose of TPHE (TT + HD), V. Tumor treated with tamoxifen (TT + TAM). Values are given as mean ± SEM for six rats in each group. One-way ANOVA followed by post hoc test. *p<0.05 compared with the control group.

<table>
<thead>
<tr>
<th>Group</th>
<th>Treatment groups</th>
<th>Body weight (g)</th>
<th>Tumor volume (mm³)</th>
<th>Reduction of tumor percentage (%)</th>
</tr>
</thead>
<tbody>
<tr>
<td>I</td>
<td>NC</td>
<td>300.57 ± 10.63</td>
<td>0</td>
<td>0</td>
</tr>
<tr>
<td>II</td>
<td>TC</td>
<td>285.16 ± 8.57</td>
<td>2137 ± 324</td>
<td>0</td>
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<tr>
<td>III</td>
<td>TT + LD</td>
<td>273.61 ± 10.41</td>
<td>1781 ± 248</td>
<td>16.65</td>
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<tr>
<td>IV</td>
<td>TT + HD</td>
<td>254.68 ± 15.38</td>
<td>*457 ± 132</td>
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</tr>
<tr>
<td>V</td>
<td>TT + TAM</td>
<td>212.93 ± 14.42</td>
<td>*396 ± 96</td>
<td>*81.46</td>
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Fig. 1. Histological study of normal and treated breast cancer tissues. A) Normal breast tissue. B) Breast tumor induced by LA7. C) Breast tissue from low dose of TPHE treatment group. D) Breast tissue from high dose of TPHE treatment group. E) Breast tissue from tamoxifen treated group. H and E staining of the tumor tissues shows morphological differences between normal breast tissues and cancerous breast tissue. The treatment of the tumor with TPHE shows reorganisation if the tumor part to the normal tissues. Magnification 40x.
Histopathology

Pleomorphism of the cells confirmed the induction of breast cancer tumours by LA7 cells (Fig. 1). Based on these results, we conclude that the development of the tumour was based on the invasive adenocarcinoma subtype with no tubular structure. In normal tissue, the duct was healthy and there was no sign of variation in size and shape; however, the untreated tumour showed a significant variation in the size and shape of the duct, followed by the disruption of the ductal structure (Fig. 1). Groups treated with high doses of TPHE had reduced mitotic events and smoothening of the tissue that led to the reorganization of the cells compared to the untreated group. However, normal rats treated with TPHE showed no changes in mammary gland morphology. Based on the morphology of the cells, the improved mammary gland structure was significantly more visible after treatment with high doses of TPHE and tamoxifen compared to the untreated cancer rats and there were no significant differences in the low dose treatment group.

Apoptosis

The TUNEL assay was used to determine whether inhibition of the tumour cells occurred through apoptosis. The tumour sections of the rats injected with breast cancer inducing LA7 showed significantly higher numbers of apoptotic cells after treatment with TPHE than the normal control and mammary tumour control rats. Indeed, the control groups showed very little or no obvious signs of apoptosis in the mammary glands (Fig. 2). Apoptotic cells were apparent in mammary tumour sections of all of the high dose TPHE- and tamoxifen-treated rats (Fig. 2).

Immunohistochemistry

The expression of cell nuclear antigen (PCNA) and Ki67 protein are closely associated to the proliferation of the tumour cells. The high levels of PCNA and Ki67 indicate an aggressive tumour prognosis. Immunohistochemical analysis of the cancer control group showed higher expressions of PCNA and Ki67 compared to the TPHE treatment groups. These findings demonstrate the promising chemopreventive Table 2. NMR spectroscopic data (500 MHz, CDCl3) for compound 1 (δ in ppm, J in Hz)

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<th>Position</th>
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<td>3α</td>
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Fig. 2. Induced apoptosis of breast cancer cells as determined by the TUNEL assay. A) Control tumor tissue. B) Breast tissue from low dose of TPHE treatment group C) Breast tissue from high dose of TPHE treatment group. D) Breast tissue from tamoxifen treated group. Breast cancer tissue sections from the control and treatment group were subjected to the TUNEL assay. Tissue from the control group did not shows any significant changes in the color of the tissues While tissue from the high dose TPHE and tamoxifen treated groups showed a very high number of apoptotic cells. The low dose TPHE treated group did not show significant apoptotic activity. Magnification 40x.
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The potential of TPHE, which was proven by the reduced protein expressions of tumor markers (Fig. 3).

Because the potency of TPHE in inducing apoptosis in breast cancer cells was previously reported to occur through the mitochondrial pathway [16], we investigated the expression of Bax, Bcl-2, caspase 3, p53, p21 and p27 proteins, as these proteins are implicated in the induction of apoptosis through intrinsic apoptosis pathways. The microscopic observations of the slides in Fig. 4 demonstrate the significant differences between the TPHE high dose and tamoxifen-treated groups in comparison to the tumor control group. The results represent the up-regulation of Bax, p53 and caspase 3 proteins and the down-regulation of the expression of Bcl-2 proteins. The low dose TPHE treated group was not significantly different compared to the positive control group (Fig 4). Immunohistochemical observations...
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Programmed cell death can be triggered through p53 expression associated with a transcriptional modulator, which can be closely mediated by the suppression or activation of other gene transcriptions [24]. The function of p53 is also closely related to the ratio of Bax/Bcl-2 proteins [25]. Tumours with the loss of function of p53 are expected to have low levels of Bax and high levels of Bcl-2. In addition, the G1-S phase of cell cycle arrest can be closely mediated by the suppression or activation of the p53 gene, which has the ability to induce the cyclin-dependent kinase genes p21/p27. Our results showed the up-regulation of the Bax protein and the down regulation of the Bcl-2 protein after treatment with a high dose of TPHE, while the tumour control group had low expressions of Bax and high expressions of Bcl-2 protein (Fig. 4), suggesting that apoptosis occurred through the mitochondrial pathway. The expression level of p53 was elevated in the high dose TPHE treatment group compared to the untreated control. The up-regulation of p21/p27 in the high dose TPHE treatment group compared to the normal control group could be due to an arrest in the cell cycle progression (Fig. 5).

**Sesquiterpene lactones**

Sesquiterpene lactones, an extensive and diverse class of biologically active compounds, have been widely isolated from various plants belonging to the Asteraceae family [26]. Sesquiterpene lactones are a group of naturally occurring plant terpenes representing crucial constituents of essential oils [27]. In recent years, numerous studies have

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**Table 3.** HDZC IC_{50} concentration against MCF7 and MCF10A cells after treatment for 12, 24 and 48 h. The data represent the means ± SEM of three independent experiments

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<thead>
<tr>
<th>Cell line</th>
<th>HDZC 12 h</th>
<th>Tamoxifen 12 h</th>
<th>HDZC 24 h</th>
<th>Tamoxifen 24 h</th>
<th>HDZC 48 h</th>
<th>Tamoxifen 48 h</th>
</tr>
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<tbody>
<tr>
<td>MCF7</td>
<td>6.42 ± 1.73</td>
<td>3.30 ± 0.58</td>
<td>3.80 ± 0.32</td>
<td>2.60 ± 0.63</td>
<td>2.47 ± 0.21</td>
<td>1.5 ± 0.15</td>
</tr>
<tr>
<td>MCF10A</td>
<td>90.85 ± 2.64</td>
<td>70.09 ± 1.79</td>
<td>85.21 ± 2.82</td>
<td>54.59 ± 1.08</td>
<td>69.46 ± 0.17</td>
<td>32.46 ± 2.54</td>
</tr>
</tbody>
</table>

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**Fig. 5.** Immunohistochemistry staining of p21 and p27 A) Tumor control B) Low dose treatment C) High dose treatment D) Tamoxifen treatment. Cell cycle arrest was confirmed by high expression of p21 and p27 proteins in treatment groups when compare to control group.

**Fig. 6.** Chemical structure of the compound HDZC.
demonstrated that different sesquiterpene lactones possess a notable anticancer potential [27, 28]. For instance, costunolide isolated from Saussurea lappa roots [29], parthenolide isolated from Tanacetum parthenium [30-32] and helenalin isolated from Arnica species [33, 34] were reported to be anti-cancer sesquiterpene lactones. In this study, 8β-hydroxy-4β,15-dihydrozaluzanin C (Fig. 6) was isolated from T. polycephalum to assess its biological activity on human breast cancer cells.

**Fig. 7.** Flow chart of bioassay-guided isolation of HDZC from the hexane extract of T. polycephalum leaves. Each fraction was subjected to MTT assay for the evaluation of its suppressive effect on MCF7 cells. IC50 values represent the means ± SEM of three independent experiments.

**Fig. 8.** Cell cycle analysis. Cell cycle analysis of MCF7 cells treated with (A) vehicle 0.1% DMSO (control) for 48 h and HDZC (IC50 concentration) for (B) 12, (C) 24 and (D) 48 h. (E) The representative bar chart illustrated the significant G1 cell cycle arrest after 12, 24 and 48 h. The data represent the means ± SEM of three independent experiments. *p<0.05 compared with the control group.

**HDZC selectively suppressed the growth of MCF7 cells**

The antiproliferative effect of HDZC against a breast cancer cell line (MCF7) and a non-cancer cell line (MCF10A) was screened using the MTT assay. As shown in Table 3, HDZC
suppressed the growth of MCF7 cells at IC$_{50}$ values of 6.42 ± 1.73, 3.80 ± 0.32 and 2.47 ± 0.21 µg/mL after 12, 24 and 48 h, respectively. These results were comparable to the tamoxifen cytotoxic effect (3.30 ± 0.58, 2.60 ± 0.63, 1.5 ± 0.15 µg/mL after 12, 24 and 48 h, respectively). HDZC elicited a powerful effect against MCF7 cells compared to the hexane extract (IC$_{50}$ value of 6.42 ± 0.35 µg/mL at 48 h of treatment) [16]. Meanwhile, proliferation of normal cells (MCF10A) was affected by HDZC solely at higher concentrations.

**HDZC induced G1 cell cycle arrest**

Flow cytometry analysis of MCF7 treated with HDZC indicated significant increases of the cell population in the G1 phase after 12, 24 and 48 h (Fig. 8). Arrest of the cell cycle progression is an effective approach for inhibiting the proliferation of cancer cells [35-37]. PI dye binds to DNA molecules at all cell cycle stages when applied to apoptotic cells, and thereafter, the emitted fluorescent intensity representing the DNA content of the cells is measured using flow cytometry analysis. Fig 8 demonstrates the cell cycle phase distribution in MCF7 cells specific to each phase after treatment with HDZC at the IC$_{50}$ concentration. Expression of the G1 phase was most evident in the MCF7 cells treated with HDZC (12 h: 61.13%; 24 h: 62.64%; 48 h: 64.25%) relative to the control cells. The arrest of MCF7 cells at the G1 phase was associated with a consequent reduction in the percentage of cells at the S and G2/M phases.

A previous study showed that costunolide sesquiterpene lactone induced G1 cell cycle arrest in hormone dependent (LNCaP) and independent (PC-3 and DU-145) prostate cancer cells [38]. Similarly, parthenolide sesquiterpene, in combination with NS398, resulted in cell cycle arrest at the G0/G1 phase in Hep3B, HepG2 and PLC human hepatocellular carcinoma cells [39]. Jang and co-workers reported a dose-dependent elevation in the sub G0 populations of human renal carcinoma Caki cells treated with helenalin sesquiterpene [34]. Consistently, our findings showed that HDZC, like other sesquiterpene lactones, arrested the cell cycle of the breast cancer cell line at the G1 phase.

**HDZC induced apoptosis in MCF7 cells**

The aforementioned results of the selective cytotoxic effect of HDZC towards MCF7 cells encouraged us to examine the early events of apoptosis. It has been well established...
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that during early apoptosis, plasma membrane asymmetry is lost because of the phosphatidylserine externalization [40]. Cancer is generally defined as a disease of abnormal cell growth; therefore, disruption or deformity of the cell plasma membrane means losing the cell architecture, leading to the discontinuity of their life cycle. Annexin V-FITC/PI followed by flow cytometry analysis is a methodology that is applied to examine the integrity of the cell external membrane [41]. We observed MCF7 cell membrane leakage with double staining with different intensities in accordance to the phase of apoptosis. As shown in Fig. 9, in treated cells, early apoptotic cells (Annexin V+/PI−) and late apoptotic cells (Annexin V+/PI+) began to emerge as early as 12 h at proportions of 24.2% and 3.7%, respectively, while control cells supplemented with a 0.1% DMSO vehicle elicited low or negative staining with both Annexin V-FITC and PI for visualized and well-survived viable cells. The number of the apoptotic cells increased significantly from 12 to 48 h, and the different stages of the programmed cell death, from early apoptosis till necrosis, were clearly observed in a time-dependent manner. However, our findings revealed a dominance of the late apoptotic phase (after 24 h), suggesting that the HDZC mode of action may rely on a correlation between the cell cycle arrest and the loss of cell integrity. It is noteworthy that the elevation of necrotic cells (Annexin V+/PI+) reached 22.7% after 48 h of treatment.

**Fig. 10.** Effect of HDZC on the activation of caspase 9 and caspase 7. The caspase 7 and 9 on MCF7 cells was determined using a bioluminescent analysis. MCF7 cells were treated with the vehicle 0.1% DMSO (control) for 48 h and HDZC (IC_{50} concentration) for 3, 6, 12, 24 and 48 h. Caspase 9 and caspase 7 were significantly activated after 12, 24 and 48 h treatment. The data represent the means ± SEM of three independent experiments. *p<0.05 compared with the control group.

**Fig. 11.** Q-PCR analysis of apoptosis- and cell cycle-associated molecules, viz. Bcl-2, Bax, caspase 9, caspase 7, p21 and p27. MCF7 cells were exposed to HDZC at IC_{50} concentration for 12, 24 and 48 h. The mRNA expression of genes was represented after normalization with β-actin housekeeping gene. The data represent the means ± SEM of three independent experiments. *p<0.05 compared with the control group.
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Fig. 12. Western blot analysis of apoptosis- and cell cycle-associated molecules, viz. Bcl-2, Bax, caspase 9, caspase 7, p21 and p27. MCF7 cells were exposed to HDZC at IC50 concentration for 12, 24 and 48 h. β-actin was used as a positive control. The data represent the means ± SEM of three independent experiments.

HDZC activated caspase 7 and caspase 9

Caspases, a family of cysteine proteases including at least 14 members, form a complex and energy-dependent cascade that executes apoptotic cells after a fatal insult [42]. The hierarchical organization of caspases generally consists of several initiator proteases (caspase 2, 8, 9, and 10), executioner or effector proteases (caspase 3, 6, and 7) and inflammatory proteases (caspase 1, 4, and 5) [43]. Caspase 9, in particular, is a key mediator of mitochondrial-mediated apoptosis, and the up-regulation of caspase 9 follows the upstream activation of caspase 7 prior to apoptosome formation and the fluctuation of the mitochondrial membrane potential [44]. To determine whether the induced apoptosis was caspase-mediated, we measured the activity of caspase 9 and caspase 7 in MCF7 cells using a bioluminescent analysis. Our studies indicated a significant time-dependent elevation of caspase 9 and caspase 7 (for example, the mRNA expression of caspase 9 and caspase 7 was 8.30 and 5.85-fold, respectively, after 48 h) after treatment with HDZC in comparison to the control group (Fig. 10). However, the elevation of caspase 9 was higher than caspase 7, as observed in Fig. 11. Western blot analysis also confirmed the time-dependent enhancement of the expression of caspase 9 and caspase 7 (Fig. 12). Together, these results highlight the involvement of the intrinsic pathway of mitochondrial-derived apoptosis mediated by caspase 9 and caspase 7 dysregulation in MCF7 cells.

The effect of HDZC on Bax/Bcl-2 expression

Translocation of the pro-apoptotic Bax protein from the cytosol to the outer mitochondrial membrane and its oligomerization leads to changes in the mitochondrial membrane potential, both of which are associated with cytochrome c leakage [45]. On the other hand, the suppression of cytochrome c release is controlled by the anti-apoptotic Bcl-2 protein [46, 47]. Both the pro-apoptotic and anti-apoptotic proteins cause perturbations among mitochondrial-activated functions, such as those of caspase 9 [48]. Because HDZC elicited the ability to activate caspase 9, it was of interest to determine the possible role of the two proteins, Bax and Bcl-2, in neutralizing this effect. Hence, we examined the mRNA and protein expression of these proteins using Q-PCR and western blot analysis. As shown in Fig. 11, Bax mRNA expression was elevated after 12 h and reached a 9-fold significant increase in the experimental groups compared to the control. Bcl-2 mRNA expression was found to decrease in a time-dependent fashion (Fig. 11). The results of the western blot analysis indicated that the up-regulation of Bax and the down regulation of Bcl-2 are additionally observed on the protein level (Fig. 12). These findings demonstrate that HDZC induced apoptosis by altering the regulation of apoptotic-related pro-proteins at both the mRNA and protein levels.
The effect of HDZC on p21/p27 expression

As accelerators of the cell cycle, cyclin-dependent kinases (CDKs) and their cyclin partners have a pivotal role in cell cycle progression and, accordingly, have been proposed to be key players in the suppression of accelerators [49, 50]. Thus, the activation of cyclin dependent kinase inhibitors (CKIs) effectively adjusts the abnormal proliferation of cancer cells. The CKIs proteins p21 (CIP1) and p27 (KIP1) are members of the CIP/KIP family and are known to form heterotrimeric complexes that promote cell cycle arrest via the G1/S CDKs cascade [51]. Because HDZC possessed the ability to induce G1 cell cycle arrest in MCF7 cells, the next step was to elucidate the possible mechanism behind this outcome and to determine whether it includes the targeted accelerators/inhibitors. Hence, we examined the mRNA and protein expression level of p21 and p27 using Q-PCR and western blot analysis, respectively. The Q-PCR assay showed a significant up-regulation of the mRNA expression of p21 and p27, which reached 2.83 (p21) and 3.57 (p27)-fold higher levels compared to the controls after 48 h of HDZC treatment. As depicted in Fig. 12, western blot confirmed the perturbations in the mRNA expression of p21 and p27 at the protein level in a time-dependent manner. In sum, the up-regulation of p21 and p27 mediated by the introduction of HDZC treatment is part of the intrinsic pathway governed by mitochondrial genes and leads to the programmed death of cancer cells.

Conclusions

Our study demonstrates the in vivo chemopreventive activity of TPHE. Our results reveal the capability of TPHE to induce apoptosis in LA7-induced tumours in rats accompanied by caspase activation and the up-regulation of Bax and down-regulation of Bcl-2. Moreover, the exposure of tumour cells to TPHE caused cell cycle arrest, as was revealed by p21/p27 protein expression accompanied by p53 activation. The isolation of 8β-hydroxy-4β,15-dihydrozaluzanin C sesquiterpene lactone from T. polycephalum is reported, to the best of our knowledge, for the first time. The bioassay guided approach revealed the potent cytotoxic effect of the sesquiterpene lactone. The accumulating body of data has shown the capacity of this class of compounds to act as anti-cancer agents. HDZC displayed a powerful and selective effect on cell survival, life cycles and integrity. In-depth molecular investigations were performed to illustrate the possible mechanisms of action. The elevated activity of caspase 9 associated with the up-regulation of Bax and the down-regulation of Bcl-2 proved that the induction of apoptosis in MCF7 is through the mitochondrial pathway. Additionally, HDZC caused cell cycle arrest at the G1 phase via alterations of the mRNA and protein expression of p21/p27. Collectively, our work suggests the promise of finding a renewable source of novel anti-cancer therapeutic agents from medicinal plants.

Disclosure Statement

No conflict of interest.

Acknowledgements

Financial support from the University of Malaya, the high impact research grant (UM-MOHE UM.C/625/1/HIR/MOHE/SC/09) and the IPPP research grant (PG053/2012B) are greatly appreciated.

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