Ferulago angulata activates intrinsic pathway of apoptosis in MCF-7 cells associated with G₁ cell cycle arrest via involvement of p21/p27

Abstract: Ferulago angulata is a medicinal plant that is traditionally known for its anti-inflammatory and antiulcer properties. The present study was aimed to evaluate its anticancer activity and the possible mechanism of action using MCF-7 as an in vitro model. F. angulata leaf extracts were prepared using solvents in the order of increasing polarity. As determined by MTT assay, F. angulata leaves hexane extract (FALHE) revealed the strongest cytotoxicity against MCF-7 cells with the half maximal inhibitory concentration (IC₅₀) value of 5.3±0.82 μg/mL. The acute toxicity study of FALHE provided evidence of the safety of the plant extract. Microscopic and flow cytometric analysis using annexin-V probe showed an induction of apoptosis in MCF-7 by FALHE. Treatment of MCF-7 cells with FALHE encouraged the intrinsic pathway of apoptosis, with cell death transducing signals that reduced the mitochondrial membrane potential with cytochrome c release from mitochondria to cytosol. The released cytochrome c triggered the activation of caspase-9. Meanwhile, the overexpression of caspase-8 suggested the involvement of an extrinsic pathway in the induced apoptosis at the late stage of treatment. Moreover, flow cytometric analysis showed that FALHE treatment significantly arrested MCF-7 cells in the G₁ phase, which was associated with upregulation of p21 and p27 assessed by quantitative polymerase chain reaction. Immunofluorescence and the quantitative polymerase chain reaction analysis of MCF-7 cells after treatment with FALHE revealed an upregulation of Bax and a downregulation of Bcl-2 proteins. These findings proposed that FALHE suppressed the proliferation of MCF-7 cells via cell cycle arrest and the induction of apoptosis through intrinsic pathway.

Keywords: Ferulago angulata, apoptosis, cancer, MCF-7, cell cycle, p21/p27

Introduction
Cancer is a debilitating disease that has caused a markedly high death rate in all generations. Despite noteworthy progress in reducing cancer incidence and the developments in chemotherapeutic strategies, it is still a major public health problem in many parts of the world. The available chemotherapeutic agents only provide minimal survival benefits in advanced-stage cancer due to various factors, namely toxicity, side effects, and drug resistance. The main modes of therapy used for the treatment of this disease are surgery and chemotherapy, with varying degrees of failure due to high resistance to chemotherapy. Treatment responses are especially gloomy for hormone-independent cancers. Hence, research into new therapeutic agents for these cancers is clearly warranted.

Natural products have been used for the treatment of different ailments and diseases for thousands of years. Thus, because of the similarity between body composition and plant ingredients and the long history of folk medicine from traditional plants,
there is a rising interest in plant-derived medicines. The World Health Organization estimates that roughly 80% of the world’s inhabitants rely on folk medicine for their primary health care. Natural products’ contribution to cancer therapy is very evident considering the fact that they were involved in the development of approximately 75% of all novel anticancer agents from 1981–2012. Plant products induce cell death by a variety of mechanisms. The most prevalent mechanisms are programmed cell death (PCD)-type I (apoptosis), PCD-type II autophagic cell death, and necrosis.

Apoptosis is a mode of self-cannibalism, which involves individual cells and does not cause inflammation to the neighboring cells. Thus, apoptosis, which is characterized by a series of biochemical and morphological changes, including deoxyribonucleic acid (DNA) fragmentation, nuclear condensation, externalization of phosphatidylserine (PS), membrane blebbing, and loss of mitochondrial membrane potential (MMP), has become a mechanism of interest in drug research. Mitochondria are found to be a crucial factor in the regulation of apoptosis. A loss in MMP leads to the translocation of proapoptotic Bax to mitochondria, which results in the release of cytochrome c and the activation of caspase cascades. In addition, excessive production of reactive oxygen species leading to oxidative stress and the depletion of the glutathione level has been reported to be a trigger to apoptotic signaling.

Ferulago angulata, locally known as chavir, is a member of the Apiaceae family, which is mostly indigenous to Iraq, Turkey, and western Iran. This perennial shrub, with a height of 60–150 cm, is widely used in folk medicine to treat digestive pains, hemorrhoids, snake bites, and ulcers. It is very evident considering the fact that they were involved in the development of approximately 75% of all novel anticancer agents from 1981–2012. Plant products induce cell death by a variety of mechanisms. The most prevalent mechanisms are programmed cell death (PCD)-type I (apoptosis), PCD-type II autophagic cell death, and necrosis.

Materials and methods

Plant materials

F. angulata plants were collected from Shahrekord, Chaharmahal and Bakhtiari province, Iran, in March 2012, and a voucher specimen of this plant has been deposited at the Herbarium, Biological Institute, Shahrekord Azad University, Iran. The leaves of F. angulata were cut into thin slices and dried at 25°C. The dried leaves (1.5 kg) were then ground with a mill grinder into coarse powder and were first extracted with n-hexane (3,000 mL, three times) in conical flasks. The leaves’ residues were then dried and sequentially reextracted with dichloromethane and methanol using the same method. The resulting filtrates were dried under reduced pressure by a rotary evaporator (Büchi Labortechnik GmbH, Essen, Germany) at 35°C to produce n-hexane (14 g), dichloromethane (10 g), and methanol (9 g) extracts, respectively. The extracts were dissolved in dimethyl sulfoxide (DMSO), (Sigma, St Louis, MO, USA) and stored in −20°C until further experiments.

Cell culture and cell viability assay

A549 (human lung cancer cells), CCD841 (normal human colon epithelial cells), HepG2 (human hepatoma cells), HT29 (human colon cancer cells), MCF-7, and MDA-MB-231 (human breast cancer cells), PC3 (human prostate cancer cells), and WRL-68 (human hepatic cells) cell lines were purchased from American Type Culture Collection (ATCC) (Manassas, VA, USA). The cells were maintained in Roswell Park Memorial Institute 1640 or Dulbecco’s Modified Eagle’s Medium (Sigma), supplemented with 10% fetal bovine serum (PAA Laboratories GmbH, Pasching, Austria) and 1% penicillin/streptomycin (PAA; 100 µg/mL) in a 37°C incubator with 5% CO₂ saturation. Negative control for all the assays was represented by the untreated medium containing vehicle DMSO (0.1%).

Cell viability was determined using MTT assay. In brief, cells (5×10⁴ cells/mL) were treated with three extracts at different concentrations in 96-well plates and incubated for 48 hours at 37°C. MTT was added to the wells and incubated for 3 hours. Cell media was replaced by DMSO and pipetted to dissolve the formed formazan crystals. Then, optical density of the colored solution was quantified at 570 nm wavelength using a microplate reader (UVM340; Asys-Hitech GmbH, Eugendorf, Austria). The potency of cell growth inhibition for each of the three extracts was expressed as an IC₅₀ value. MCF-7 cells treated with F. angulata leaves hexane extract (FALHE) revealed the lowest IC₅₀ when compared to cells treated with the other extracts; therefore, we only used FALHE for further studies. The percentage of cell viability = (absorbance of treated cells/absorbance of untreated cells) ×100%.
Animal experiments and acute toxicity assay

This experiment was carried out after approval by the University of Malaya Institutional Ethics Committee (Ethic #: FAR/26/07/2013/HK [R]). In addition, 6–8 week old rats (150–180 g) were obtained from the Experimental Animal House facility, Faculty of Medicine, University of Malaya. All animals received care, according to the current guidelines for the care of laboratory animals prepared by the National Academy of Sciences and published by the National Institutes of Health Sciences. Also, 18 female rats were divided into three groups and placed in cages that were labeled as: “low dose group” (FALHE, 2 g/kg); “high dose group” (FALHE, 5 g/kg); and “vehicle control group” (Tween-20 10% weight/volume; 5 mL/kg). Before dosing, the rats were fasted overnight but allowed access to water. After fasting, each group was administered with its respective compound, further deprived of food for 3–4 hours, and monitored for 14 days for any sign of toxicity and mortality. Histological, hematological, and serum biochemical parameters were assessed after sacrificing the animals on the 15th day.

Chemical analysis assay

To determine the chemical constituents of FALHE, we carried out the gas chromatography (GC)–mass spectrometry (MS)–time of flight analysis (TOF) analysis, as previously described.27 The analysis of the FALHE was performed using an Agilent and LECO GC-MS, with the following features: RESTEK, Rxi-5MS capillary column (30 minutes; 0.25 µm film thickness) and a mass spectrometer Pegasus HT High Throughput TOFMS. The carrier gas was helium at a flow rate of 1 mL/minute. Column temperature was initially 40°C for 5 minutes, then gradually increased to 160°C at 4°C/minute, and finally increased to 280°C at 5°C/minute and held for 10 minutes. For GC–MS detection, an electron ionization system was used with ionization energy of 70 eV. The fraction was diluted 1:100 (volume/volume) with ethyl acetate, and 1.0 µL of the diluted sample was injected automatically in splitless mode. The injector temperature was set at 250°C. The detected compounds were identified from their mass spectra by comparison of the retention times of peaks with interpretation of MS fragmentation patterns from data library.

Annexin-V-fluorescein isothiocyanate (FITC) assay

Annexin-V, as a Ca2+-dependent phospholipid-binding protein, detects the plasma membrane alterations, such as the PS externalization during the early stages of apoptosis.28 The effect of FALHE on the induction of early apoptosis was analyzed in MCF-7 cells using the annexin-V probe. The MCF-7 cells (5×10^4 cells/mL) at the exponential phase of growth were seeded into a chamber slide plate. After 48 hours of FALHE treatment at IC_{50} doses, they were washed with phosphate buffered saline (PBS) and stained with fluorescein-labeled annexin-V containing a binding buffer. The MCF-7 cells were observed under fluorescent microscopy, Olympus BX51 (Olympus Corporation, Tokyo, Japan).

To confirm PS translocation suggesting an induction of apoptosis by FALHE observed in microscopic investigation, we carried out the flow cytometric analysis using the BD Pharmingen Annexin V-FITC Apoptosis Detection Kit (APOAlert® Annexin V; Clontech, Mountain View, CA, USA). In brief, MCF-7 cells (1×10^5 cells/mL) at the exponential phase of growth were exposed to FALHE at IC_{50} concentration for 24, 47, and 72 hours. After centrifuging the cells and removing the media, the MCF-7 cells were washed twice with PBS and resuspended in the annexin-V binding buffer. The cells were then stained with annexin-V-FITC and propidium iodide (PI), according to the manufacturer’s instructions. The flow cytometric analysis was performed using BD FACSCanto II flow cytometer (BD Biosciences, San Jose, CA, USA) followed by quadrant statistics analysis for the detection of early and late apoptotic cells and necrotic cells.

Cell cycle analysis

Cell cycle analysis was carried out using a flow cytometer as previously described.29 In brief, the MCF-7 cells (5×10^4 cells/mL) at the exponential phase of growth were treated with FALHE for 24, 48, and 72 hours. After incubation, treated MCF-7 cells were centrifuged and washed with PBS twice prior fixing with 95% cold ethanol (700 µL) at 4°C overnight. After fixing, the cells were washed again and resuspended in PBS (500 µL). The fixed MCF-7 cells were treated with PI (80 µL; 1 mg/mL) for 45 minutes, and the binding ability of PI was restricted to DNA by adding the ribonuclease A (20 µL; 10 mg/mL) for degradation of RNAs. The DNA content of the cells was analyzed by using a BD FACSCanto II flow cytometer (BD Biosciences, San Jose, CA, USA) and a ModFit LT software (Verity Software House, Inc., Topsham, ME, USA).

Caspase analysis

The effect of FALHE on the activation of caspase-8 and caspase-9 was investigated using a commercial kit.
Multiple cytotoxicity assay

Multiple cytotoxicity was determined using the Cellomics Multiparameter Cytotoxicity 3 Kit (Cellomics, Pittsburgh, PA, USA), as previously described in detail. This kit provides simultaneous measurements of four independent cell health parameters in a single cell, i.e., changes in nuclear intensity, cell membrane permeability, cytochrome c release, and MMP. In this experiment, MCF-7 cells were stained with the nuclear dye (Hoechst 33342), cell permeability dye (FITC), mitochondrial membrane potential dye (Cy5), and cytochrome c dye (Cy3). The plates were analyzed using the ArrayScan HCS system (Cellomics).

Apoptosis proteome profiler array

To determine the probable mechanism of action for induced apoptosis by FALHE, we examined the proteins involved in different pathways of apoptosis using the Proteome Profiler Array (RayBio Human Apoptosis Antibody Array Kit, RayBiotech, Inc., Norcross, GA, USA), according to the manufacturer’s protocol. Briefly, after treatment of MCF-7 cells with FALHE at IC₅₀ dose for 48 hours, the proteins were extracted from each sample, and 300 µg of the extracted proteins were incubated with the human apoptosis array for 24 hours. The membrane was then used to quantify the result via scanning on a Biospectrum AC ChemiHR 40 (UVP, Upland, CA, USA). An image analysis software (Progenesis SameSpots software v.2.0.2733.19819; Nonlinear Dynamics, Newcastle, UK) was used to analyze the array image file.

Gene expression analysis of Bax/Bcl-2 and p21/p27

The gene expression of four proteins, namely Bax, Bcl-2, p21, and p27, which demonstrated the highest changes in proteome profiler array when compared to the control, was investigated by quantitative polymerase chain reaction analysis. After treatment of MCF-7 cells with FALHE at an IC₅₀ dose for 24, 48, and 72 hours, total RNAs were isolated using a Quick-RNA MiniPrep kit (Zymo Research, Freiburg, Germany), followed by the synthesisization of complementary DNAs using a High-Capacity RNA-to-cDNA kit (Applied Biosystems, Foster City, CA, USA), according to the manufacturer’s instructions. The gene expression was measured, based on the relative expression of β-actin as a positive reference. Applied Biosystems StepOnePlus system (Applied Biosystems), Solaris qPCR Gene Expression Master Mix and Solaris Gene Expression (Thermo Fisher Scientific, Waltham, MA, USA) Assays were used for the gene expression analysis of: Bcl-2, AX-003307-00-0100; Bax, AX-003308-00-0100; p21, AX-003471-00-0100; p27, AX-003472-00-0100; GAPDH, AX-004253-00-0100; and β-actin, AX-003451-00-0100.

Immunofluorescence analysis of Bax/Bcl-2

To determine the changes in the expression of Bax and Bcl-2 at the protein level, immunofluorescence analysis was carried out using the Cell Reporter™ (Molecular Devices, Wokingham, UK). The MCF-7 cells (5×10⁴ cells/well) at the exponential phase of growth were seeded in 96-well plate and treated with FALHE at IC₅₀ dose for 24, 48, and 72 hours. After incubation, the cells were fixed using 4% paraformaldehyde at 25°C for 15 minutes. Then, each well was supplied with blocking buffer (0.03% Triton X-100/ PBS and normal serum) for 1 hour. After blocking, the cells were treated with diluted primary antibody solution (1× PBS; BSA/0.3% Triton X-100) and incubated overnight at 4°C. Bax and Bcl-2 fluorochrome-conjugated secondary antibodies (Santa Cruz Biototechnology, Santa Cruz, CA, USA) were diluted in PBS and added to each well and incubated for 1 hour at room temperature. Cells were treated with 4′,6-diamidino-2-phenylindole (DAPI) and examined using the Cell Reporter™ cytofluorimetric system. During the process, cells were washed twice with PBS after every single step.

Statistical analysis

Each assay was carried out three times independently. Statistical analysis was performed using the statistical package GraphPad Prism Version 5 (GraphPad Software Inc., San Diego, USA) and the SPSS-16.0 package (SPSS Inc., Chicago, IL, USA, USA). One-way analysis of variance (Dunnett’s multiple comparison test) was used to distinguish the difference among groups. The results are presented as the means ± standard error of the mean of three
Results and discussion
Cytotoxic effects of FALHE on different cancer cells
The cytotoxic effects of three *F. angulata* extracts on different cancer cells were determined using the MTT assay. Dichloromethane and methanol extracts did not show significant cytotoxic effect toward cancer cells compared to the hexane extract. Meanwhile, the FALHE treatment of cancer cells for 48 hours decreased cell proliferation with a wide range of suppressive effects (IC$_{50}$ ranged from 5.3±0.82 to 70.25±7.42 µg/mL), due to the various range of resistance shown by cancer cells. As shown in Figure 1A, FALHE elicited the strongest cytotoxicity effect on the MCF-7 cells with the IC$_{50}$ value of 5.3±0.82 µg/mL at 48 hours.

In addition, the suppressive effect of FALHE on MCF-7 cells was proved to be concentration- and time-dependent (Figure 1B). In contrast, even the highest concentration of FALHE (50 µg/mL) did not significantly affect the proliferation of normal cells (WRL-68 and CCD-841). The cell viability implied FALHE’s suppressive effect is selective for cancer cells. Following these results, MCF-7 cells and FALHE were chosen for subsequent experiments. Findings from this study were comparable with a previous study, which showed significant cytotoxic effects of *F. angulata* against the Jurkat cell line (IC$_{50}$ <8 µg/mL).\textsuperscript{25} However, an investigation on its suppressive effect on Raji (B-cell lymphoma), U937 (human leukemic monocyte lymphoma),

![Graph A](image1.png)

![Graph B](image2.png)

**Figure 1** Suppressive effect of FALHE against cancer and normal cells.

**Notes:** (A) IC$_{50}$ concentration of FALHE on eight different cell lines after 48 hours of treatment. (B) FALHE suppressed MCF-7 cell proliferation in a concentration- and time-dependent manner. *P*<0.05.

**Abbreviations:** IC$_{50}$, half maximal inhibitory concentration; FALHE, *Ferulago angulata* leaves hexane extract.
and KG-1A (human acute myelocytic leukemia) cell lines showed relatively high IC\textsubscript{50} values of 96.81, 158.68, and 128.47 µg/mL, respectively.\textsuperscript{26}

**Safety of FALHE**

The extensive traditional use of plant remedies by natives for a long period of time without any report of toxicity or adverse side effects can generally provide some evidence for the safety of the respective plant.\textsuperscript{32} However, previous studies showed that some chemical constituents or secondary metabolites present in traditional medicines have toxic or even carcinogenic effects on human body.\textsuperscript{32–34} Therefore, investigation on the safety of plants prior to performing any experiment on their biological activity is highly recommended to avoid any undesirable adverse side effects.

The acute toxicity study of FALHE demonstrated the safety of this extract, since during the 14 days period all the rats remained alive and did not show any sign of toxicity even at a high dose of 5 g/kg. In addition, histological, hematological, and serum biochemical analysis did not show any sign of hepatic or renal toxicity in the treatment groups (Figure 2; Table S1).

**Chemical analysis of FALHE**

The chemical composition of FALHE was characterized using GC-MS-TOF analysis (Figure 3). The chromatogram showed two monoterprenoid phenols as major compounds, ie, thymol (3, C\textsubscript{10}H\textsubscript{14}O) and carvacrol (4, C\textsubscript{10}H\textsubscript{14}O) (Table 1). Previous studies showed that these two phenolic compounds have promising anticancer effects against cancer cell lines.\textsuperscript{35} Thymol was found to induce apoptosis in HL-60 cells through both caspase-dependent and caspase-independent pathways.\textsuperscript{35} The marked antiproliferative effect of carvacrol and its mechanism of action toward different cancer cells,
Figure 3 Chemical constituents of FALHE determined by a gas chromatogram analysis. 
**Note:** 3 (Thymol) and 4 (Carvacrol) are the major compounds based on their intensity. 
**Abbreviations:** FALHE, *Ferulago angulata* leaves hexane extract; AIC, analytical ion chromatogram.

Figure 4 Microscopic evaluation of the PS externalization. 
**Notes:** (A) Untreated cells (control) did not show any sign of PS translocation as an early marker for apoptosis. After 48 hours of treatment with IC_{50} concentration of FALHE, annexin-V-FITC binding to PS residues shown in the light green stain demonstrated the externalization of PS. The treated cells were shown with (B) 20× and (C) 40× magnifications. The white arrows show the PS externalization in treated cells. 
**Abbreviations:** PS, phosphatidylserine; IC_{50}, half maximal inhibitory concentration; FALHE, *Ferulago angulata* leaves hexane extract; FITC, fluorescein isothiocyanate.
including MDA-MB-231, HepG-2, and A549 cells, has previously been investigated. The suppressive effects of these two phenolic compounds on cancer cells are closely correlated with the IC$_{50}$ value of FALHE toward different cancer cells. However, a study with a bioassay-guided approach on FALHE is further required to certainly determine the major bioactive compounds responsible for the observed cytotoxic effects, which is ongoing in our laboratory.

**Externalization of PS induced by FALHE**

Apoptotic properties of cancer cells are the critical key to most of the approaches applied in radiation and chemotherapy.

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**Figure 5** Flow cytometric analysis of apoptosis in MCF7 cells. **Notes:** (A) Shows the untreated MCF-7 cells (control) after 72 hours. After treatment with IC$_{50}$ concentration of FALHE for (B) 24, (C) 48 and (D) 72 hours, flow cytometry analysis was carried out on MCF7 treated and untreated cells. Meanwhile, (E) representative bar chart showed a significant elevation in the number of apoptotic and necrotic cells after 72 hours. *P*-value 0.05.

**Abbreviation:** FALHE, Ferulago angulata leaves hexane extract.
Thus, it has considerable therapeutic implications. Apoptosis is characterized by various biochemical changes in cells, mainly nuclear condensation, DNA fragmentation, PS externalization, MMP changes, membrane blebbing, and regulations in caspases’ enzymes. These changes are considered as the biomarkers of apoptosis.

PCD (apoptosis) has two distinct phases, namely, early and late apoptosis. A simple and noninvasive assay of annexin-V-FITC is designed for early detection of apoptosis. PS, which is normally located on the inner surface of the plasma membrane, translocates to the outer surface of the plasma membrane after an induction of apoptosis, apparently through an active mechanism. Annexin-V-FITC is a probe with a high affinity for PS, which has provided an easy method for the detection of apoptosis, since PS exposure is a widespread event during apoptosis, which occurs prior to membrane leakage and DNA-associated changes. As shown in Figure 4, the treatment of MCF-7 cells with the IC_{50} concentration of FALHE-induced translocation of PS to the outer membrane after 48 hours, while there was no sign of translocation in the control cells.

To confirm the microscopic detection of PS externalization, we carried out the flow cytometric analysis using annexin-V-FITC and PI double staining. Treatment with FALHE for 24 hours significantly elevated the number of early (annexin-V positive; PI negative) and late (annexin positive; PI positive)

![Figure 6](image_url)  
**Figure 6** Effect of FALHE on cell cycle distribution of MCF-7 cells.  
**Notes:** (A) Shows the untreated MCF-7 cells (control) after 72 hours. After treatment with {\textit{IC}_{50}} concentration of FALHE for (B) 24, (C) 48 and (D) 72 hours, flow cytometry analysis was carried out on MCF7 treated and untreated cells. (E) The quantitative analysis shows the cell cycle arrest at G_{0}/G_{1} phase. (*P<0.05).

**Abbreviations:** FALHE, Ferulago angulata leaves hexane extract; {\textit{IC}_{50}}, half maximal inhibitory concentration; G, gap; S, synthesis; M, mitosis.
PI positive) apoptotic cells to 17.7% and 12.45%, respectively (Figure 5B). These numbers reached to 20.5% and 16.15% after 48 hours of treatment, respectively (Figure 5C). Meanwhile, the number of necrotic cells also significantly increased after 48 hours and 72 hours (Figure 5D), which was expected since due to the long incubation time with FALHE and its cytoxic effects, some cells showed necrotic characterizations. The result of flow cytometric analysis confirmed the induction of apoptosis by FALHE in a time-dependent manner (Figure 5E).

**Cell cycle arrest at G1 phase induced by FALHE**

Previous studies have reported that the regulation of cell death and cell proliferation shares prevalent proteins and molecules in multicellular organisms. Indeed, cell cycle regulation and apoptosis are regulated by several mutual genes. A direct correlation between apoptosis and cell cycle shows that apoptosis and mitosis have very close morphological characteristics. During either of the processes, after the cells detach, they become rounded, condense their chromatin, and display membrane blebbing. In spite of some shared features, there are critical differences between mitosis and apoptosis. For example, cell membrane proteins are cross-linked during apoptosis, resulting in the rigidity of the membrane, and apoptotic cells are normally phagocytized by macrophages or adjacent cells. In contrast, during mitosis, the segregation of DNA and the division of the cell by cytokinesis leading to two healthy and viable daughter cells, are observed. To analyze the respective connection, we investigated the cell cycle progression in MCF-7 cells upon treatment with FALHE in a time-dependent pattern for 24, 48, and 72 hours. This experiment was carried out to determine the effect of FALHE on the DNA content of MCF-7 cells by cell cycle phase distribution (gap [G0/G1], synthesis [S], and G2/M) (Figure 6). The results demonstrated that the FALHE significantly arrested the cell cycle progression at the G1 phase in a time-dependent manner. The number of MCF-7 cells at this phase increased from 58.93% (untreated control) to 67.31% (72 hours). Meanwhile, the number of cells in the S and G2/M phases reduced after 72 hours of treatment compared to the control. Furthermore, the significant and time-dependent elevation in the cell population at sub-G1 phase presenting the hypodiploid DNA in MCF-7 cells elicited the number of cells undergoing apoptosis.

**Caspase-8 and caspase-9 activation induced by FALHE**

The caspase cascade signaling system is a critical factor in the process of apoptosis, which is precisely controlled by a variety of proapoptotic and antiapoptotic molecules. After treatment of MCF-7 cells with IC50 concentration of FALHE for 3, 6, 12, 24, 48, and 72 hours, caspase-8 and caspase-9 enzyme activity was evaluated. The activity of caspase-8 significantly increased at 72 hours, while caspase-9 activity significantly rose at 12, 24, 48, and 72 hours of treatment (Figure 7). The apoptosis-inducing potential of FALHE through the mitochondrial pathway was confirmed by the activation of caspase-9 after 12 hours. Furthermore, the elevated activity of caspase-8 after treatment with FALHE has thrown light on the possible contribution of extrinsic pathway to the induction of apoptosis by FALHE at the late stage of treatment. Involvement of more than a single apoptotic pathway in the induction of PCD by anticancer compounds is well-established, especially by plant-derived natural agents, such as curcumin. In this study, untreated cells

**Figure 7 Luminescence assay.**

**Notes:** The luminescence assay revealed an activation of caspase-9 after 12 hours and caspase-8 after 72 hours of treatment with FALHE. *P* < 0.05.

**Abbreviations:** FALHE, Ferulago angulata leaves hexane extract; RLU, Relative luminescence unit.
did not show any increase in the activity of caspase-8 and caspase-9; however, activity was significantly and time dependently elevated after treatment with FALHE. The role of caspase-9 has previously been determined to be a step prior to the activation cascade of the mitochondrial pathway that results in apoptosis. Caspase-9 activation is heavily regulated by apoptosome, which converts procaspase-9 to caspase-9 by enzymatic cleavage.

Multiparameter cytotoxicity analysis

To determine the possible mechanism of action for the induced apoptosis by FALHE, we investigated critical biochemical and morphological changes in MCF-7 cells, in specific nuclear condensation, and in mitochondrial-initiated events. Hoechst 33342 staining demonstrated that numerous cells treated with FALHE displayed nuclear condensation shown with blue fluorescent intensity (Figure 8).

Figure 8 Images of MCF-7 cells.
Notes: These images of MCF-7 cells (>20) were treated with FALHE at IC_{50} concentration, stained with different specific dyes, detecting total nuclear intensity, MMP, cytochrome c release, and cell membrane permeability. The images from each row were captured from the specific field of treated sample.
Abbreviations: FALHE, Ferulago angulata leaves hexane extract; IC_{50}, half maximal inhibitory concentration; MMP, mitochondrial membrane potential.

Figure 9 Quantitative analysis of multiparameter cytotoxicity assay revealed changes in total nuclear intensity, cell membrane permeability, MMP, and cytochrome c release.
Notes: Following treatment with FALHE for 48 hours, significant elevation in total nuclear intensity, cell membrane permeability, cytochrome c release associated with reduction in MMP were detected. (*P<0.05).
Abbreviations: MMP, mitochondrial membrane potential; FALHE, Ferulago angulata leaves hexane extract.
Nuclear intensity (which directly corresponds to apoptotic chromatin changes), cell membrane permeability, MMP, and cytochrome c release were quantified (Figure 9). Nuclear intensity significantly increased after 48 hours of treatment with FALHE. A concurrent elevation in the cell membrane permeability shown with green fluorescent intensity was also exhibited for the same period (Figure 8). MMP was significantly decreased in the MCF-7 cells treated with FALHE, as shown by a conspicuous attenuation in the red fluorescent intensity (Figure 8).

Furthermore, FALHE triggered a significant translocation of cytochrome c from mitochondria into cytosol. At 48 hours, FALHE triggered approximately threefold cytochrome c release represented by cyan fluorescent intensity in Figure 8.

Apoptosis proteome profiler analysis
To further examine the detailed pathways responsible for induced apoptosis by FALHE toward MCF-7 cells, screening of critical proteins that are involved in various pathways of apoptosis was carried out. As shown in Figure 10, critical proteins involved in the intrinsic pathway of apoptosis, including Bad, Bax, cytochrome c, second mitochondria-derived activator of caspases (SMAC)/DIABLO, and serine protease HtrA2/Omi were significantly overexpressed after 48 hours of treatment, while the expression of antiapoptotic proteins of Bcl-2, survivin, X-linked inhibitor of apoptosis protein (XIAP) were downregulated. Additionally, p21 and p27, which are cell proliferation suppressor proteins, were upregulated. Different insulin-like growth factor binding proteins (IGFBPs) were also upregulated while treatments. Specific ligands of the tumor necrosis factor (TNF)-α involved in the extrinsic pathway and death receptors, such as soluble TNF receptor (R) 2, TNF-related apoptosis-inducing ligand (TRAIL)R-3, and TRAILR-4 were slightly induced by treatment with FALHE.

The induced apoptosis by FALHE was associated with significant changes in the expression level of proteins from Bcl-2 family. We found out that the expression of the antiapoptotic protein of Bcl-2 was significantly reduced, while proapoptotic proteins of the Bcl-2 family, including Bax and Bad, were significantly expressed after treatment with FALHE. Changes in the expression of the Bcl-2 family proteins led to the overexpression of cytochrome c, which resulted in the translocation of cytochrome c from the mitochondria to the cytosol as proved by the multiparameter cytotoxicity assay.52 Besides cytochrome c, there are other proteins, such as SMAC/DIABLO and the serine protease HtrA2/Omi, that are released through the mitochondrial permeability transition pores provided by Bax protein in the inner mitochondrial membrane.53

Cytochrome c release has a critical role in the activation of the intrinsic pathway of apoptosis as an important factor in the formation of the apoptosome, while SMAC/DIABLO and HtrA2/Omi suppress the activity of inhibitors of apoptosis proteins (IAPs), such as survivin and XIAP, whose expression was significantly reduced in our experiment.44 IGFBPs have been proved to significantly suppress the proliferation of different cell types. Previous studies have shown that IGFBPs are involved in the induction of apoptosis in MCF-7 cells via a death receptor-mediated pathway.55 The activation of apoptosis by a death receptor-mediated pathway was also suggested by the expression of TNF-α as the activator ligand for the death receptors, including TNFR1 and TNFR2, which are involved in the extrinsic pathway.56 In addition, the activation of caspase-8 after
72 hours of treatment with FALHE compared to the activation of caspase-9 after 12 hours of treatment, as shown in the luminescence analysis, suggests an activation of the intrinsic pathway followed by an activation of the extrinsic pathway at the late stage of treatment. Additionally, the cell cycle has been shown to be involved in the suppression of cyclin-dependent kinases, which are responsible for the G1-to-S transition.57

FALHE upregulated the gene expression of Bax, p21, and p27

Since the protein array analysis showed the most significant changes in the expression of the Bcl-2, Bax, p21, and p27 proteins, we investigated the expression of these proteins at the gene level, using the quantitative polymerase chain reaction analysis. The result showed a significant elevation in the gene expression of Bax, p21, and p27, which supported our findings.
from the protein array analysis. However, Bcl-2 gene expression did not significantly change during 72 hours of treatment (Figure 11). These results suggest that FAHLE treatment induces cell cycle arrest at the G1 phase via overexpression of p21 and p27 at the gene level. Recent studies on the p21 and p27 proteins have determined the critical role of these cell-cycle inhibitors on cell division and, subsequently, on cancer therapy approaches, since various cancer cells contain mutations in members of the proteins that regulate cyclin-dependent kinase activity.58,59 Therefore, apoptosis-inducing agents with the ability to upregulate p21 and p27 proteins at the gene and protein level can be identified as promising anticancer agents.

**FALHE upregulated the protein expression of Bax and downregulated the protein expression of Bcl-2**

Due to the important role of Bcl-2 and Bax proteins in the mitochondrial-dependent apoptosis, expression of these proteins was investigated using immunofluorescence analysis. DAPI dye was used to stain the nucleus of the MCF-7 cells and in determining the number of cells. As shown in Figure 12, the number of MCF-7 cells decreased in a time-dependent manner in the treatment group as compared to the control group. The fluorescent intensity of the Bax-conjugated antibody was markedly increased after increasing the time of treatment with FAHLE (Figure 12A). In contrast, the fluorescent intensity of Bcl-2 was conspicuously decreased in the same time period (Figure 12B). Quantitative analysis of the result showed a significant upregulation of Bax associated with significant downregulation of Bcl-2 after 24, 48, and 72 hours of treatment (Figure 12C). The perturbation in the protein expression of Bcl-2 compared to its unchanged gene expression after 72 hours of treatment suggested that in spite of FAHLE treatment, the Bcl-2 gene was still expressed and its activity was only significantly suppressed at the protein level. However, Bax expression was upregulated both at the gene and protein level.

**Conclusion**

In conclusion, the plant *F. angulata* has been found to have the potential of inducing mitochondrial-mediated apoptosis in MCF-7 cells through the activation of caspase enzymes.
Furthermore, the active role of mitochondria in the induction of apoptosis was confirmed by the release of cytochrome c and reduction in MMP. The cell cycle arrest in G1 phase was proved to be one of the possible mechanisms for suppressing cell proliferation, which suggests to be induced by the upregulation of p21 and p27 proteins. The suggested molecular pathway for the induced apoptosis is summarized in Figure 13. Our result demonstrated that F. angulata is a promising plant for cancer therapy. However, further in vitro and in vivo studies on the active compounds of this plant are still required.

Acknowledgments

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Disclosure

The authors report no conflicts of interest in this work.

References


Supplementary material

Table S1 Serum biochemical analysis of rats

<table>
<thead>
<tr>
<th>Animal groups</th>
<th>Kidney function test</th>
<th>Liver function test</th>
<th>Hematological test</th>
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<tr>
<td></td>
<td>Sodium (mM/L)</td>
<td>Potassium (mM/L)</td>
<td>Chloride (mM/L)</td>
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<tr>
<td>Vehicle</td>
<td>138.56±0.76</td>
<td>3.7±0.01</td>
<td>100.2±0.33</td>
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<tr>
<td>FALHE (2 g/kg)</td>
<td>142.21±0.51</td>
<td>4.1±0.3</td>
<td>101.2±0.4</td>
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<tr>
<td>FALHE (5 g/kg)</td>
<td>140.21±0.77</td>
<td>4.0±0.02</td>
<td>105.2±0.33</td>
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</table>

<table>
<thead>
<tr>
<th>Animal groups</th>
<th>Total protein (g/L)</th>
<th>Albumin (g/L)</th>
<th>Globulin (g/L)</th>
<th>TB (µmol/L)</th>
<th>AP (U/L)</th>
<th>ALT (U/L)</th>
<th>AST (U/L)</th>
<th>GGT (U/L)</th>
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<tr>
<td>Vehicle</td>
<td>70.6±1.6</td>
<td>39.9±0.9</td>
<td>29.7±1.8</td>
<td>1.1±0.05</td>
<td>161.5±5.7</td>
<td>30.9±1.4</td>
<td>24.2±2.8</td>
<td>0.45±0.07</td>
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<tr>
<td>FALHE (2 g/kg)</td>
<td>67.9±1.8</td>
<td>38.5±0.3</td>
<td>30.6±1.2</td>
<td>1.4±0.06</td>
<td>163.8±4.1</td>
<td>35.5±1.7</td>
<td>25.48±2.9</td>
<td>0.57±0.09</td>
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<tr>
<td>FALHE (5 g/kg)</td>
<td>69.7±1.6</td>
<td>38.7±0.9</td>
<td>31.5±1.5</td>
<td>1.4±0.04</td>
<td>163.3±5.2</td>
<td>36.7±1.8</td>
<td>27.7±4.2</td>
<td>0.6±0.03</td>
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<table>
<thead>
<tr>
<th>Animal Groups</th>
<th>HGB (g/dL)</th>
<th>HCT (%)</th>
<th>RBC (10⁶ cells/µL)</th>
<th>MCV (fl)</th>
<th>MCH (pg)</th>
<th>MCHC (g/dL)</th>
<th>RDW (%)</th>
<th>WBC (10³/µL)</th>
<th>Platelet (10³/µL)</th>
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<tbody>
<tr>
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<td>16.03±0.7</td>
<td>39.00</td>
<td>9.3±0.1</td>
<td>60.2±0.5</td>
<td>17.5±0.3</td>
<td>33.34±0.7</td>
<td>14.1±0.4</td>
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<tr>
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<td>59.8±0.7</td>
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<td>32.93±0.9</td>
<td>12.7±0.4</td>
<td>5.9±0.2</td>
<td>969.2±23.8</td>
</tr>
</tbody>
</table>

Notes: Serum biochemical analysis of rats did not show any sign of Falhe toxicity on (A) renal function test, (B) liver function test, and (C) hematological parameters of the rodents, when compared with the control group. Values expressed as mean±standard deviation.

Abbreviations: ALT, alanine aminotransferase; AST, aspartate aminotransferase; AP, alkaline phosphatase; GGT, G-glutamyl transferase; HCT, hematocrit; HGB, hemoglobin; MCH, mean corpuscular hemoglobin; MCHC, mean corpuscular hemoglobin concentration; MCV, mean corpuscular volume; RBC, red cell count; RDW, red cell distribution width; TB, total bilirubin; WBC, white cell. Falhe, Ferulago angulata leaves hexane extract.