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Gynura divaricata on $NF-\kappa B$ pathway
in human HeLa cells

Effects of ethanolic leaf extract of *Gynura divaricata* on NF- κ B pathway in human HeLa cells

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Abstract

This study aims to evaluate the anti-cancer activity of the ethanolic leaf extract of *Gynura divaricata* in human cervical cancer HeLa cells. Cells were treated with various concentrations of extract, and cell viability was assessed to determine the half-maximal inhibitory concentration (IC₅₀). The extract significantly reduced HeLa cell viability in a concentration-dependent manner, demonstrating marked cytotoxic activity. Treatment with 2x IC₅₀ extract markedly suppressed cellular metabolic activity and significantly downregulated the expression of NF- κ B-related genes compared with control and DMSO-treated cells. The extract also reduced the transcription of inflammatory and apoptosis-associated factors, including TNF- α , Bcl-2, Bax, COX-2, IKK α , I κ B α , IL-8, IL-6, and IL-1 β . These findings indicate that *G. divaricata* inhibits HeLa cell growth by suppressing NF- κ B-mediated signaling and inflammatory responses, suggesting its potential as a natural therapeutic agent against cervical cancer.

Introduction

The use of natural antioxidants as pharmaceutical products and functional foods has gained considerable attention, and the antioxidant ability and biological properties of various natural extracts are correlated with the plant sources from which they are derived (Lin et al., 2019). One of these plants, the Taiwanese native species *Gynura divaricata* Formosana, is an edible flowering perennial plant in the family Asteraceae. It is mostly planted in southern Taiwan. It is a valuable and economically significant herbal plant with notable economic potential. Young leaves of *G. divaricata* are commonly consumed in Taiwan as tea or salad garnish, as a natural health food, and historically used in tradi-

tional Chinese medicine to treat hepatitis and liver cancer (Kantawong et al., 2021). Furthermore, the components of *G. divaricata* have been studied for their various bioactivities, and are widely treated as a folk medicinal tonic in Asia (Chao et al., 2014; Nadechanok et al., 2015; Xu et al., 2017; Ma et al., 2018). For example, *G. divaricata* extracts have been used to treat diabetes, hypertension, hyperlipidemia, hyperglycemia, and many other ailments for their anti-cancer effects (Chou et al., 2012; Chen et al., 2014; He et al., 2014; Jin et al., 2018; Yen et al., 2018; Dong et al., 2019; Hong et al., 2020; Ye et al., 2022; Xu et al., 2024; Rifai et al., 2024; Shen et al., 2025).

Transcription factor nuclear factor (NF)- κ B plays a key



role in inflammatory responses and cancer development by regulating genes involved in angiogenesis and tumorigenesis during molecular signaling in the NF- κ B pathway. Elevated NF- κ B activity contributes to increased levels of pro-inflammatory cytokines; therefore, the NF- κ B signaling pathway is considered a potential therapeutic target for cancer therapy (Feng et al., 2023). The expression levels of NF- κ B, cyclooxygenase (COX)-2, and multigenes have shown significant associations with poor survival and increased tumor aggressiveness in human cervical cancer samples (Pasha et al., 2024). Black soybean seed coat extracts inhibit tumor necrosis factor (TNF)- α -induced COX-2 levels through an NF- κ B-dependent pathway, and have anti-inflammatory effects on an immortalized epidermal keratinocyte cell line (Kim et al., 2012). It has anti-inflammatory and anti-atherogenic effects on TNF- α -induced cell adhesion in human aortic endothelial cells (HAEC) by regulating NF- κ B (Chao et al., 2013a). In addition, TNF- α -induced HAEC adhesion is significantly suppressed by mulberry leaf extracts, resulting in reduced expression of NF- κ B (Chao et al., 2013b). Moreover, extract of sweet potato leaves shows anti-inflammatory effects via reducing mitogen-activated protein kinase (MAPK) and NF κ B expressions, inhibiting TNF- α -induced monocyte-endothelial cell adhesion, and attenuating interleukin-8 (IL-8) and vascular cell adhesion molecule-1 (VCAM-1) expressions (Lee et al., 2015). The consumption of purple sweet potato leaves modulates various immune functions, including the secretion of cytokines IL-2 and IL-4 of natural killer cells and induces an increase in proliferation responsiveness in peripheral blood mononuclear cells (Chen et al., 2005). Moreover, purple leaf extracts also depress neuroinflammatory responses in lipopolysaccharide-activated BV-2 microglia cells by inhibiting the production of inflammatory mediators such as inducible COX-2 and TNF- α (Kang et al., 2014). *G. divaricata* water extract inhibits NF- κ B activation and reduces insulin pathway deficiencies, such as IRS1, AKT, and GLUT1 (Li et al., 2018). In addition, aqueous extracts reduce the levels of IL-6 and TNF- α and increase the levels of VEGFA mRNA and protein expression in rats (Sun et al., 2023). It has hypoglycemic effects *in vivo* by regulating the genes at PI3K/AKT and fatty acid metabolism signaling pathway, and the Bcl-2/Bax expression ratio in the liver of rats is significantly up-regulated after *G. divaricata* intervention (Xu et al., 2020).

However, the cytotoxic effects of *G. divaricata* leaf extract on human cervical cancer HeLa cells and the underlying molecular mechanisms remain largely unknown. NF- κ B signaling plays an important role in cancer progression and oxidative stress regulation. Therefore, this study investigated the effects of ethanolic *G. divaricata* leaf extract on HeLa cell viability, antioxidant activity, and NF- κ B-related gene expression. Antioxidant capacity was evaluated using 2,2-diphenyl-

1-picrylhydrazyl (DPPH), ferric reducing antioxidant power (FRAP), malondialdehyde (MDA), hydrogen peroxide (H₂O₂), and antioxidant enzyme assays. The findings provide insights into the antioxidant and anticancer potential of *G. divaricata* and its possible application as a plant-derived complementary therapeutic agent.

Materials and Methods

Plant material and culture practice

Young *G. divaricata* plantlets were originally collected from a local farmer's field in the Peito mountain area, Taipei, Taiwan (25°07'12" N, 121°30'36" E). The subspecies was authenticated and identified according to the List of Plants of Formosa at the National Taiwan University herbarium (<https://tai2.ntu.edu.tw/species/>). The surfaces of stem explants (15 cm long) with 3-paired leaves were propagated in 5-inch plastic pots, one plant per pot, filled with commercial potting soil with a substrate mixture of peat moss, perlite, and vermiculite (3:1:1 v/v/v, Known-You Co., Taiwan). They were then placed in a growth chamber for two months. The average temperature in the growth chamber was 25°C, with day lengths of 16 hours in 1,600 lux/m² and 80% relative humidity. Plants were watered twice daily, and a compound fertilizer solution (N-P₂O₅-K₂O, 20-20-20) was applied once a week. Uniformly sized plants were randomly chosen for further experiments.

Extraction of leaves

Fresh leaves of *G. divaricata* were extracted by maceration using 95% ethanol as the solvent at a ratio of 1:10 (w/v) for 2 hours (Kalsoom et al., 2024) with a shaker at 100 rpm under in a 50 L glass bottle at room temperature, followed by centrifuging (Kubota 3740, Japan) at 12,000 rpm for 10 min at 4°C (Hou et al., 2005). Supernatants were filtered and freeze-dried using a vacuum evaporator (FDM-2000, Eyela, Japan) to remove the solvent until a fat with viscous wax was obtained. The ethanol-free extract was then stored at -20°C until used.

Cell culture treatment and cell viability test

Human cervical epithelioid carcinoma HeLa cells (#60005) were purchased from the Bioresource Collection and Research Center of the Institute of Food Industry, Hsinchu City, Taiwan. The cells were cultured and maintained in Dulbecco's modified Eagle's medium (DMEM) (D5648, Sigma-Aldrich, USA) supplemented with 10% fetal bovine serum (FBS) and 1.5% penicillin-streptomycin antibiotic solutions in 10 cm Petri dishes at 37°C with a 5% CO₂ atmosphere as previously described (Ma et al., 2018) within an incubator (SCA150-Astec, Japan). After the cultured cells reached 80% confluency, they were treated with trypsin, and the

cleaved cells were counted under an inverted microscope (AMEX1000, Life Technologies, USA) using a cell counting plate. Plates were divided according to the number of cells and seeded into new culture plates at a density of 10^5 cells/ mL for sub-culturing.

The 3-(4,5-dimethylthiazol-2-yl)-2,5-diphenyltetrazolium bromide (MTT) assay, by the reduction of MTT in active mitochondrion to purple formazan, was used to determine the effects of potential agents or compounds on the proliferation of cells to assess cell viability (Kumar et al., 2018). Briefly, 5,000 cells in a well were plated in a 96-well plate and kept at 37°C with 5% CO₂ for a day. After the medium was removed, HeLa cells were exposed to different concentrations of *G. divaricata* extract for another 24 hours. HeLa cells that cultured in normal complete DMEM (not treated with extract) served as the control. When incubation was completed, media were discarded, and cells were post-treated by exposure to 0.5 mg/mL MTT (Invitrogen Inc., USA) and further incubated for 4 hours at 37°C. After incubation, the media were discarded, and cells were then washed with phosphate-buffered saline (PBS), followed by dissolving in 100 µL of dimethyl sulfoxide (DMSO) in the well and then incubated for 1 hour. Absorbance was subsequently determined using an enzyme-linked immunosorbent assay (ELISA) micro-plate reader (Spectra Max 190, USA) at a wavelength of 570 nm. Absorbance readings were converted into percentages of cells viable by comparing the treated group of each concentration with untreated controls. The IC₅₀ value was determined by performing linear regression between the concentration (x) and % cell viability (y). Cell viability % was measured as

$$[(\text{control} - \text{blank}) / (\text{sample} - \text{blank})] \times 100\%$$

where control, sample, and blank were the extract-untreated group, extract-treated group, and DMSO-only group, respectively

The IC₅₀ value represents the concentration that inhibited 50% of cell proliferation, and the experimental data for this research was obtained from three independent replication experiments. After percent cell viability was observed, their values were used for making the standard curve with a linear equation.

Cell migration assay

Cell migration is an important factor to consider when investigating the metastatic potential of cancer cells, and cell adhesion to extracellular matrices is considered to be a pivotal step in the invasive process of metastatic cells. The abovementioned DMSO (1 µL/mL), and extract dissolved in DMSO to concentrations of 500, 1000, and 2000 ng/mL as 0.5x IC₅₀, 1x IC₅₀, and 2x IC₅₀ concentrations, respectively, were cultured for 24 hours at 37°C.

Crude protein was extracted based on the protocol from

Yoon and Moon (2021). Centrifugation was performed to remove the medium, and cells were harvested and washed with cold PBS. Samples were incubated with 1 mL RIPA lysis buffer (containing 20 mM Tris, 5 mM EDTA, 10 mM Na₃PO₄, 100 mM NaF, 1% Nonidet P-40, and 1 mM PMSF at pH 7.4, RB4475, ProTech, Taiwan) and 10 µL of protease inhibitor (P8340, Sigma-Aldrich, USA). Thereafter, the cell layer was scratched with a cell scraper (BD Falcon, Corning, USA) to remove cells, placed into a 1.5 mL tube, and vortexed for 1 min. The crude protein was centrifuged at 13,000 rpm and 4°C for 1 min, and the supernatant then collected. Total soluble protein was quantified with a Bradford Protein Assay Kit (BioRad, USA). The amount of crude protein concentration was determined at an optical density of 595 nm using an ELISA reader. Commercial bovine serum albumin (BSA-A9418, Sigma) was used as a standard of linear regression equation.

HeLa cells were left untreated or treated with 0x, 0.5x, 1x, and 2x IC₅₀ concentrations of extract for 24 hours at 37°C, seeded in a 24-well plate, and grown overnight to confluency in DME-containing medium for 48 hours. After removing the medium, the confluent cell layer was scratched with a p200 Eppendorf pipette tip (Justus et al., 2014). Scraped cells were then washed away with PBS, and extract was added at 0x, 0.5x, 1x, and 2x IC₅₀ concentrations (with a final concentration of 4 µL/mL in the micro-plate wells). The plates were incubated for 0, 24, 48, and 72 hours, followed by counting the cells using ImageJ (1.05d, 64-bit Java 8, NIH, USA). The closure of the scratch wound area was photographed from three different images captured at a magnification of 10-fold using an inverted microscope. Triplicate controls and triplicate test samples were prepared for each time point in duration of 72 hours.

Determination of H₂O₂ content and lipid peroxidation

H₂O₂ content was determined according to Rahman et al. (2023). Briefly, 0.1 mL extract was homogenized in 0.1 mL of sodium phosphate buffer (10 mM, pH 7.0) containing 1 mM hydroxylamine and centrifuged at 8,000 rpm for 25 min at 4°C. One mL of the supernatant was then mixed with 0.5 mL of titanium chloride (TiCl₄, 0.1% v/v) dissolved in 20% (v/v) sulfuric acid (H₂SO₄), followed by centrifuging again at 3,300 rpm for 15 min at room temperature. The supernatant was transferred to a 96-well microplate (SpectraMax-190) and measured at 390 nm (A₃₉₀). Distilled-deionized water was used as the blank, and H₂O₂ concentration (µmol/mg protein) was calculated based on the standard calibration curve with 5, 10, 15, 20, 25 µmol H₂O₂.

Lipid peroxidation was evaluated by quantifying MDA, a commonly used indicator, based on Kosugi and Kikugawa (1985). In brief, 0.1 mL extract was homogenized

in 1.5 mL 5% (w/v) trichloroacetic acid (TCA) and centrifuged at 10,000 rpm for 5 min at 20°C. Supernatant (0.2 mL) was thoroughly mixed with 0.8 mL of 0.5% (W/V) thiobarbituric acid prepared in 20% (W/V) TCA, and incubated in a circulating dry bath (Sahara Rocker, Taiwan) at 95°C for 30 min. After cooling, samples were centrifuged again at 6,000 rpm for 10 min, and the supernatant then measured at 532 nm (A_{532}) using a spectrophotometer. TCA (5% w/v) was used as the blank. MDA concentration ($\mu\text{mol}/\text{mg}$) was calculated using the following equation:

$$A_{532} / 1.56 \times 10^5 \text{ (L mol}^{-1} \text{ cm}^{-1}) \times 1 \text{ (dilution factor)} / \text{protein (mg)}$$

Determination of antioxidant activity

To analyze the activity of ascorbate peroxidase (EC 1.11.1.11) and catalase (EC 1.11.1.6), 100 mg of leaf powder was extracted with 100 mM sodium phosphate buffer (pH 6.8) containing 1 mM phenylmethylsulfonyl fluoride (Sigma). Ascorbate peroxidase and catalase activity was measured according to the protocols of Nakano and Asada (1981). Absorbance was measured at 290 nm in a UV-visible spectrophotometer, and one unit was defined as micromoles of ascorbate oxidized per minute (extinction coefficient 2.8 mM cm^{-1}). Further-more, one unit of catalase activity was defined as the amount of enzyme catalyzing the decomposition of 1 nmol H_2O_2 per minute, with an extinction coefficient of 36 mM cm^{-1} . Glutathione reductase (EC 1.6.4.2) activity was measured by the GSH-dependent oxidation of NADPH according to Foyer et al. (1997). Superoxide dismutase (EC 1.15.1.1) activity was assayed by using the protocol of Dhindsa et al. (1981), and absorbance was recorded at 560 nm in a spectrophotometer.

Measurements of DPPH radical-scavenging ability and reducing power capacity

The antioxidant activity of extract was determined by DPPH scavenging capacity and reducing power ability. The assessment of scavenging activity against free DPPH radicals followed the protocols outlined by Wan et al. (2011). Briefly, 100 μL of the abovementioned control, DMSO, 0.5x IC_{50} , 1x IC_{50} , and 2x IC_{50} in culture medium DMEM were mixed with 100 μL 0.2 mM of DPPH (Sigma) and incubated in the dark at room temperature for 30 min. After centrifuging at 13,000 rpm for 1 min, the supernatant was placed into a 96-well ELISA microplate reader. The mixture was kept in the dark for 20 min, during which its color changed from purple (DPPH radical) to yellow (DPPH-H) following reduction. This change in color, quantified by a decrease in absorbance at wavelength 517 nm, indicates DPPH scavenging activity. A lower absorbance denotes higher DPPH-scavenging activity, and radical-scavenging activity is directly proportional to the concentrations of the extracts. Optical densities were calculated for % inhibition using the following

equation:

$$\text{DPPH radical scavenging activity (\%)} = [1 - (\text{sample}) / (\text{blank})] \times 100\%$$

Blanks were prepared in the same manner, using ethanol in place of extract samples. Triplicate controls and samples were prepared for each concentration presented in the form of mean \pm SD in the DPPH test.

G. divaricata extract reducing power activity was determined by the ferric reducing antioxidant power (FRAP) assay of Oyaizu et al. (1998). Briefly, 100 μL of control and extract samples were mixed with 100 μL of 0.2 M sodium phosphate buffer (pH 6.6) and 100 μL of 1% potassium hexacyanoferrate. The mixture was incubated at 50 °C for 20 min. After cooling on ice, 100 μL of 10% TCA was added to the mixture, vibrated for 5 min, and then centrifuged at 10,000 rpm for 10 min. The supernatant was then mixed with 100 μL 1% ferric chloride. After 10 min, absorbance at 700 nm using an ELISA plate reader was measured against a blank. FRAP (mmol vitamin C/ μg protein) was assessed using the following equation:

$$\text{FRAP} = \text{OD}_{700} / 1.77 \times 10^4 \text{ (L/mol/cm)} \times 0.5 \text{ (total volume, mL)} \times 0.1 \text{ (sample volume, mL)} \times 10 \text{ (dilution fold)} \times 0.1 \text{ (sample extract, g / mL)}$$

A blank was prepared using water without sample, and vitamin C used as a standard. The existence of reductants in the solution causes the reduction of the ferric to the ferrous form, and an increase in absorbance of the sample with various concentrations indicates a high reducing power capacity of the samples.

Western blotting

The effect of extract on NF- κ B was determined by immune-blotting according to Xu et al. (2020). Briefly, total proteins were extracted from the abovementioned controls, DMSO, 0.5x IC_{50} , 1x IC_{50} , and 2x IC_{50} of extract, by incubating with RIPA lysis buffer (pH 7.4) for 30 min on ice, with agitation every 10 min. After centrifugation for 15 min at 10,000 rpm and 4°C, the supernatant was collected and the protein concentration determined based on the Bradford protein assay (1976). Equal amounts of crude protein (20 $\mu\text{g}/\text{lane}$) were resolved in 10% sodium dodecyl sulfate-polyacrylamide gel electrophoresis (SDS-PAGE) gels (Mini-Protein Tetra, Bio-Rad, USA), separated at 20 mA for 2 hours, and transferred onto polyvinylidene difluoride membrane (Immobilon®-P, Millipore, USA) at 200 V and 4°C for 2 hours. After blocking with 5% skimmed milk at 4°C for 2 hours, membranes were incubated overnight at 4°C with primary antibodies against glyceraldehyde 3-phosphate dehydrogenase (GAPDH) XP® rabbit mAb (5174S, Cell Signaling Technology, USA) and anti-NF- κ B p65 (ab16502, Abcam, UK) at 1:1000 dilution. Subsequently, the blot was then incubated with goat anti-rabbit IgG H&L horseradish

peroxidase conjugated secondary antibody (ab205718, Abcam, UK) at 1:5000 dilution for 2 hours. Membranes were visualized and detected with a gel documentation system after incubating with chemiluminescent substrate solution for 10 sec (LumiFlash Mltima, Taiwan). Signal intensities were quantified using ImageJ software.

RNA isolation and gene expression analysis

Total RNA was extracted from control and extract - treated HeLa cells using the NucleoZOL extraction kit (Macherey-Nagel, Germany), and treated with DNase I to remove genomic DNA according to the instruction manual. RNA quality and the quality of each sample were determined by agarose electrophoresis and NanoDrop 2000 (Thermo Fisher Scientific, USA), respectively. Moreover, mRNAs were purified from 3 µg of total RNA using a MMLV Reverse Transcription Kit (Protech-Bio, Taiwan), followed by RNA converted to complementary (c)DNA under reaction buffers with 0.5 µL (75 ng) cDNA product, 1 µL 10 µM forward and reverse primers (Table I), 4 µL dNTP (2.5 mM), 5 µL Taq DNA polymerase, and 5 µL 10x Taq buffer (Yoon et al., 2021). PCR was carried out in an Eppen-

dorf Mastercycler Gradient Thermal Cycler (Germany) at the following thermal program: initial denaturation at 94°C for 10 min, followed by 30 cycles at 94°C for 30 sec, 57°C for 30 sec, and 72°C for 1 min, with a final extension at 72°C for 10 min. The products were electrophoretically separated on 1.5% agarose gels, predicted size of each gene (Table I) was verified with a 100bp DNA ladder marker (PT-M1-100T, ProTech, Taiwan), and sequences checked.

The relative changes in NF-κB pathway related gene expressions in response to various extract treatments were monitored by real-time (RT) quantitative (q)PCR and quantification of RNA levels. RT-qPCR reactions were assayed with 1 µL of cDNA product, 10 µL of 2X SyGreen mix (BIO Lo-ROX Biosystem Ltc., UK), and 0.8 µL of forward and reverse primers (the final concentration of each the primer was 10 µM) in ddH₂O to a total volume of 20 µL. The RT-qPCR was performed on the ABI 7500 Real-time system (ABI, USA), and the PCR program was set as denature at 95°C for 5 sec, annealing at 60°C for 30 sec, and extension at 72°C performed for 45 cycles. The sequences of paired primers used in this study are presented in Table I. GAPDH was used as a reference gene for normalization, and differential gene expression calculated as follows (Livak and Schmittgen, 2001):

$$2^{-\Delta\Delta Ct} = (Ct_{\text{target gene}} - Ct_{\text{GAPDH}})_{\text{treatment}} - (Ct_{\text{target gene}} - Ct_{\text{GAPD}})_{\text{control}}$$

The control was the cells cultured in complete DMEM, while the treatments were cells cultured in DMSO containing 500, 1,000, and 2,000 ng/mL of extract at 0.5 IC₅₀, 1 IC₅₀, and 2 IC₅₀ concentrations. Triplicate control samples and triplicate test samples were prepared and presented as mean ± SD. The independent Student's *t*-test was used to determine differences between control and test groups, and statistical significance was considered when *p*<0.05.

Statistical analysis

All experiments in this study are tabulated as the mean value with the standard deviation (SD) of at least three independent replications, as stated in each figure legend. Student's *t*-test was performed using SAS statistical analysis software Version 9.4 (SAS Institute Inc, USA) to evaluate the statistical comparison of the mean values between two groups. A *p*-value of less than 0.05 was considered statistically significant.

Results

Effects on the viability of HeLa cells

After measuring absorbance using an ELISA reader, the untreated controls showed the highest cell viability at 100%, and extract significantly reduced HeLa cell viability in a concentration-dependent manner (Figure 1A). In the group treated with 250 ng of extract, the

Table I

Paired primers for RT-qPCR used

Gene	Oligonucleotide sequence	Size (bp)
GAPDH	F 5'-TGTAGGCTCATTTCAGGGG-3'	317
	R 5'-TCCCATTCCCAGCTCTCAT-3'	
Bcl-2	F 5'-AAAGCCAGCTTCCCCAATGA-3'	596
	R 5'-CCCAACTGCAGGATGCCTTT-3'	
Bax	F 5'-AACCCCTTCAGGGAGTCA-3'	423
	R 5'-CTGCAGGGGTTGATACCAC-3'	
IL-8	F 5'-GGAAAGTTCAGGTGTAGGA-3'	256
	R 5'-CACCCAGTTTTCTTGGGGT-3'	
TNF-α	F 5'-GAGACAGATGTGGGGTGTGAG-3'	642
	R 5'-TCCTAGCCCTCCAAGTTCCA-3'	
IL-1β	F 5'-GACTTCCCCATGACGGCTAC-3'	507
	R 5'-AACCTCTTCGAGGCACAAGG-3'	
Iκβa	F 5'-TGGCAAATAGCAGAGGCTCC-3'	345
	R 5'-TGGGGTTTTCCCTCTCTCC-3'	
COX-2	F 5'-AATGGCTGAGTATGGCACCC-3'	783
	R 5'-GCTGTCCTTTAATTGCAGCAAATCC-3'	
IL-6	F 5'-GCTTCCCTCAGGATGCTTGT-3'	405
	R 5'-ATTAAGTGGGGTGGCTGCTC-3'	
NF-κB	F 5'-ATGTGGAGATCATTGAGCAGC-3'	151
	R 5'-CCTGGTCTGTGTAGCCATT-3'	
IKKα	F 5'-GGCTTCGGGAACGCTGTGTC-3'	87
	R 5'-TTTGGTACTTAGCTCTAGGCCGA-3'	

F, Forward; R, Reverse. All genes were searched from NCBI (<http://www.ncbi.nlm.nih.gov/>) with *Homo sapiens* (human), and shown as: GAPDH (NC_000012.12), Bcl-2 (NC_000018.10), Bax (NC_000019.10), IL-8 (NC_000004.12), TNF-α (NC_000006.12), IL-1β (NC_000002.12), Iκβa (NC_000014.9), COX-2 (NC_000001.11), IL-6 (NC_000007.14), NF-κB

HeLa cell viability level was reduced to 88.7%, and subsequent concentrations showed a decreasing trend, indicating that extract exerted different cytotoxicity effects on the proliferation of HeLa cells. Extract from 500 to 2,750 ng/mL significantly inhibited the viability of HeLa cells, showing that higher concentrations of extract treatments resulted in progressively lower HeLa cell viability values with higher inhibition rates (%) of repressed cell proliferation. In particular, treatment with 2,000 ng/mL of extract tremendously reduced the viability of HeLa cells (17.9%) compared with the control group (100%). The IC_{50} value was found at 1,000 ng/mL. The HeLa cell inhibition rates ranged from 250 to 2,750 ng/mL of extract treatment, the linear curve of regression was $y = 0.00484x - 0.3557$, with $R^2 = 0.9375$ (Figure 1B). The inhibitory ability of extract on HeLa cells at a concentration of 1,500 ng/mL entered a plateau which was much stronger than that of other concentrations with 80% of cell inhibition.

Potential effect of extract on metastasis

The inhibition of HeLa cell migration was examined using a wound healing assay. The morphologies of HeLa cells were different between controls and extract-treated samples for 0 ~ 72 hours, and all HeLa cells exposed for 24, 48, and 72 hours had significant reductions in adhesion compared to 0 hour exposure (Figure 2A). Significant reductions in adhesion were observed in all HeLa cells exposed for 24, 48, and 72

hours ($20.7 \pm 0.7\%$, $0.8 \pm 0.1\%$, and $0.02 \pm 0.01\%$, respectively) compared to 0 hour at 100% relative wound closure (Figure 2B). After exposure of HeLa cells to DMSO (without extract-treated group) for 24, 48, and 72 hours, significant reduction in adhesion was observed in all cells at $22.9 \pm 1.3\%$, $1.1 \pm 0.1\%$, and $0.01 \pm 0.002\%$, respectively, compared to 0 hour at 100%. The extract at $0.5x IC_{50}$ (500 ng/mL) significantly reduced the relative wound closure for 24-, 48-, and 72-hours periods at $34.8 \pm 1.2\%$, $10.0 \pm 1.7\%$, and $2.1 \pm 0.1\%$, respectively, compared to 0-hour culture period. After 24, 48, and 72 hours of incubation, $1x IC_{50}$ extract (1,000 ng/mL)-treated cells showed significantly decreased levels of adhesion ($57.2 \pm 2.9\%$, $21.8 \pm 1.0\%$, and $28.9 \pm 1.4\%$) compared to 0 hour of incubation. HeLa cells treated with $2x IC_{50}$ extract (2,000 ng/mL) for 24, 48, and 72 hours showed less wound healing ($56.4 \pm 2.9\%$, $70.7 \pm 2.5\%$, and $73.6 \pm 3.4\%$) than that of the 0 hour group.

Contents of H_2O_2 , MDA, antioxidant activity, and ability of DPPH-scavenging and FRAP

Table II illustrates *G. divaricata* extract significantly altered oxidative stress markers in HeLa cells in a concentration- and time-dependent manner. H_2O_2 levels increased in the $0.5x IC_{50}$ and $1x IC_{50}$ treatments after 0.5 hour, whereas prolonged exposure resulted in higher H_2O_2 accumulation in the $1x IC_{50}$ and $2x IC_{50}$ groups. MDA contents generally decreased after 24 hours of incubation, with the lowest level observed in the $0.5x IC_{50}$ treatment, indicating reduced lipid peroxidation.

Among antioxidant enzymes, ascorbate peroxidase activity was not significantly affected by the extract treatment. In contrast, catalase and glutathione reductase activities were reduced at higher extract concentrations, whereas superoxide dismutase activity increased significantly and showed a dose-dependent response. The highest superoxide dismutase activity was detected in the $2x IC_{50}$ treatment after 24 hours of exposure.

The antioxidant potential of extract was further supported by DPPH and FRAP assays. The extract-treated cells exhibited enhanced radical-scavenging activity, particularly in the $2x IC_{50}$ treatment, demonstrating that the extract effectively modulated cellular redox status and antioxidant defense mechanisms in HeLa cells.

Western blot of NF- κ B from extract

To further elucidate NF- κ B mechanisms in extract, the relative amounts (%) of NF- κ B accumulation in extract were analyzed by western blot (Figure 3). NF- κ B proteins responded differently in the various concentrations of extract-treated HeLa cells, and a clear NF- κ B band (60 kDa) appeared in all concentrations (DMSO, $0.5x IC_{50}$, $1x IC_{50}$, and $2x IC_{50}$) and controls. Extract at all concentrations ($0.5x$, $1x$, and $2x IC_{50}$) was significantly lower than in controls. As extract concentration increased, the NF- κ B expression level decreased

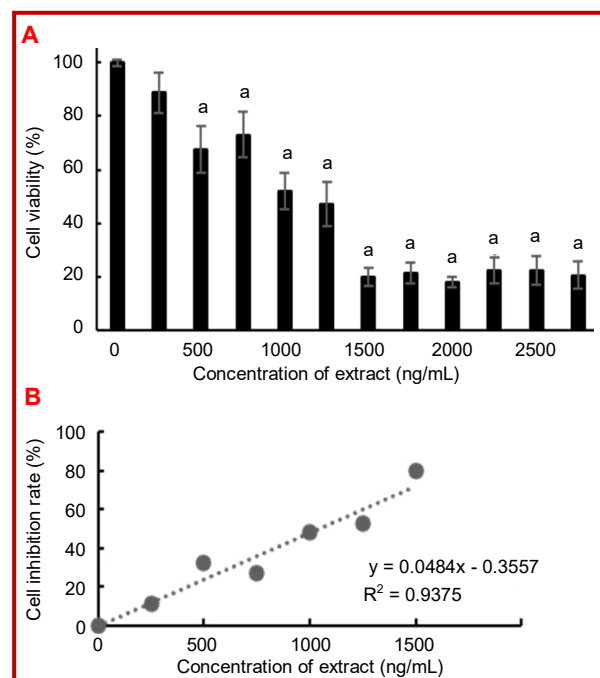


Figure 1: Effect of different concentrations (250 ~ 2,750 ng/mL) of extract treatments on human HeLa cell viability (A) and cell inhibition rate (B) at 48 hours, evaluated by MTT assay. Data are mean \pm SD; $n=4$. $p<0.05$ was considered as statistically significant (superscript "a") compared between control and each concentration of extract treatment using Student's *t*-test

from controls at 100%, dropping to 20.0% for 2-IC₅₀ extract.

Effects of extract on NF- κ B-related gene expressions in the NF- κ B signaling pathway

Different NF- κ B related genes involved with extract show differential gene expressions. Human HeLa cells treated with 0.5x, 1x, and 2x IC₅₀ of extract showed significant down-regulation of Bcl-2, Bax, IL-8, IKK α ,

NF- κ B, and IL-1 β genes (Figure 4A, B, E, G, H, and K, respectively) compared to controls. However, RNA expression of the Bcl-2/Bax gene ratio (Figure 4C) was significantly lower in human HeLa cells treated with 0.5-IC₅₀ of extract than in controls. Significant down-regulations of IL-6, COX-2, and TNF- α genes (Figure 4D, F, and J, respectively) were found in human HeLa cells treated with 1- and 2-IC₅₀ of extract compared to controls. Figure 4I shows that RNA abundance of the

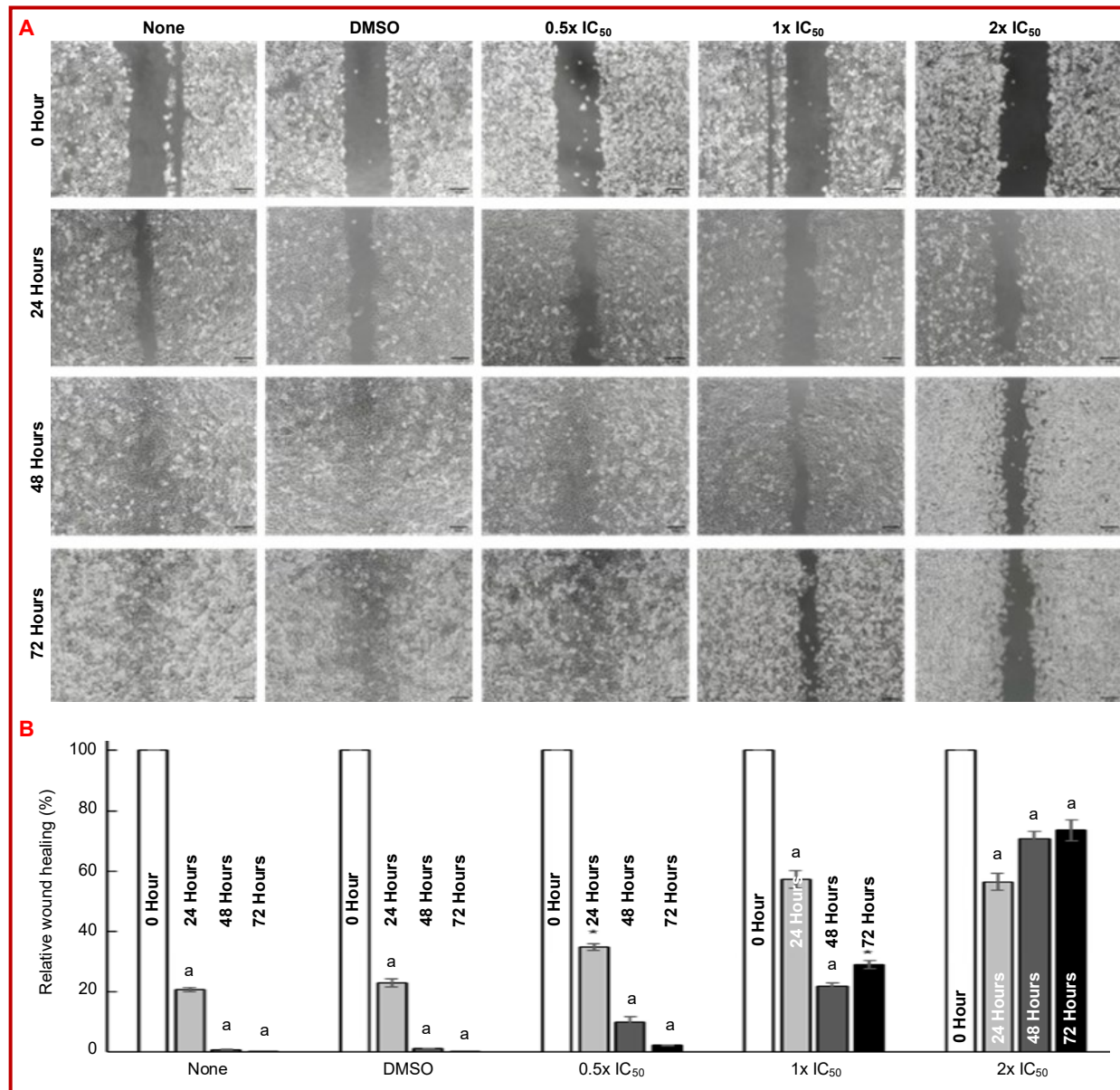


Figure 2: Cell migration assay. Cultured HeLa cells were scratched with a pipette tip, followed by culturing in complete DMEM supplemented with extract at concentrations of 0.5x, 1x, and 2x IC₅₀ (calculated as 500, 1,000, and 2,000 ng/mL of extract) and incubated for 72 hours. Controls were cells cultured in complete DMEM, while DMSO was a maintenance medium of extract - untreated cells. Data were collected on incubation day (0 hour) and after 72 hours of incubation, and photographed at 100x magnification. The scale bar at the right corner indicates 20 μ m (A). The closure of the scratch relative wound area (%) from cell morphology photographs was quantified using ImageJ (B). Among each extract concentration treatment, period durations of 24, 48, and 72 hours were compared to 0 (100% as control) using Student's t-test. Triplicate control and triplicate test samples were prepared for each time point, and superscript "a" indicate significance levels of $p \leq 0.05$. Vertical bars indicate standard deviations

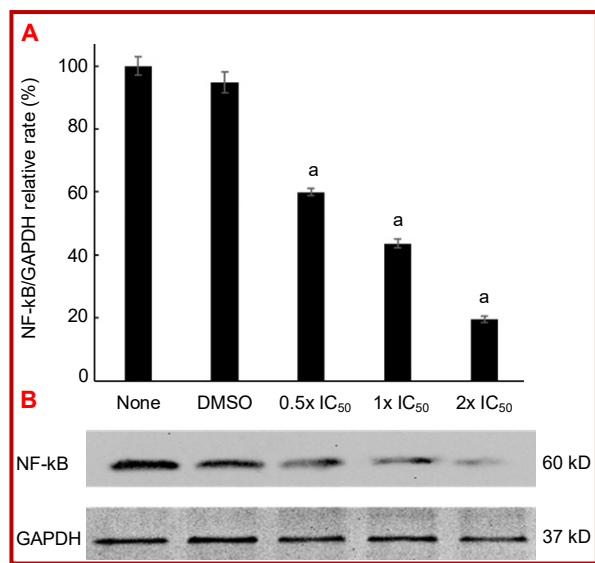


Figure 3: Western blot analysis of HeLa cells cultured in control, DMSO (complete DMEM), and DMSO containing 500, 1,000, and 2,000 ng/mL of extract as 0.5x IC₅₀, 1x IC₅₀, and 2x IC₅₀ concentrations, respectively, probed by NF-κB and GAPDH as antibodies. Forty micrograms of total protein were used for each sample during SDS-PAGE separation, and the molecular weights of NF-κB and GAPDH (used as an internal control) are 60 and 37 kDa, respectively, as indicated by two arrowheads. A representative image of three similar results is shown. Bands with NF-κB/GAPDH relative rate (%) were visualized and detected with a chemiluminescent substrate, and intensities are quantified using ImageJ software, followed by representation on the bar graphs. ^ap<0.05 compared to the control is expressed as 100%

IκBα gene was significantly up-regulated in DMSO and 0.5-IC₅₀ of extract-treated cells in comparison to controls, while the IκBα gene significantly decreased its expression in 1x and 2x IC₅₀ of extract treatments compared to controls. In human HeLa cells treated with 0.5x IC₅₀ of extract, the expression of the Bcl-2/Bax gene decreased by 0.5x compared to controls (Figure 4C). Nevertheless, under human HeLa cells treated with 1-IC₅₀ of extract, the expressions of Bcl-2, Bax, IL-6, IL-8, COX-2, IKKα, NF-κB, IκBα, TNF-α, and IL-1β genes decreased by 0.034, 0.037, 0.119, 0.011, 0.048, 0.055, 0.065, 0.101, 0.561, and 0.083 times (Figure 4A, B, and D-K), respectively, compared with controls.

Discussion

The present study demonstrates that the extract possesses significant anti-cancer and antioxidant activities in HeLa cells. Treatment reduced cell viability, inhibited

Table II

Effects of *Gynura divaricata* on oxidative stress and antioxidant parameters

Parameter	Time (hour)	Control	DMSO (Vehicle control)	0.5x IC ₅₀	1x IC ₅₀	2x IC ₅₀
H ₂ O ₂ (μmole/μg protein)	0.5	2.2 ± 0.0	2.4 ± 0.0	3.8 ± 0.0 ^a	2.5 ± 0.1 ^a	2.0 ± 0.0 ^a
	24	1.6 ± 0.1	1.7 ± 0.1	1.6 ± 0.0	2.7 ± 0.0 ^a	2.6 ± 0.1 ^a
MDA (μmole/mg protein)	0.5	14.2 ± 0.2	15.9 ± 0.4 ^a	19.0 ± 0.3 ^a	17.7 ± 0.4 ^a	14.4 ± 0.3
	24	12.4 ± 0.1	13.8 ± 0.0 ^a	11.2 ± 0.2 [*]	12.4 ± 0.0	14.1 ± 0.3 ^a
Ascorbate peroxidase (Unit/μg protein)	0.5	94.1 ± 33.2	83.9 ± 24.4	69.7 ± 23.1	73.3 ± 13.0	95.9 ± 24.0
	24	124.4 ± 26.7	133.6 ± 12.8	136.8 ± 21.9	140.1 ± 17.4	126.1 ± 36.1
Catalase (Unit/μg protein)	0.5	4.7 ± 0.2	6.1 ± 0.2	4.5 ± 0.3	4.3 ± 0.4	3.8 ± 0.2 ^a
	24	11.9 ± 0.5	16.2 ± 1.2	6.4 ± 0.7 ^a	4.7 ± 0.8 ^a	3.7 ± 0.5 ^a
Glutathione reductase (Unit/μg protein)	0.5	3.0 ± 0.1	3.4 ± 0.1	2.8 ± 0.2	2.4 ± 0.2 ^a	2.6 ± 0.4
	24	6.7 ± 0.4	6.9 ± 0.5	6.0 ± 0.5	5.4 ± 0.3 ^a	5.4 ± 0.2 ^a
Superoxide dismutase (Unit/μg protein)	0.5	3.4 ± 0.5	4.5 ± 0.7	6.4 ± 0.4 ^a	6.7 ± 0.7 ^a	9.5 ± 1.2 ^a
	24	7.8 ± 0.7	7.5 ± 0.8	8.1 ± 0.9	8.4 ± 0.7	12.3 ± 1.2 ^a
DPPH scavenging (%)	0.5	6.2 ± 0.9	7.9 ± 1.1	10.1 ± 0.9 ^a	10.1 ± 0.9 ^a	9.8 ± 1.8 ^a
	24	8.1 ± 0.5	6.1 ± 0.8	7.7 ± 0.8	6.1 ± 0.9	10.9 ± 0.7 ^a
FRAP (μM Fe ²⁺ equivalent)	0.5	24.8 ± 0.3	26.7 ± 0.3 ^a	22.1 ± 0.4 ^a	29.7 ± 0.7 ^a	26.1 ± 0.4 ^a
	24	24.9 ± 0.2	23.2 ± 0.2 ^a	20.6 ± 0.2 ^a	22.2 ± 0.3 ^a	22.3 ± 0.2 ^a

Values are mean ± SD; n=9; HeLa cell culture time period of 0.5 hour or 24 hours. Controls (without extract) were cells cultured in complete DMEM, while treatments were cells cultured in DMSO containing 500, 1,000, and 2,000 ng/mL of extract as 0.5x IC₅₀, 1x IC₅₀, and 2x IC₅₀ concentrations; ^ap<0.05 is considered statistically significant in comparing controls and treatments of extract with HeLa cells across the same culture time period at 0.5 hour or 24 hours by Student's *t*-test

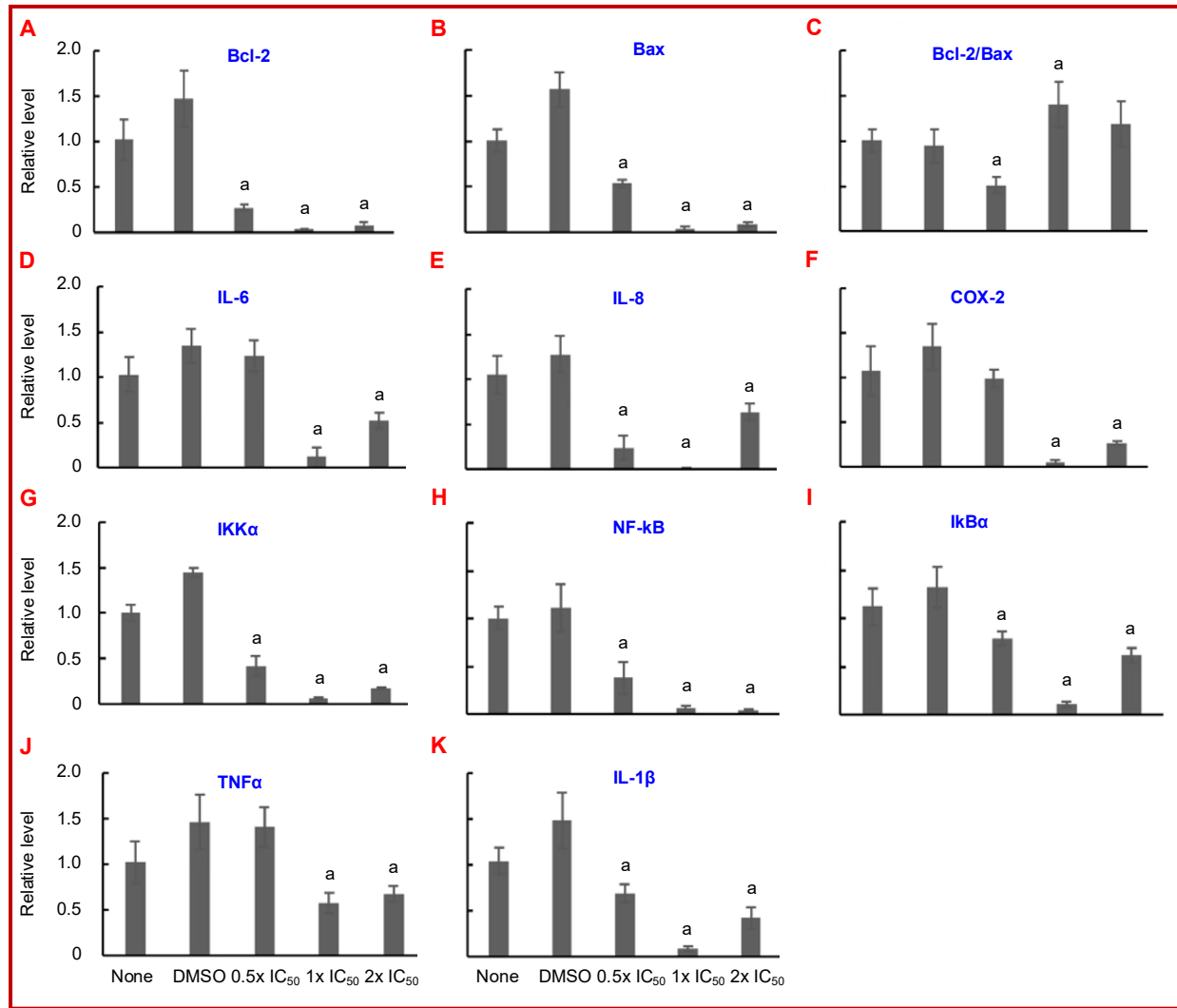


Figure 4: Relative RNA expressions of Bcl-2, Bax, Bcl-2/Bax ratio, IL-6, IL-8, COX-2, IKK α , NF- κ B, I κ B α , TNF- α , and IL-1 β genes in control, DMSO, and extract-treated HeLa cells. Relative amounts were calculated using the $2^{-\Delta\Delta C_t}$ method and normalized with respect to GAPDH co-amplified as an internal control. Values are the means of three replicates with the corresponding standard deviation. The gene expression is compared to controls using Student's t-test, and superscript "a" indicates a significance level of $p \leq 0.05$. Controls were cells cultured in complete DMEM, while the treatments were cells cultured in DMSO containing 500, 1,000, and 2,000 ng/mL of extract as 0.5x IC₅₀, 1x IC₅₀, and 2x IC₅₀ concentrations

cell migration, modulated oxidative stress biomarkers, enhanced antioxidant defense responses, and suppressed NF- κ B signaling. The extract also regulated the expression of apoptosis- and inflammation-related genes, including Bcl-2, Bax, TNF- α , IL-6, IL-8, COX-2, IKK α , I κ B α , and IL-1 β . Collectively, these findings suggest that the extract may inhibit cervical cancer progression through coordinated regulation of oxidative stress, inflammatory signaling, and cell survival pathways.

The observed cytotoxic and anti-migratory effects are consistent with previous reports showing that plant-derived bioactive compounds suppress HeLa cell proliferation and migration (Rifai et al., 2024). Similar to

earlier studies, the extract enhanced antioxidant defense mechanisms and reduced oxidative damage, supporting the role of antioxidant enzymes in protection against reactive oxygen species (Nguyen et al., 2018; Arora and Bhalata, 2017; Lin et al., 2020). The reduction of NF- κ B signaling and inflammatory gene expression also agrees with reports indicating that inhibition of NF- κ B is associated with decreased tumor growth, inflammation, and metastatic potential (Garcia-Pineres et al., 2009; Napetschnig and Wu, 2013; Li et al., 2016). Furthermore, the modulation of apoptosis-related genes is in accordance with studies showing that alterations in the Bcl-2/Bax regulatory network contribute to cancer cell death and growth suppression (Ryu et al., 2017; Yao et al., 2017; Maqbool et al., 2025). However, unlike some

studies that reported selective activation of pro-apoptotic signaling, the present study observed simultaneous regulation of both pro- and anti-apoptotic genes, suggesting a more complex mechanism of cellular adaptation and stress response.

The extract appears to influence intracellular redox balance by scavenging reactive oxygen species (ROS) and enhancing endogenous antioxidant defenses, thereby limiting lipid peroxidation and oxidative injury (Kumaran and Karunakaran, 2006; Dantu et al., 2012). Reduced oxidative stress may subsequently suppress NF- κ B activation, leading to decreased expression of inflammatory mediators and adhesion-related molecules involved in cancer progression (Chao et al., 2013b; Cruceriu et al., 2020; Hayden and Ghosh, 2011). In addition, modulation of the Bcl-2/Bax signaling network may contribute to the regulation of apoptosis and cellular homeostasis (Gansukh et al., 2019; Kang et al., 2011; Chung and Yu, 2013). The coordinated down-regulation of NF- κ B-associated genes may further impair cancer cell migration and metastatic behavior. These observations support the hypothesis that the extract exerts its anti-cancer effects through integrated regulation of oxidative stress, inflammation, apoptosis, and NF- κ B-dependent signaling pathways. Nevertheless, the active constituents responsible for these effects remain unclear and require further investigation through mechanistic and *in vivo* studies.

Extract mitigates ROS accumulation and protects cellular components through the regulation of NF- κ B-associated signaling, thereby suppressing cancer cell progression. Although extract demonstrates promising anti-cancer potential as a plant-derived therapeutic agent, its active constituents and precise molecular mechanisms within the NF- κ B pathway remain unclear.

Limitation of this study includes *in vivo* studies that requires to elucidate its bioactivity, optimal dosage, and therapeutic efficacy as a safe natural antioxidant for clinical applications.

Conclusion

HeLa cell viability % significantly decreased upon exposure to increasing extract concentrations, showing that extract inhibited cell proliferation and increased the death rate of HeLa cells at high extract concentrations. Various extract dose-dependent manners also improved the antioxidant profiles of HeLa cells and exhibited relatively better DPPH radical scavenging activity and reducing power capacity to cope with oxidative stress.

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Ethical Issue

The development, acquisition, authentication, cryopreservation, and transfer of cell lines between laboratories were followed according to the guidelines published in British Journal of Cancer, 2014

Conflict of Interest

Authors declare no conflict of interest

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