

Galangin (3,5,7-Trihydroxyflavone) Shields Human Keratinocytes from Ultraviolet B-Induced Oxidative Stress

Susara Ruwan Kumara Madduma Hewage¹, Mei Jing Piao¹, Ki Cheon Kim¹, Ji Won Cha¹, Xia Han¹, Yung Hyun Choi², Sungwook Chae³ and Jin Won Hyun^{1,*}

¹Jeju National University School of Medicine, Jeju 690-756, ²Department of Biochemistry, College of Oriental Medicine, Donggeui University, Busan 614-052, ³Aging Research Center, Korea Institute of Oriental Medicine, Daejeon 305-811, Republic of Korea

Abstract

Most skin damage caused by ultraviolet B (UVB) radiation is owing to the generation of reactive oxygen species. Phytochemicals can act as antioxidants against UVB-induced oxidative stress. This study investigated the protective effects of the flavone galangin against UVB-induced oxidative damage in human keratinocytes. Galangin efficiently scavenged free radicals and reduced UVB-induced damage to cellular macromolecules, such as DNA, lipids, and proteins. Furthermore, galangin rescued cells undergoing apoptosis induced by UVB radiation *via* recovering mitochondrial polarization and down-regulating apoptotic proteins. These results showed that galangin protects human keratinocytes against UVB radiation-induced cellular damage and apoptosis *via* its antioxidant effects.

Key Words: Apoptosis, Galangin, Human keratinocytes, Oxidative damage, Ultraviolet B

INTRODUCTION

Solar radiation can be divided into three main types based on its wavelength, namely, ultraviolet (UV), visible light, and infrared (Lyons and O'Brien, 2002). Of these, UV radiation is the most responsible for photoaging and skin cancer. UV radiation is subdivided into UVA (320-400 nm), UVB (280-320 nm), and UVC (200-280 nm). The ozone layer reflects UVC radiation, meaning only UVA and UVB reach the earth's surface. UVB radiation is particularly absorbed by human skin and causes erythema, burns, immune suppression, and skin cancer (Park *et al.*, 2013). Although UVA accounts for the majority of UV radiation that reaches the earth's surface and can penetrate the skin deeper than UVB, it is weakly carcinogenic and causes aging and wrinkling of the skin (Yoshikawa *et al.*, 1990; Donawho *et al.*, 1996; Matsumura and Ananthaswamy, 2004).

UVB directly or indirectly damages skin cells *via* the formation of cyclobutane pyrimidine dimers (CPDs) and pyrimidine-pyrimidone(6-4) photoproducts or *via* the generation of reactive oxygen species (ROS), such as hydroxyl radicals ($\cdot\text{OH}$), superoxide anions ($\cdot\text{O}_2^-$), hydrogen peroxide (H_2O_2), and singlet oxygen ($^1\text{O}_2$) (Cunningham *et al.*, 1985; Hattori *et al.*, 1996; Meeran *et al.*, 2008). UVB-exposed cells generate ROS by activating specific small molecules such as riboflavin, tryptophan, and porphyrin (Ikehata and Ono, 2011). The antioxidant defense system in cells balances ROS production; however, when levels of ROS are elevated, this antioxidant defense system is overwhelmed, resulting in oxidative stress. Uncontrolled release of ROS causes single- and double-strand DNA breaks and DNA-protein cross-linking (Caldwell *et al.*, 2007). ROS also attack important cellular structural and functional molecules, such as lipids and proteins, causing the malfunction of cellular activities, finally leading to apoptosis, a process of programmed cell death (Tsoyi *et al.*, 2008; Dhumrongvaraporn and Chanvorachote, 2013).

With the increased occurrence of skin cancers and other damaging effects of UVB exposure, the protection of skin from UVB-induced oxidative cellular damage has become a key consideration in the pharmaceutical industry. Phytochemicals are well-known for their protective effects against oxidative stress in the skin (Sumiyoshi and Kimura, 2009). Galangin (3,5,7-trihydroxyflavone, Fig. 1) is a type of flavonoid that is commonly found in *Alpinia officinarum* and *Helichrysum aureonitens* (Afolayan and Meyer, 1997; Ciolino and Yeh, 1999). Galangin has antibacterial (Cushnie and Lamb, 2005; 2006)

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Received Nov 25, 2014 Revised Jan 13, 2015 Accepted Jan 20, 2015
Published online Mar 1, 2015

***Corresponding Author**

E-mail: jinwonh@jejunu.ac.kr

Tel: +82-64-754-3838, Fax: +82-64-702-2687

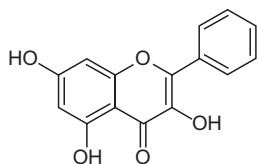


Fig. 1. Chemical structure of galangin (3,5,7-trihydroxyflavone).

and antiviral (Afolayan *et al.*, 1997) properties and suppresses breast tumor cell growth (So *et al.*, 1996; Diffey, 2004). However, the cytoprotective effects of galangin against UVB-induced oxidative damage in human keratinocytes have not been studied. Therefore, the objective of this study was to investigate the protective effects of galangin against UVB-induced oxidative stress in human keratinocytes.

MATERIALS AND METHODS

Reagents

Galangin was purchased from Santa Cruz Biotechnology Inc. (Dallas, TX, USA). [3-(4,5-Dimethylthiazol-2-yl)-2,5-diphenyltetrazolium] bromide (MTT), 1,1-diphenyl-2-picrylhydrazyl (DPPH), N-acetyl cysteine (NAC), 5,5-dimethyl-1-pyrroline-N oxide (DMPO), 2',7'-dichlorodihydrofluorescein diacetate (DCF-DA), and Hoechst 33342 were purchased from Sigma-Aldrich Inc. (St. Louis, MO, USA). 5,5',6,6'-Tetrachloro-1,1',3,3'-tetraethyl-benzimidazolylcarbocyanine iodide (JC-1) was purchased from Invitrogen (Carlsbad, CA, USA). Primary antibodies for Bax and Bcl-2 were purchased from Santa Cruz Biotechnology Inc. (Dallas, TX, USA). Primary antibodies for caspase-3 and caspase-9 were purchased from Cell Signaling Technology (Danvers, MA, USA). All other chemicals and reagents were of analytical grade.

Cell culture and UVB exposure

The human keratinocyte cell line HaCaT was obtained from Amore Pacific Company (Yongin, Korea) and maintained at 37°C in an incubator with a humidified atmosphere of 5% CO₂. Cells were cultured in RPMI 1640 medium containing 10% heat-inactivated fetal bovine serum, streptomycin (100 µg/ml), and penicillin (100 units/ml). Cells were exposed to UVB at a dose of 30 mJ/cm². The CL-1000M UV Crosslinker (UVP, Upland, CA, USA) was used as the UVB source and delivered a UVB energy spectrum of 280-320 nm.

Cell viability assay

The influence of galangin on cell viability was examined using the MTT assay. Cells were seeded in a 96-well plate at a density of 1×10⁵ cells/ml. After 24 h, galangin was added to a final concentration of 20, 40, 80, or 100 µM and cells were incubated for 24 h. MTT stock solution (50 µl, 2 mg/ml) was added to each well to yield a final reaction volume of 200 µl. The supernatant was aspirated 4 h later and formazan crystals were dissolved in 150 µl of dimethylsulfoxide (DMSO). The absorbance at 540 nm was read using a scanning multi-well spectrophotometer (Carmichael *et al.*, 1987).

Detection of DPPH radicals

The ability of galangin to scavenge DPPH radicals was as-

essed. Cells in a 96-well plate were treated with 20, 40, 80, or 100 µM of galangin or 1 mM of NAC. DPPH dissolved in ethanol (0.1 mM) was added to each well to yield a total volume of 200 µl. After shaking for 3 h, unreacted DPPH was detected by measuring the absorbance at 520 nm using a spectrophotometer.

Detection of hydroxyl radicals

Hydroxyl radicals generated by the Fenton reaction (H₂O₂+FeSO₄) were reacted with DMPO. The resultant DMPO/•OH adducts were detected using an electron spin resonance (ESR) spectrometer (Li *et al.*, 2004). The ESR spectrum was recorded 2.5 min after a phosphate buffer solution (pH 7.4) was mixed with 0.02 ml each of 0.3 M DMPO, 10 mM FeSO₄, 10 mM H₂O₂, and 40 µM of galangin. The ESR spectrometer parameters were set as follows: central magnetic field, 336.8 mT; power, 1.00 mW; frequency, 9.4380 GHz; modulation width, 0.2 mT; amplitude, 600; sweep width, 10 mT; sweep time, 0.5 min; time constant, 0.03 sec; and temperature, 25°C.

Detection of intracellular ROS

DCF-DA fluorescence was detected to measure intracellular ROS generated by H₂O₂ or UVB (Rosenkranz *et al.*, 1992). Cells were seeded at a density of 1.5×10⁵ cells/ml and incubated at 37°C for 24 h. Galangin (40 µM) or NAC (1 mM) was added to each well. After 1 h, cells were treated with H₂O₂ (1 mM) or exposed to UVB. After 30 min, H₂O₂-treated cells were treated with DCF-DA (25 µM) and incubated for another 20 min. UVB-treated cells were incubated for 24 h, treated with DCF-DA (50 µM), and incubated for a further 30 min. Fluorescence of 2',7'-dichlorofluorescein (DCF) was detected and quantified using a PerkinElmer LS-5B spectrofluorometer (PerkinElmer, Waltham, MA, USA). Intracellular ROS scavenging effect of galangin (%)=(absorbance of control cells - absorbance of galangin- or NAC-treated cells)/absorbance of control cells×100. Only H₂O₂-or UVB-treated cells were considered as controls.

Single cell gel electrophoresis (Comet assay)

DNA damage caused by oxidative stress was detected by the comet assay (Singh, 2000; Rajagopalan *et al.*, 2003). Cells were seeded at a density of 1×10⁵ cells/ml and incubated at 37°C for 24 h. Cells were treated with galangin (40 µM) and after 1 h, exposed to UVB (30 mJ/cm²). The cell suspension was collected and mixed with 120 µl of 0.7% low melting agarose (LMA) at 37°C. The mixture was spread on a fully frosted microscope slide pre-coated with 200 µl of 1% normal melting agarose. After this had solidified, a further 170 µl of LMA was applied to the slide. Slides were immersed in lysis solution (2.5 M NaCl, 100 mM Na-EDTA, 10 mM Tris, 1% Trion X-100, and 10% DMSO, pH 10) for 90 min at 4°C. Slides were then immersed in an unwinding buffer (300 mM NaOH and 10 mM Na-EDTA, pH 13) for 30 min at 4°C. Slides were subjected to electrophoresis in unwinding buffer solution with an electrical field of 300 mA and 25 V for 20 min at room temperature. The slides were washed three times with neutralizing buffer (0.4 M Tris, pH 7.5) for 10 min each time and then washed with 70% ethanol for 5 min. Slides were stained with 70 µl of ethidium bromide and observed under a fluorescence microscope using an image analyzer (Kinetic Imaging, Komet 5.5, UK). The tail length and the percentage of fluorescence in comet tails were recorded for 50 cells per slide.

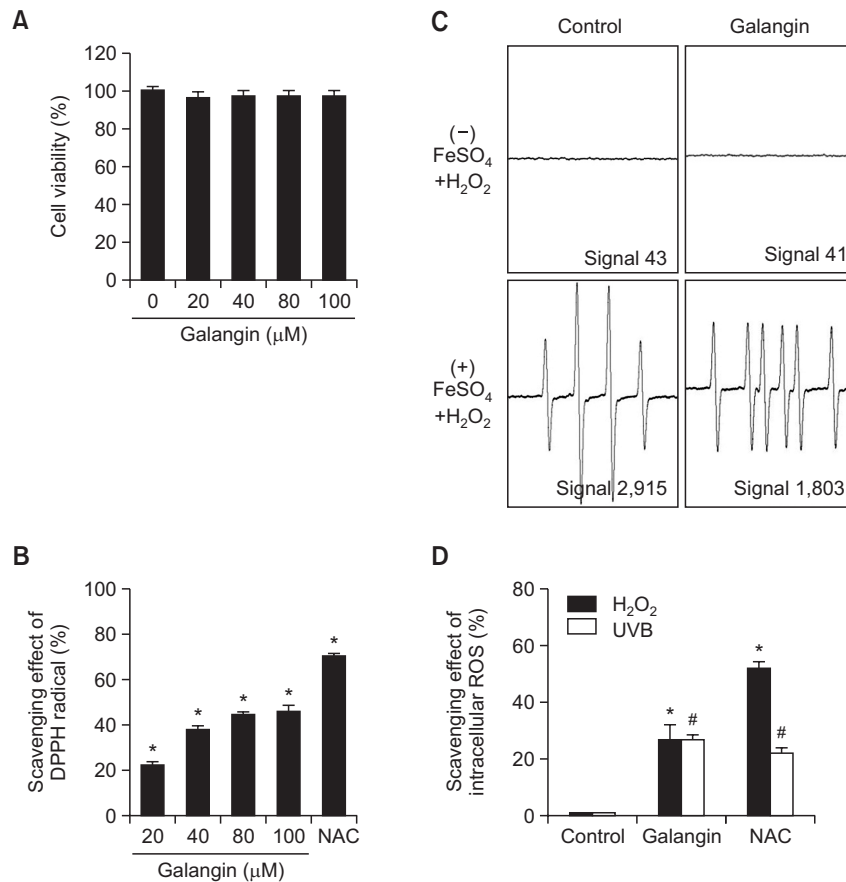


Fig. 2. Galangin scavenges reactive oxygen species (ROS). (A) HaCaT cells were treated with various concentrations of galangin for 24 hours and cell viability was measured by the [3-(4,5-dimethylthiazol-2-yl)-2,5-diphenyltetrazolium] bromide (MTT) assay. (B) Levels of the 1-diphenyl-2-picrylhydrazyl (DPPH) radical were measured spectrophotometrically at 520 nm. N-acetyl cysteine (NAC, 1 mM) was used as the positive control. *Significantly different from the control group ($p < 0.05$). (C) Hydroxyl radical-scavenging ability was evaluated using the Fenton reaction system. (D) The ability of galangin to scavenge intracellular ROS generated by H₂O₂ or UVB was assessed using a spectrofluorometer after DCF-DA staining. *#Significantly different from cells treated with H₂O₂ or UVB alone ($p < 0.05$).

8-Isoprostane assay

Cells were seeded at a density of 1×10^5 cells/ml and incubated at 37°C for 24 h. Cells were treated with galangin (40 μM) and after 1 h, exposed to UVB (30 mJ/cm²), and incubated at 37°C for another 24 h. Lipid peroxidation was assayed by colorimetric determination of the amount of 8-isoprostane secreted into the culture medium by HaCaT keratinocytes (Beauchamp *et al.*, 2002). A commercial enzyme immunoassay (Cayman Chemical, Ann Arbor, MI, USA) was used according to the manufacturer's instructions.

Protein carbonylation assay

Cells were seeded at a density of 1×10^5 cells/ml and incubated at 37°C for 24 h. Cells were treated with galangin (40 μM) and after 1 h, exposed to UVB (30 mJ/cm²), and incubated at 37°C for another 24 h. The extent of protein carbonyl formation was determined using an OxiSelect™ Protein Carbonyl ELISA Kit from Cell Biolabs (San Diego, CA, USA) according to the manufacturer's instructions.

Nuclear staining with Hoechst 33342

Cells were seeded at a density of 1×10^5 cells/ml and incu-

bated at 37°C for 24 h. Cells were treated with galangin (40 μM) and after 1 h, exposed to UVB (30 mJ/cm²), and incubated at 37°C for another 24 h. The DNA-specific fluorescent dye Hoechst 33342 was added to each well and cells were incubated for 10 min at 37°C. Stained cells were visualized under a fluorescence microscope equipped with a CoolSNAP-Pro color digital camera. The degree of nuclear condensation was evaluated and apoptotic cells were counted.

Analysis of mitochondrial membrane potential ($\Delta\psi_m$)

JC-1 was added to each well and cells were incubated for 30 min at 37°C. Stained cells were washed with phosphate-buffered saline (PBS), coverslips were mounted onto microscope slides in mounting medium (DAKO, Carpinteria, CA, USA), and slides were examined using a confocal microscope. Microscopic images were collected using the Laser Scanning Microscope 5 PASCAL program (Carl Zeiss, Jena, Germany) (Cossarizza *et al.*, 1993). In addition, mitochondrial membrane potential was analyzed by flow cytometry. Cells were harvested, washed, suspended in PBS containing JC-1 (10 μg/ml), incubated for 30 min at 37°C, and analyzed using a flow cytometer (Troiano *et al.*, 2007).

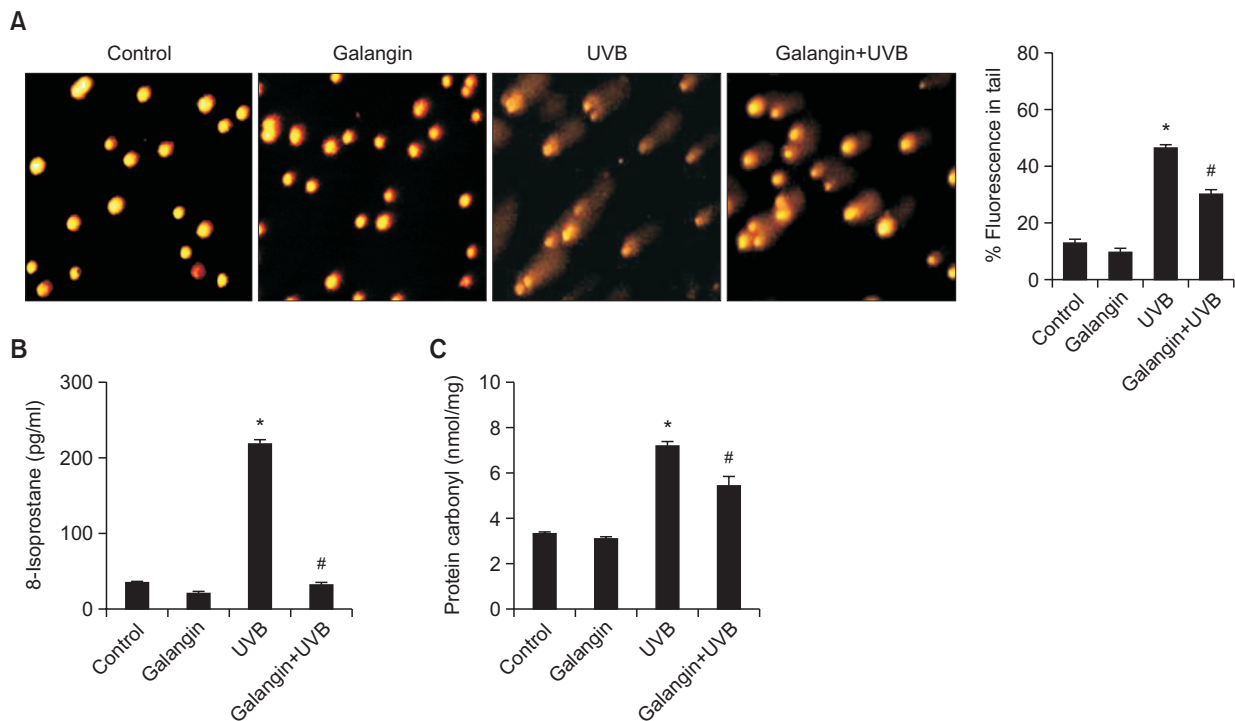


Fig. 3. Galangin attenuates UVB-induced macromolecule damage. Cells were treated with 40 μ M of galangin and exposed to UVB (30 mJ cm^{-2}) 1 hour later. After 24 hours, (A) The comet assay was performed to assess DNA damage. Representative images and the percentage of cellular fluorescence within comet tails are shown. *Significantly different from control cells ($p < 0.05$); #significantly different from UVB only-exposed cells ($p < 0.05$). (B) Lipid peroxidation was assessed by measuring the concentration of 8-isoprostane in the conditioned medium. *Significantly different from control cells ($p < 0.05$); #significantly different from UVB only-exposed cells ($p < 0.05$). (C) Protein oxidation was determined by measuring the amount of carbonyls formed. *Significantly different from control cells ($p < 0.05$); #significantly different from UVB only-exposed cells ($p < 0.05$).

Western blot analysis

Harvested cells were lysed by incubation on ice for 10 min in 150 μ l of lysis buffer containing 120 mM NaCl, 40 mM Tris (pH 8), and 0.1% NP 40. The resultant cell lysates were centrifuged at 13,000 rpm for 5 min. Supernatants were collected and protein concentrations were determined. Aliquots were boiled for 5 min and electrophoresed on 12% SDS-polyacrylamide gels. Protein blots of the gels were transferred onto nitrocellulose membranes. The membranes were incubated with the appropriate primary antibodies (1:1,000) followed by horseradish peroxidaseconjugated anti-IgG secondary antibodies (1:5,000) (Pierce, Rockford, IL, USA). Protein bands were detected using an enhanced chemiluminescence Western blotting detection kit (Amersham, Little Chalfont, Buckinghamshire, UK).

Statistical analysis

All measurements were performed in triplicate and all values are expressed as means \pm standard error. The results were subjected to an analysis of variance using Tukey's test to analyze differences between means. In each case, a p -value of < 0.05 was considered statistically significant.

RESULTS

Galangin attenuates UVB-induced ROS generation

The MTT assay showed that galangin was not toxic to

HaCaT cells at any concentration used. Following the treatment with each of the concentrations of galangin tested, cell viability was more than 96% of that of control cells (Fig. 2A). Galangin showed DPPH radical-scavenging activity at concentrations of 20-100 μ M compared to treatment with NAC (1 mM), a well-known antioxidant (Fig. 2B). Therefore, 40 μ M was selected as the optimal concentration of galangin for further experiments. To assess the ability of galangin (40 μ M) to scavenge hydroxyl radicals, ESR spectrometry was performed. In the Fenton reaction ($\text{Fe}^{2+} + \text{H}_2\text{O}_2 \rightarrow \text{Fe}^{3+} + \cdot\text{OH} + \text{OH}^-$), DMPO/ $\cdot\text{OH}$ adducts generated a signal of 2,915 in control cells and this was reduced to 1,803 in the presence of galangin (Fig. 2C). Next, the intracellular ROS scavenging activity of galangin was assessed using the DCF-DA fluorescent probe. Galangin scavenged 27% of ROS (versus 52% for NAC) in H_2O_2 -treated cells and 27% of ROS (versus 22% for NAC) in UVB-treated cells (Fig. 2D). Taken together, these results show that galangin efficiently scavenges ROS.

Galangin significantly attenuates UVB-induced damage of cellular macromolecules

We next investigated whether galangin can protect macromolecules, such as DNA, lipids, and proteins, from UVB-induced oxidative damage. First, UVB-induced DNA damage and the protective effects of galangin were studied using the comet assay. Representative microscopy images indicating the length of comet tails and the percentage of fluorescence in the tails are shown in Fig. 3A. UVB treatment increased

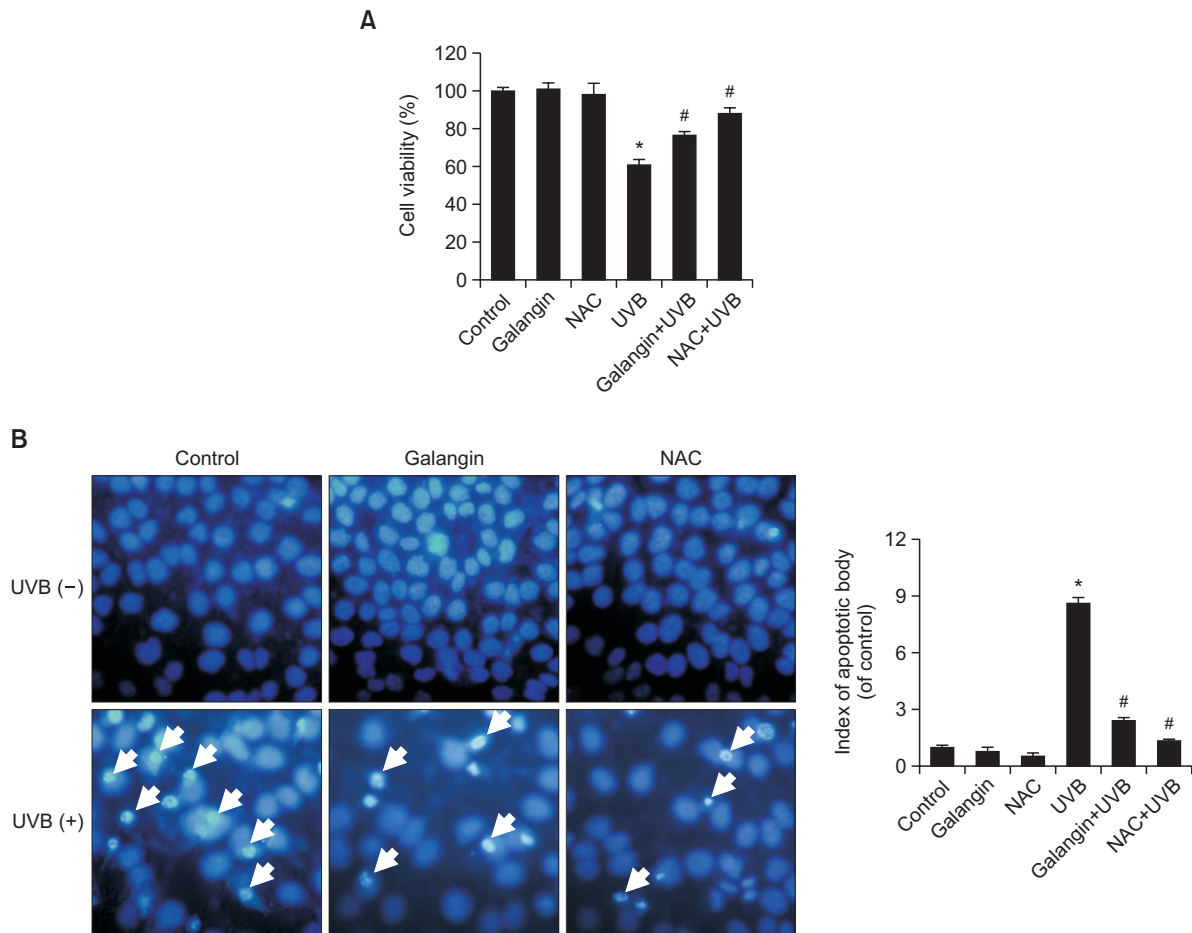


Fig. 4. Galangin increases the viability of UVB-irradiated cells and reduces apoptosis. Cells were pretreated with 40 μ M of galangin and exposed to UVB (30 mJ/cm^2) 1 hour later. (A) After 24 hours, the MTT assay was performed. *Significantly different from control cells ($p < 0.05$); #significantly different from UVB only-exposed cells ($p < 0.05$). (B) Cells were stained with Hoechst 33342 and observed by fluorescence microscopy. The ratio between apoptotic bodies (arrows) and total number of cells were determined within randomly selected 0.3 mm^2 area of each well. *Significantly different from control cells ($p < 0.05$); #significantly different from UVB only-exposed cells ($p < 0.05$). JC-1 staining was performed to observe the mitochondrial membrane potential via (C) confocal microscopy and (D) flow cytometry. (E) Western blot analysis was performed to detect the expression levels of cleaved caspase-9, -3, Bcl-2 and Bax proteins. (Continued on 170page)

the comet tail length and increased the percentage of fluorescence in the tail to 52%, indicating an increased level of DNA damage in keratinocytes, while galangin pretreatment significantly reduced this to 34%. UVB-induced cellular membrane damage was measured by detecting lipid peroxidation via colorimetric determination of the level of 8-isoprostane secreted into the culture medium by HaCaT cells. The concentration of 8-isoprostane was 219 pg/ml in the culture medium of UVB-exposed cells and 32 pg/ml in that of cells pretreated with galangin prior to UVB exposure (Fig. 3B). Finally, protein carbonylation was measured to assess the degree of protein damage. Oxidative stress usually modifies the amino acid side chains of proteins to carbonyl derivatives, which can be used to quantify protein damage caused by UVB-induced oxidative stress (Dalle-Donne, 2006). Protein carbonylation was higher in UVB-treated cells than in control cells, whereas this increase was significantly reduced in cells pretreated with galangin prior to UVB exposure (Fig. 3C). These findings confirm that galangin can protect cellular macromolecules from UVB-induced oxidative damage.

Galangin reduces UVB-induced apoptosis

The viability of HaCaT cells was assessed using the MTT assay (Fig. 4A). UVB exposure reduced the viability of keratinocytes to 60% in comparison to that of control cells. However, pretreatment with galangin increased cell viability to 76%, while pretreatment with the well-known antioxidant NAC increased cell viability to 88%. Next, cells were stained with Hoechst 33342 to visualize nuclear condensation and the formation of apoptotic bodies, which are characteristic of apoptosis. Normal nuclei were observed in control and galangin-treated cells, whereas notable nuclear condensation was found in UVB-exposed cells (Fig. 4B). The nuclear condensation was significantly lowered in galangin- and NAC-pretreated cells than in UVB only-exposed cells. Mitochondrial membrane permeability plays a crucial role in the mitochondria-mediated apoptosis pathway. To elucidate the mechanism underlying how galangin protects keratinocytes against UVB-induced apoptosis, changes in the mitochondrial membrane potential were assessed. UVB treatment strongly increased the level of green fluorescence caused by JC-1 monomers, indicat-

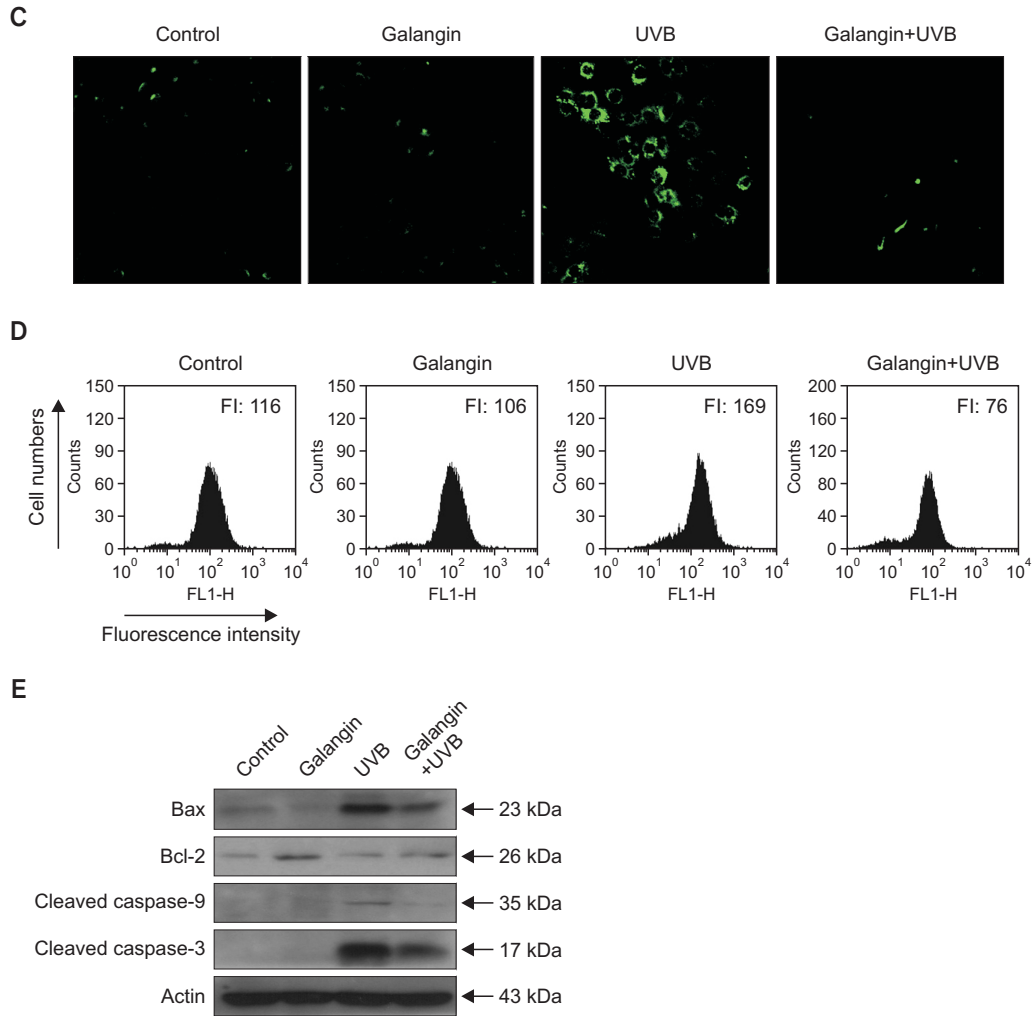


Fig. 4. Continued.

ing mitochondrial depolarization (Fig. 4C). However, galangin treatment prior to UVB exposure significantly reduced the intensity of green fluorescence, showing that galangin attenuated UVB-induced mitochondrial depolarization. To confirm these results, flow cytometric analysis of JC-1 was performed. As expected, the green fluorescence peak was increased in UVB-treated cells. However, galangin pretreatment notably reduced the peak of this fluorescence (Fig. 4D). Caspase-9, an initiator caspase, is cleaved to activate followed by the releasing cytochrome c from the depolarized mitochondria into the cytosol, cleaved caspase-9 propagates further activation of downstream apoptotic proteins which causes for cleavage of procaspase-3, activated caspase-3 triggers the apoptotic process (Adrain and Martin, 2001). To assess the caspase-9 and -3 expression levels, a western blot analysis was performed. UVB exposure resulted up-regulation of cleaved caspase-9 and -3 in HaCaT cells, interestingly galangin pretreatment significantly down-regulated the levels of cleaved caspase-9 and -3. The vulnerability of cells towards apoptosis is ultimately determined by the pro-apoptotic and anti-apoptotic proteins in the B-cell leukaemia/lymphoma-2 (Bcl-2) family (Basu and Haldar, 1998). Bcl-2 itself is consider as an anti-apoptosis pro-

tein while Bcl-2-associated x protein (Bax) is known as a pro-apoptotic protein in the Bcl-2 protein family (Wei *et al.*, 2001) Therefore we next investigated the level of Bcl-2 and Bax proteins after galangin treatment (Fig. 4E). Galangin treatment prior to UVB exposure resulted up-regulation of Bcl-2 proteins whereas down- regulation of Bax proteins. These results indicate that galangin inhibits apoptotic process *via* establishing mitochondrial membrane polarity and regulating apoptotic proteins.

DISCUSSION

Chronic solar UVB exposure, also known as photoaging, is the most well understood skin aging mechanism (Nichols and Katiyar, 2010; Svobodova and Vostalova, 2010). Although the skin possesses a complex enzymatic and non-enzymatic defense system to protect it from adverse effects, prolonged UVB exposure can overwhelm this system. Solar UVB radiation has become one of the most common carcinogens in the environment and excessive UVB exposure leads to skin cancer. Meanwhile, due to the depletion of the stratospheric ozone

layer and climate change, the earth receives elevated levels of UV radiation (De Grujil *et al.*, 2003; Diffey, 2004). UVB damages DNA directly via the formation of pyrimidine dimers and indirectly via ROS generation (Ichihashi *et al.*, 2003). DNA is the chief information molecule in the cell; therefore, nuclear DNA must exist for the entire life time of the cell. DNA damage can critically affect cellular functions. ROS attack not only nucleic acids but also proteins and lipids, thereby interrupting cellular metabolism. In this regard, prevention of the adverse effects caused by exposure to harmful UV radiation has become a popular theme in the cosmetic and pharmaceutical industries. Phytochemicals are well-known for their abilities to protect the skin against harmful UV radiation. Galangin is a flavone whose antioxidant ability depends on the donation of hydrogen atoms to free radicals (Sim *et al.*, 2007). In this study, galangin scavenged DPPH radicals and hydroxyl radicals (Fig. 2B, C). DCF-DA staining revealed that intracellular ROS generated via H₂O₂ and UVB (Fig. 2D) were removed by galangin. Flavonoid is known to possess antioxidant activity by donation of a hydrogen to radicals, depending on the substitution pattern of hydroxyl groups. Hydroxylated position is related with stability of the resulting phenoxyl radical by hydrogen donation or electron delocalization. The presence of 3-OH and 5-OH and C2-C3 double bond conjugated with the 4-oxo function increase antioxidant activity (Rice-Evans *et al.*, 1996; Jung *et al.*, 2005). In general, O-dihydroxyl groups known as catechol structure in B ring increases radical scavenging activity, however it is reported that absence of catechol structure in the B ring like galangin is not always related with low antioxidant activity by compensating lack of catechol structure through combination C2-C3 double bond with the 3-OH and 4-keto group (Amic *et al.*, 2003), demonstrating antioxidant effect as shown in our results.

UVB-induced ROS cause oxidative damage to DNA, lipids, and proteins in keratinocytes. Formation of CPDs and 6-4 photoproducts are considered as the main types of lesions caused by UVB radiation in DNA, and these adducts cause DNA strand breaks and induction of mutations (You *et al.*, 2000). ROS especially hydroxyl radicals abstract allylic hydrogen forming carbon centered lipid radicals which will rapidly react with oxygen to form lipid peroxy radicals, interestingly these peroxy radicals are capable of abstract hydrogens from another lipid molecules generating lipid radicals initiating chain reactions (Ayala *et al.*, 2014). ROS attack proteins and cause reversible and/or irreversible modifications, such as protein carbonylation, formation of adducts with lipid peroxidation products and protein protein cross-linking (Perluigi *et al.*, 2010). Formation of protein carbonyls could be either by oxidative cleavage of proteins or by direct oxidation of lysine, arginine, proline, and threonine residues (Sander *et al.*, 2002). The comet assay showed that galangin significantly suppressed DNA strand breaks induced by UVB (Fig. 3A). Detection of 8-isoprostane released by cells into the extracellular environment is an accurate means of assaying lipid peroxidation. Galangin-pretreated cells showed notably lower levels of 8-isoprostane in the culture medium than UVB alone-treated cells (Fig. 3B), suggesting that galangin attenuated UVB-induced lipid peroxidation in skin cells. Proteins are oxidized by ROS and formation of protein carbonyls is a hallmark of oxidative stress. Our results illustrated that the level of carbonyls was significantly attenuated in cells pretreated with galangin prior to UVB exposure (Fig. 3C).

At elevated concentration of ROS overwhelms the antioxidant defense system and interrupts cellular metabolism. In response to DNA damage, cells can undergo apoptosis (Barzilai and Yamamoto, 2004).

Recent studies show that UVB induced cell death is basically governed through apoptotic pathway (Ji *et al.*, 2012). In addition to DNA damage and cell cycle arrest by UVB induced oxidative stress, following the membrane damage via lipid peroxidation and protein carbonylation, ROS weaken the inner cellular structures especially mitochondrial membranes which causes to release cytochrome c into the cytosol and activates apoptotic proteins (Kulms *et al.*, 2002). Galangin pretreatment significantly increased the viability of UVB-exposed cells (Fig. 4A). Hoechst 33342 staining revealed that galangin pretreatment effectively reduced the formation of apoptotic bodies and DNA condensation induced by UVB exposure (Fig. 4B). Furthermore, we investigated the impact of galangin pretreatment on mitochondria-mediated apoptosis via JC-1 staining. Confocal microscopy (Fig. 4C) and flow cytometric (Fig. 4D) analyses confirmed that galangin restored mitochondrial membrane polarization. These data showed that galangin reduced apoptosis of UVB-radiated cells by establishing mitochondrial polarization. Pro-apoptotic protein Bax and anti-apoptotic protein Bcl-2 regulate the release of cytochrome c into the cytosol (Atan *et al.*, 1999). Caspases are synthesized in their inactive forms in the cells and activated in respond to apoptotic signals (Thornberry and Lazebnik, 1998). Released cytochrome c into the cytosol binds to apoptotic protease activating factor 1, which then activates caspase-9 by cleaving procaspase-9 (Bossy-Wetzel *et al.*, 1998). Cleaved caspase-9 activates caspase-3 and which then triggers downstream caspase cascades to execute apoptosis (Soengas *et al.*, 1999). Our results elucidated that galangin suppressed the expression of Bax, cleaved caspase-9 and -3 while increased the expression of Bcl-2 protein level (Fig. 4E).

In summary, our results confirmed that galangin possesses antioxidant properties and it interferes intrinsic pathway of apoptosis via down regulating key apoptotic proteins, which it protected HaCaT cells from UVB-induced oxidative damage.

ACKNOWLEDGMENTS

This work was supported by a National Research Foundation of Korea Grant funded by the Korean Government (MEST) (NRF-C1ABA001-2011-0021037).

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