Chemical Constituents from Leaves of *Hydrangea serrata* and Their Anti-photoaging Effects on UVB-Irradiated Human Fibroblasts

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Received September 25, 2018; accepted November 28, 2018.

**Abstract**

*Hydrangea serrata* (Thunb.) Ser. (Hydrangeaceae) leaves have been used as herbal teas in Korea and Japan. The objective of this study was to identify anti-photoaging compounds in aqueous EtOH extract prepared from leaves of *H. serrata* and their effects on UVB-irradiated Hs68 human foreskin fibroblasts. Phytochemical study on *H. serrata* leaves led to the isolation and characterization of ten compounds: hydrangenol, thunberginol A, thunberginol C, hydrangenoside A, hydrangenoside C, cudrabibenzyl A, 2,3,4-trihydroxystilbene, thunberginol F, quercetin 3-O-β-D-xylopyranosyl (1-2)-β-D-galactopyranosyl (1-2)-β-D-o-galactopranoside, quercetin 3-O-β-D-xylopyranosyl (1-2)-β-D-o-glucopyranoside, cudrabibenzyl A, 2,3,4-trihydroxyxylotilbene, quercetin 3-O-β-D-xylopyranosyl (1-2)-β-D-o-galactopranoside, quercetin 3-O-β-D-xylopyranosyl (1-2)-β-D-o-glucopyranoside were firstly isolated from *H. serrata*. We estimated the effects of 10 compounds on cell viability and production of pro-collagen Type I, matrix metalloproteinase (MMP)-1, and hyaluronic acid (HA) after UVB irradiation. Of these compounds, hydrangenol showed potent preventive activities against reduced cell viability and degradation of pro-collagen Type I in UVB-irradiated Hs68 fibroblasts. Hydrangenol had outstanding inductive activities on HA production. It suppressed mRNA expression levels of MMP-1, MMP-3, hyaluronidase (HYAL)-1, HYAL-2, cyclooxygenase-2 (COX-2), interleukin (IL)-6, IL-8, and IL-1β in UVB-irradiated Hs68 fibroblasts. When Hs68 fibroblasts were exposed to hydrangenol after UVB irradiation, UVB-induced reactive oxygen species (ROS) production was suppressed. Hydrangenol also inhibited the activation of activator protein-1 (AP-1) and signal transduction and activation of transcription 1 (STAT-1) by downregulating phosphorylation of p38 and extracellular signal-regulated kinase (ERK). Our data indicate that hydrangenol isolated from *H. serrata* leaves has potential protective effects on UVB-induced skin photoaging.

**Key words** *Hydrangea serrata*; photoaging; matrix metalloproteinase (MMP); procollagen type 1; skin fibroblast.

**INTRODUCTION**

Human skin comprises the epidermis, dermis, and hypodermis. The epidermal extracellular matrix is a basement membrane composed of keratinocytes that produce keratins. The dermis is composed of fibrous structural collagen-rich extracellular matrix (ECM) which is essentially responsible for the skin’s tensile strength and mechanical properties, together with elastin and hyaluronic acid. Upon exposure to solar UV, especially UVB, degradation of ECM is being processed through activation of matrix metalloproteinases (MMPs) with decreased pro-collagen synthesis, resulting in wrinkle formation and sagging skin (skin photoaging). A great number of plant extracts and phytocompounds including polyphenols (flavonoids and non-flavonoids) and non-phenolic derivatives possess anti-oxidant and anti-inflammatory properties. They have received growing attention as potential dietary supplements for anti-aging interventions. Natural sources of photoprotective phytocompounds including green tea, coffee, berries, and pomegranate have been reported. Hydrangea (Hydrangeaceae) leaves have been consumed as medicine and tea in far-east Asian countries such as Korea, China, and Japan. Some researchers have reported that extracts of hydrangea leaves possess anti-inflammatory, hepato-protective, and antimarial activities. To date, a number of bioactive compounds such as dihydroisocoumarins, secoiridoids, and stilbenes (hydrangenol, phyllodulcin, macrophylline, and their glucosides) have been isolated from hydrangea plants. Phyllodulcin has been identified as the compound responsible for the sweet taste of its infusion. *Hydrangea serrata* (Thunb.) Ser. is native to Korean mountains known as “San-soogook, Mountain Hydrangea, or Tea of Heaven.” *H. serrata* is small deciduous shrubs similar to *H. macrophylla* and usually with lace cap flowers that are flat with smaller florets in the center surrounded by larger florets. The blooms are either blue or pink depending on the acidity of the soil. The flowers will be blue in higher acidic soils and lilac in alkaline soils. Growing to 1–2 m tall and broad, it features dark green, serrated (toothed), and ovate leaves to 15 cm long. Its leaves are used to make herbal teas. However, few studies have determined the chemical compositions of *H. serrata* and their anti-photoaging activities. Therefore, the objective of this study was to isolate constituents from *H. serrata* and determine their anti-photoaging properties in UVB-irradiated skin fibroblast.
MATERIALS AND METHODS

Isolation of Compounds from Ethanol Extracts from *H. serrata* Leaves  The leaves of *Hydrangea serrata* (THUNB.) Sir. (20 g) were extracted twice with 50 or 70% ethanol (375 mL) at 50°C for 3 h to give dried extract residue 50% ethanol extracts from *H. serrata* (EHS50, 5.8 g, 29%) and 70% ethanol extracts from *H. serrata* (EHS70, 5.6 g, 28%), respectively. EHS50 was fractionated on Diaion HP-20 column, starting with 30% MeOH followed by 50% (50M), 70% (70M), 100% MeOH (100M) and CH<sub>3</sub>Cl<sub>2</sub>–MeOH (CM), 1:1 as an eluent. Fraction 70E-100M was subjected to sephadex LH-20 CC eluted with MeOH to afford seven fractions (70E100M1–70E100M7). Subfraction 70E100M4 was further purified by toyporeal HW-40 eluted with MeOH to give hydrgenanol (31.1 mg). Fraction 70E-70M was subjected to silica gel CC using CH<sub>3</sub>Cl<sub>2</sub>–MeOH (7:1) mixtures to give nine fractions (70E70M1–70E70M9). Subfraction 70E70M2 was further purified by preparative HPLC using a gradient mobile phase system CH<sub>3</sub>CN–H<sub>2</sub>O (30.70:70.30, v/v) at flow rate 4 mL/min to give hydrgenanol A (24.8 mg, 12.5 min) and hydrgenanol C (2.7 mg, 28 min). Preparative HPLC was performed on a Waters pump (model 1525) with photodiode array detector (Waters model 2996) using a Luna C18 (10 mm × 250 mm, Phenomenex) column. Fraction 70E-50M was separated by silica gel eluted with CH<sub>3</sub>Cl<sub>2</sub>–MeOH–H<sub>2</sub>O (4:12:0.1) to give ten fractions (70E50M1–70E50M10). Subfraction 70E50M9 (38.7 mg) was purified by preparative HPLC using 32% MeOH to afford cudrabibenzyl A (4.8 mg, 3.5 min), quercetin-3-O-β-D-glucopyranoside (2-β-D-galactopyanoside (4.4 mg, 21 min) and quercetin-3-O-β-D glucopyranoside (4.4 mg, 21 min). EHS50 was fractionated on Diaion HP-20 column, starting with 30% MeOH followed by 50% (50M), 70% (70M), 100% MeOH (100M) and CH<sub>3</sub>Cl<sub>2</sub>–MeOH (CM), 1:1 as an eluent. Fraction 50E-100M (731.1 mg) was subjected to sephadex LH-20 CC eluted with MeOH to afford five fractions (50E100M1–50E100M5). Subfraction 50E100M5 was further purified by preparative HPLC using a gradient mobile phase system CH<sub>3</sub>CN–H<sub>2</sub>O (20:70→75:25, v/v) as described above to give hydrgenanol (29.1 mg, 24 min), thunberginol A (2.5 mg, 22 min), thunberginol C (18.1 mg, 17 min), 2,3,4,5,6-pentahydroxystilbene (1.5 mg, 8.2 min) and thunberginol F (4.3 mg, 15.3 min).

**Cell Culture and UVB Exposure** Human skin fibroblast Hs68 cells were obtained from the American Type Culture Collection (ATCC, Manassas, VA, U.S.A.). Cells were grown in Dulbecco’s modified Eagle’s medium (DMEM) containing 10% fetal bovine serum (FBS) and penicillin–streptomycin, 100 U/mL penicillin and 100 µg/mL streptomycin in a humidified atmosphere of 5% CO<sub>2</sub>. Hs68 cells were plated at 1 × 10<sup>5</sup> cells/mL in culture vessels and incubated overnight. Cells were then washed with phosphate-buffered saline (PBS) and irradiated with UVB at 15 mJ/cm<sup>2</sup> using UVP Crosslinker (Analytik Jena AG, Jena, Germany). Cells were washed with PBS and further incubated in fresh DMEM in the presence or absence of the indicated concentration of compounds. Isolated compounds were dissolved in dimethyl sulfoxide (DMSO).

**Determination of Cell Viability** After UVB irradiation, cells were treated with indicated concentration of compounds for 24 h. Epigallocatechin gallate (EGCG, 10 µM) was used as a positive control. Fifty micrometers of 3-(4,5-dimethylthiazol-2-yl)-2,5-diphenyltetrazolium bromide (MTT) was then added to the wells followed by 4 h. Formazan crystals were completely dissolved by DMSO and the optical density at 595 nm was measured using a microplate reader (Molecular Devices, U.S.A.).

**Enzyme-Linked Immunosorbent Assays (ELISAs) for Pro-collagen Type I, Matrix Metalloproteinase (MMP)-1, Hyaluronic Acid (HA)** Following UVB irradiation, cells were treated indicated the concentration of compounds for 24 h. Epigallocatechin gallate (10 µM) was used as a positive control for type I procollagen and MMP-1. The level of pro-collagen type I, MMP-1, hyaluronic acid in culture media was determined using specific ELISA kit (Procollagen Type I C-Peptide ELISA Kit, TaKaRa, Shiga, Japan; human MMP-1 ELISA kit, Abcam, Cambridge, UK; HA ELISA kit, R&D Systems, MN, U.S.A.).

**RNA Preparation and Real-Time Quantitative (q)RT-PCR** Total cellular RNA from compound-treated cells was isolated using Easy Blue<sup>®</sup> kits (Intron Biotechnology, Seoul, Korea) and cDNA was prepared using random oligonucleotide primers (Promega, Madison, WI, U.S.A.) and TOPO<sup>TM</sup> cDNA synthesis kit (Enzymics, Daejeon, Korea). PCR amplification was performed using the incorporation of SYBR green (TaKaRa, Shiga, Japan) and specific primers. Specific gene primer was listed in Supplementary Table 1. Steady-state mRNA levels of MMP-1, MMP-3, HA synthase (HAS) 2, hyaluronidase (HYAL)-1, HYAL-2, cyclooxygenase-2 (COX-2), interleukin (IL)-6, IL-8, IL-1β, and glyceraldehyde-3-phosphate dehydrogenase (GAPDH) were determined by real-time PCR using the Light Cycler1.5 (TaKaRa, Shiga, Japan). Dissociation curve analysis for MMPs and β-actin oligos showed a single peak for each. Mean Ct values of genes of interest were calculated from triplicate measurements and normalized versus the mean Ct of GAPDH.

**Western Blot Analysis** Total proteins from compound-treated cells were isolated using PRO-PREP (Intron Biotechnology, Seoul, Korea) and incubated at 4°C for 30 min. Cell debris was removed by microcentrifugation, followed by quick freezing of the supernatants. Nuclear extracts from compound-treated cells were prepared using a hypotonic buffer and high salt buffer. Cell pellets were resuspended in hypotonic buffer (10 mM N-(-2-hydroxyethyl)piperazine-N’-2-ethanesulfonic acid (HEPES), pH 7.9, 1.5 mM MgCl<sub>2</sub>, 10 mM KCl, 0.2 mM phenylmethylsulfonyl fluoride (PMSF), 0.5 mM dithiothreitol (DTT), 10 µg/mL aprotinin) and incubated on ice for 15 min. Cells were then lysed by adding 0.1% Nonidet P-40 and vortexed vigorously for 10 s. Nuclei were pelleted by centrifugation at 12000 × g for 1 min at 4°C and resuspended in high salt buffer (20 mM HEPES, pH 7.9, 25% glycerol, 400 mM KCl, 1.5 mM MgCl<sub>2</sub>, 0.2 mM ethylenediaminetetraacetic acid (EDTA), 0.5 mM DTT, 1 mM NaF, 1 mM sodium orthovanadate). The protein concentration of the lysates was determined using the Bio-Rad protein assay reagent.

**Determination of Reactive Oxygen Species (ROS) Generation** The generation of ROS was estimated using the redox-sensitive fluorescent dye fluorescent probes 2′,7′-dichlorodihydrofluorescein diacetate (H<sub>2</sub>DCFDA). Following UVB irradiation, cells were treated with hydrgenanol for indicated time, and further incubated with 20 µM H<sub>2</sub>DCFDA for 30 min.
Then, cells were resuspended in PBS and the fluorescence signal analyzed using flow cytometry (Beckman Coulter Inc., Brea, CA, U.S.A.).

**Statistical Analysis** Results are expressed as the mean ± standard deviation (S.D.) of triplicate experiments with similar patterns. The band density was estimated using Quantity One® program (Bio-Rad Laboratories, CA, U.S.A.). Statistically significant values were compared using ANOVA and Dunnett’s post-hoc test, and *p*-values of less than 0.05 were considered statistically significant.

![Fig. 1. Structures of Ten Compounds Isolated from the Leaves of Hydrangea serrata](image)

**RESULTS AND DISCUSSION**

Chemical investigation of aqueous EtOH extracts of *H. serrata* leaves resulted in the isolation and identification of ten compounds; dihydroisocoumarins (hydrangenol, thunberginol A, thunberginol C), secoiridoid glucosides (hydrangenoside A and hydrangenoside C), bibenzyl derivatives (cudrabibenzyl A and 2,3,4′-trihydroxystilbene), phthalide (thunberginol F) and flavonoid glycosides (quercetin 3-O-β-D-xylopyranosyl (1-2)-β-D-galactopyranoside, quercetin 3-O-β-D-xylopyranosyl (1-2)-β-D-glucopyranoside (quercetin-xylo-gluco)) (Fig. 1, Supplementary data 1). It is the first time that compounds cudrabibenzyl A, 2,3,4′-trihydroxystilbene, quercetin-xylo-galato, and quercetin-xylo-gluro were isolated from leaves of *H. serrata*. Hydrangenol was determined as the main ingredient using an HPLC/UV system with an analytical C18 column, and its content in 50 and 70% ethanol extracts was found to be 1.14 and 1.02% from HPLC profiles, respectively, while other isolates could not be measured due to low content and overlap with the others. Also, the content of hydrangenol in water extract (60°C) was found to be 0.08% by our extraction conditions (Supplementary data 1-S45). To determine whether isolated ten compounds might possess anti-

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Then, cells were resuspended in PBS and the fluorescence signal analyzed using flow cytometry (Beckman Coulter Inc., Brea, CA, U.S.A.).
photoaging properties, we assessed their effects on cell viability and productions of pro-collagen type I, MMP-1, and HA in UVB-irradiated Hs68 skin fibroblasts. EGCG (10 μM) was used as a positive control for cell viability, type I procollagen, and MMP-1.26

Cytotoxic protectors of ten compounds were evaluated using MTT assay. Hs68 cells were exposed to UVB (15 mJ/cm²) and then treated with ten compounds at 1 μM for 24h. Cell viability was significantly decreased (88.77%) after UVB-irradiation compared to that of control cells (100%). The eight compounds except hydrangenoside A and hydrangenoside C significantly recovered the reduction in cell viability induced by UVB irradiation (Table 1).

Accumulating data suggest that photoaging skin is characterized by a reduction in the deposition of collagen fibers which may be resulted from increased collagen degradation.27 To evaluate the effects of ten compounds on collagen production in UVB-irradiated Hs68 fibroblasts, UVB-irradiated cells were treated with ten compounds for 24h and type I procollagen production in cell supernatant was measured by ELISA. UVB exposure decreased pro-collagen type I production by 36.62%, while eight compounds except cudrabibenzy A and quercetin-xylo-gluco recovered UVB-induced reduction of pro-collagen type I by 22.26 to 42.67% (Table 1).

MMP-1 is a major protein-degrading enzyme in human skin that can fragment native fibrillar collagens, predominantly type I and III collagens.18 The fragments then become unfolded and degraded by MMP-2, -9, and -3.19 Therefore, MMPs are key coordinators for the formation of wrinkles in the photoaging process.20 Interestingly, all compounds had potent effects on MMP-1 production in UVB-irradiated Hs68 fibroblasts (Table 2). Compounds except quercetin-xylo-gluco inhibited UVB-induced MMP-1 production by more than 50% at 1 μM.

HA, another key component of dermal ECM, controls skin functions such as skin hydration, turgidity, elasticity, and diffusion of metabolites and nutrients.21,22 HA is synthesized by three HAS and catabolized by several HYAL enzymes.23 Loss of HA has been associated with skin aging and UVB exposure causes loss of HA because of down-regulated HAS expression.23,24 We evaluated the effects of isolated compounds on the UVB-induced loss of HA by ELISA. UVB exposure decreased HA production by 41.68%, while hydrangenol, hydrangenoside A, hydrangenoside C, and quercetin-xylo-galato significantly recovered UVB-induced HA loss. Of them, hydrangenol was the most potent inducer of HA (Table 2).

We further examined the protective effect of hydrangenol on UVB-irradiated Hs68 fibroblasts and the underlying molecular mechanism. To elucidate the mechanism of hydrangenol for pro-collagen type I recovery, we determined effects of hydrangenol (1.5, 3, or 6 μM) on mRNA expression of collagen type I chain (COL1A1), a major component of collagen type I, by qRT-PCR. The mRNA expression of COL1A1 was significantly reduced by UVB irradiation, but hydrangenol treatment failed to recover such reduction (data not shown). Therefore, increased production of pro-collagen type I by hydrangenol might be related to collagen degradation decreases

### Table 1. Effects of Compounds Isolated from Hydrangea serrata on Cell Viability and Pro-collagen Type I Productions in UVB-Irradiated Hs68 Cells

| Compounds (1 μM) | Cell viability (%) | Pro-collagen type I (ng/mL) |
|------------------|--------------------|---------------------------|
| Control          | 100.00 ± 1.19      | 725.99 ± 46.13            |
| UV               | 88.47 ± 4.86       | 460.13 ± 22.02            |
| Hydrangenol      | 98.6 ± 2.57        | 519.32 ± 25.60            |
| Thunberginol A   | 97.07 ± 0.98       | 562.46 ± 28.18            |
| Thunberginol C   | 105.92 ± 2.52      | 575.53 ± 44.90            |
| Hydrangenoside A | 88.92 ± 0.20       | 573.56 ± 21.46            |
| Hydrangenoside C | 93.12 ± 0.16       | 557.77 ± 10.66            |
| Cudrabibenzy A   | 94.69 ± 5.20       | 495.98 ± 49.49            |
| 2,3,4′-Trihydroxystilbene | 97.17 ± 3.54 | 553.03 ± 1.30***          |
| Thunberginol F   | 104.07 ± 1.74      | 572.44 ± 80.84*           |
| Quercetin-xylo-galato | 95.35 ± 2.59 | 476.87 ± 48.66            |
| Quercetin-xylo-gluco | 104.41 ± 1.58** | 528.25 ± 33.63*           |
| EGCG (10 μM)     | 93.45 ± 2.50**     | 514.57 ± 10.66*           |

* Numbers in marks of parenthesis indicated the percent (%).

### Table 2. Effects of Compounds Isolated from Hydrangea serrata on MMP-1 and Hyaluronic Acid Productions in UVB-Irradiated Hs68 Cells

| Compounds (1 μM) | MMP-1 (ng/mL) | Hyaluronic acid (ng/mL) |
|------------------|---------------|------------------------|
| Control          | 1.45 ± 0.52   | 883.85 ± 155.53        |
| UV               | 5.70 ± 0.39   | 515.22 ± 92.84         |
| Hydrangenol      | 2.51 ± 0.03***| 1045.38 ± 118.14***    |
| Thunberginol A   | 2.74 ± 0.04***| 629.74 ± 12.10         |
| Thunberginol C   | 1.58 ± 0.05***| 711.87 ± 149.17        |
| Hydrangenoside A | 1.84 ± 0.01***| 794.64 ± 156.29*       |
| Hydrangenoside C | 2.40 ± 0.03***| 893.44 ± 174.16***     |
| Cudrabibenzy A   | 2.51 ± 0.01***| 485.14 ± 76.18         |
| 2,3,4′-Trihydroxystilbene | 1.87 ± 0.01* | 551.42 ± 39.85         |
| Thunberginol F   | 1.47 ± 0.07***| 492.61 ± 48.82         |
| Quercetin-xylo-galato | 3.24 ± 0.02**| 834.43 ± 133.09**      |
| Quercetin-xylo-gluco | 1.73 ± 0.01***| 714.56 ± 171.52        |
| EGCG (10 μM)     | 0.14 ± 0.01***| (9.66 ± 0.69)***       |

* Numbers in marks of parenthesis indicated the percent (%).
rather than collagen synthesis upregulation. It has been reported that collagen is fragmented by MMP-1 from a single site in its central triple. MMP-3 not only degrades ECM components, but also activates pro MMP-1. 18,19 Based on the significant inhibitory effect of hydrangenol on UVB-induced MMP-1 production, we examined effects of hydrangenol at different concentrations (1.5, 3, or 6 \( \mu \)M) on MMP-1 and MMP-3 mRNA expression in UVB-irradiated Hs68 fibroblasts. Hydrangenol concentration-dependently reduced UVB-induced MMP-1 and MMP-3 mRNA expression (Fig. 2A). These results suggest that hydrangenol can potently recover UVB-induced collagen disruption by down-regulating expression of MMPs.

We investigated whether the recovery effect of hydrangenol on HA production might be related to HA metabolism by determining mRNA expression levels of HAS and HYALs. Results showed that mRNA expression levels of HAS-2 were significantly reduced in UVB-irradiated cells while those of HYAL-1 and HYAL-2 were significantly increased (Fig. 2B). The restoring effect of hydrangenol (1.5, 3, or 6 \( \mu \)M) on UVB-induced increase of HYAL-1 and HYAL-2 mRNA expression was superior to its effect on HAS-2 mRNA expression. Therefore, hydrangenol might increase HA production by suppressing the degradation of HA in UVB-irradiated cells.

UVB irradiation leads to skin inflammatory responses by increasing pro-inflammatory cytokines, consequently accelerating MMPs expression and collagen decomposition. 20-22 UVB-induced elevation of COX-2 is closely linked to inflammation and cancer in skin.26,27 Thus, preventing inflammation is another important strategy to protect skin against photoaging. To determine the anti-inflammatory effect of hydrangenol on skin fibroblasts, inhibitory effects of hydrangenol on UVB-induced COX-2, IL-6, IL-8, and IL-1\( \beta \) mRNA expression levels were investigated using qRT-PCR. As shown in Figs. 2C, D, hydrangenol (1.5, 3, or 6 \( \mu \)M) significantly suppressed these inflammatory mediators, suggesting that hydrangenol might have anti-inflammatory potential in UVB-irradiated Hs68 fibroblasts.

UVB-induced ROS is an initial pathogenic molecule that causes oxidative stress, skin aging-related gene expression, and subsequent damage to the skin.23-25 After exposing the cells to UVB, we evaluated the anti-oxidant properties of hydrangenol in 0.5, 1, 2, 4, 12, or 24 h by cellular ROS staining with H_2DCFDA. As shown in Fig. 3, UVB-induced ROS production increased in a typical cyclic pattern in Hs68 fibroblasts. However, hydrangenol (6 \( \mu \)M) significantly reduced ROS production in the cycle in which ROS was increased (0.5, 2, 4, 12, or 24h). Our data indicate that hydrangenol plays a defensive role against UVB-induced oxidative stress.

UVB-induced ROS activates the signal transduction pathways such as mitogen-activated protein kinase (MAPK), activator protein 1 (AP-1), nuclear factor-kappa B (NF-\( \kappa \)B), and janus kinase (JAK)/signal transduction and activation of...
transcription 1 (STAT1). AP-1 is composed of homodimers of Jun proteins or heterodimers of Jun-Fos proteins. Its activity is regulated by both expression and phosphorylation of AP-1 proteins. To determine whether hydrangenol could suppress the UVB-induced AP-1 activation, we determined the accumulation of c-Fos and c-Jun in a nuclear fraction from hydrangenol (1.5, 3, or 6µM)-treated Hs68 fibroblasts. Western blot analysis showed that hydrangenol significantly decreased UVB-induced nuclear expression of c-Fos and c-Jun, demonstrating hydrangenol could inhibit UVB-induced AP-1 activation (Fig. 4A). UVB exposure also leads to phosphorylation of STAT1 at Ser727 which in turn enhances its transcriptional activity. As shown in Fig. 4A, UVB-induced phosphorylation of STAT1 (Ser727) was diminished by hydrangenol, although total expression of STAT1 was not changed by either UVB or UVB plus hydrangenol. These results suggest that the inhibitory activity of hydrangenol on UVB-induced aging-related genes such as MMPs, HYALs, and pro-inflammatory mediators might be due to inactivation of AP-1 and STAT1 signaling.

MAPKs including p38, c-Jun N-terminal kinases (JNK), and extracellular signal-regulated kinase (ERK) have been accepted as the targets for chemoprevention of skin disease, given that they are pivotal regulation factors for UVB-induced skin aging by activation several skin aging-related transcription factors. Therefore, we assessed the phosphorylation rate of p38, JNK, and ERK by Western blot analysis, and the results showed that hydrangenol decreased UVB-induced p38 and ERK phosphorylation. No changes in phosphorylation of JNK and total expression of p38, JNK, and ERK were ob-

Fig. 3. Hydrangenol Suppressed Intracellular ROS in UVB-Irradiated Hs68 Cells

Following UVB irradiation, cells were treated with hydrangenol for the indicated time and further incubated with H$_2$DCFDA for 30min. The fluorescence signal was analyzed using flow cytometry. Data are presented as means ± S.D.s of three independent experiments. *p < 0.05 vs. the control cells; **p < 0.01 vs. UVB-exposed cells.

Fig. 4. Effects of Hydrangenol on UVB-Induced Activation of AP-1, STAT1, and MAPKs in Hs68 Cells

(A and B) Nuclear proteins (N) or whole proteins were isolated from cells treated with hydrangenol for 4h after UVB exposure. Proteins were detected using specific antibodies for p-cFos, cFos, p-cJun, cJun, p-p38, p38, p-ERK, ERK, p-JNK, and JNK. Histone H3 or β-actin were used as internal controls for nuclear or whole extraction, respectively. *p < 0.05 vs. the control cells; **p < 0.01 vs. UVB-exposed cells.
served after treatment with hydrangenol (Fig. 4B). It has been well-described that p38 and ERK can increase the transactivation and stabilization of c-Fos and c-Jun by inducing their expression and phosphorylation and can consequently result in the AP-1 activation. Moreover, inhibition of p38 or ERK by specific inhibitors or dominant negative mutant can prevent UVB-induced phosphorylation of STAT1, suggesting that p38 and ERK may play a potent role in STAT1-mediated skin inflammatory responses. Taken together, our data suggest that hydrangenol can suppress UVB-induced AP-1 and STAT1 activation via interfering with p38 and ERK-regulated pathway.

In the present study, ten compounds were isolated from *H. serrata* leaves and their protective effects on UVB-induced cell viability, pro-collagen type I, MMP1, and HA were evaluated in Hs68 fibroblasts. Hydrangenol, thunberginol A, thunberginol C, 2,3,4-trihydroxystilbene, thunberginol F, quercetin-xylo-glucos showed preventive activities against the reduced cell viability and the decreased production of pro-collagen type I and MMP-1. Hydrangenol restored the production of pro-collagen type I and HA by inhibiting expression levels one for HA induction. Hydrangenol was the most potent to reduce cell viability and the decreased production of MMP-1 and HA induction. Hydrangenol was the most potent to reduce cell viability and the decreased production of pro-collagen type I and MMP-1.

Acknowledgments This work was supported by Korea Institute of Planning and Evaluation for Technology in Food, Agriculture, Forestry and Fisheries (IPET) through High Value-added Food Technology Development Program, funded by Ministry of Agriculture, Food and Rural Affairs (MAFRA) (117070-03).

Conflict of Interest The authors declare no conflict of interest.

Supplementary Materials The online version of this article contains supplementary materials.

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