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Anti-inflammatory and ECM gene expression modulations of β -eudesmol via NF- κ B signaling pathway in normal human dermal fibroblasts

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Abstract

Background: β -eudesmol is a kind of aromatic compound belonging to sesquiterpenoid which exists within not only the bark of *magnolia* but also *Nardostachys jatamansi*, *Attractylodes lancea*, *Pterocarpus santalinus*, *Ginkgo bilobal*, *Cryptomeria japonica*, etc., and there has been progress in medical and pharmaceutic researches on antitumor, anticancer, and anti-inflammatory effects; nervous system stabilization; and vasodilator effects, etc., but not in researches on skin cares and cosmetics at all. Therefore, this study pretreated β -eudesmol with human dermal fibroblasts (HDFs) and then gave oxidative stresses with H_2O_2 to examine antioxidation, anti-inflammatory, and cell preservation effects. Through this process, it proves the possibility of β -eudesmol as cosmetic materials.

Methods: This study verified the effectiveness of β -eudesmol through cell viability analysis, reactive oxygen estimation, associated β -galactosidase assay, nuclear factor-kappa B (NF- κ B) luciferase assay, and quantitative real-time polymerase chain reaction (qRT-PCR).

Results: The cell viability which decreased due to H_2O_2 increased as per dose-dependent manners of β -eudesmol. Also, at the 2,2-diphenyl-1-picrylhydrazyl (DPPH) radical scavenging activity assay, intracellular reactive oxygen species (ROS) quantitative analysis and glutathione (GSH) estimation, the relative levels which were changed by H_2O_2 treatment, showed attenuated or protective transition forms, depending on the concentration of β -eudesmol. Additionally, reduced superoxide dismutase 1 (*SOD1*) and catalase (*CAT*) gene expression by H_2O_2 were increased by β -eudesmol. As the result of the promoter activity analysis of NF- κ B which has a key role in inflammation and skin aging, NF- κ B activity decreases as β -eudesmol concentration increases, and also this study proves that gene expression of interleukin 1 beta (*IL-1 β*) which is a downstream gene of NF- κ B related to inflammatory response decreases as well as tumor necrosis factor-alpha (*TNF- α*) gene expression depending on the concentration of β -eudesmol.

Conclusions: Through these results, this study suggests there are anti-inflammatory effects by shutting out the NF- κ B pathway. Following the results of the extracellular matrix (ECM) regulating gene expression analysis, this study proves that oxidative stress-induced increased *MMP1* levels were decreased depending on the concentration of β -eudesmol and verifies that it hinders the collapse of collagen through inhibition of transcriptional activity of NF- κ B. From the result of β -eudesmol regulating tissue inhibitors of metalloproteinase (*TIMP*)-1 gene expression which hinders matrix metalloproteinase (MMP), activation and alteration of gene expression of collagen type I alpha 1 (*COL1A1*) underpins the above consequences. Through this research, it is considered that β -eudesmol as one of natural cosmetic materials with such effects as antioxidation, anti-inflammation, and cell preservation is worthy of notice.

Keywords: β -eudesmol, Antioxidant, Anti-inflammation, Cellular senescence, Fibroblast, Extra cellular matrix

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Background

The bark of *magnolia* and *magnolia obovata* which is the dried rhizodermis have been used as a traditional medicine for medical treatment of bronchitis, asthma, stomach disease, emotional instability disorder, and allergy in Korea, Japan, and China (Hoang et al. 2010). On the basis of this folk remedy, the research on central sedation of central nervous system through *magnolia* bark extract was reported in 1973 (Watanabe et al. 1973). Various biologically active substances extracted from the bark and rhizodermis of *magnolia* are essential oils such as β -eudesmol, α -pinenes, β -pinenes, and bornyl acetate and diphenyl compounds such as magnolol, honokiol, and alkaloids; magnocurarine; and magnoflorine. It is reported that some of these ingredients has pharmacologic effects on the nervous system (Watanabe et al. 1983; Chiou et al. 1997).

In this experiment, β -eudesmol is used among those biologically active substances of *magnolia*, and β -eudesmol is a kind of aromatic compound belonging to sesquiterpenoid which exists not only in the bark of *magnolia* but also in medical herbs such as *Nardostachys jatamansi*, *Atractylodes lancea*, *Pterocarpus santalinus*, *Ginkgo biloba*, and *Cryptomeria japonica* etc. (Li et al. 2013), and its chemical formula is $C_{15}H_{26}O$ and its molecular mass is 222. β -eudesmol has antimutagenic effects (Miyazawa et al. 1996) and nervous system sedation effects too. It has been proved that β -eudesmol has effects to shut off nicotinic acetylcholine receptor in the neuromuscular junction (Kimura et al. 1991), to control neuromuscular disorder caused by neostigmine (Chiou and Chang 1992), and to control activities of Na^+ , K^+ -ATPase, and H^+ (Satoh et al. 1992). Also, it has been reported that β -eudesmol controls fatal toxicity caused by organophosphorus compound (Chiou et al. 1995), induces outgrowing of neurite in pheochromocytoma (PC12) cells through activation of mitogen-activated protein kinases (MAPK) (Obara et al. 2002), and has vasodilator effects through shutting off adrenaline α -1 receptor (Lim and Kee 2005). It has been reported that β -eudesmol controls interleukin (IL)-6 and receptor interacting protein-2 in mast cell, activates p38 MAPK, and has anti-inflammation effects through process to control caspase-1 (Seo et al. 2011). Recently, researches on effectiveness of β -eudesmol to blood vessels on the nervous system and also antitumor and anticancer effects have been largely in progress. Apoptotic effect through neovascular control effect of β -eudesmol has been reported (Ikeda and Nagase 2002; Ma et al. 2008). And it has been proved that β -eudesmol has apoptotic effect through caspase-3 via caspase-9 caused by cytochrome in HL-60 cells (Hoang et al. 2010), c-Jun N-terminal kinase (JNK)-dependent apoptotic effect through mitochondria passage in HL-60 cells (Li et al. 2013), and anticancer effects in the experiment of nude mouse to whom it implanted through

the method of heterotransplantation of human cholangiocarcinoma (Plengsuriyakarn et al. 2015).

Looking at existing studies on β -eudesmol, there have been reports about medical and pharmacologic researches on its anti-inflammation, antimutagenicity, nervous system stabilization, vasodilator, antitumor, and anticancer effects (Li et al. 2013; Miyazawa et al. 1996; Kimura et al. 1991; Chiou and Chang 1992; Satoh et al. 1992; Chiou et al. 1995; Obara et al. 2002; Lim and Kee 2005; Seo et al. 2011; Ikeda and Nagase 2002; Ma et al. 2008; Plengsuriyakarn et al. 2015). However, research on the mechanism of β -eudesmol in human dermal fibroblast has not been reported yet, and also, there have been no researches on β -eudesmol as skin care and cosmetic compounds. Therefore, this research intends to verify antioxidation, anti-inflammation, and cell preservation effects of β -eudesmol in human dermal fibroblast through studying the intracellular mechanism. Therefore, this study aims to suggest possibility of β -eudesmol as natural cosmetic materials.

Methods

Cell culture

For this research, we purchased and used human dermal fibroblasts (HDFs) from Lonza Inc. (Basel, Switzerland) and cultured it using Dulbecco's modified Eagle's medium (DMEM; Hyclone, Logan, UT, USA) as culture medium which contains 10% fetal bovine serum (FBS; Hyclone) and 1% penicillin/streptomycin (penicillin 100 IU/mL, streptomycin 100 μ g/mL; Invitrogen/Life Technologies, Carlsbad, CA, USA). Cultured cells within an incubator where we kept in a temperature of 37 °C and 5% CO_2 .

Sample treatment

We purchased β -eudesmol in the form of powder which is refined (> 90%) from Sigma-Aldrich Inc. (St. Louis, MO, USA). When we used it in the experiment, we dissolved it in dimethyl sulfoxide (DMSO; Sigma-Aldrich) in optimal concentration. After, we cultured HDFs (1×10^6 cells/well) in a 60-mm cell culture dish for 24 h; we added β -eudesmol as per indicated concentration to the culture medium and pretreated for 24 h, treated H_2O_2 in an appropriate concentration, and then used them for analysis after 3 h. For the experiment of gene level, after culturing HDFs (2×10^5 cells/well) in a cell culture medium, we treated β -eudesmol as per indicated concentration for 24 h when the plate's density is more than 85–90%. We treated H_2O_2 and collected cells after 3 h, and from these cells, we extracted RNA; then on the basis of this RNA, we identified gene expression through quantitative real-time polymerase chain reaction (qRT-PCR).

Cell viability estimation

We used the principle of water-soluble tetrazolium salt (WST-1) assay for evaluating cell viability. After we

inoculated each HDF of 100 μ L in density of 3×10^3 cells/well in 96-well plates and cultured for 24 h, we treated H_2O_2 and every kind of sample, in cell culture plates. After we added 10 μ L of EZ-Cytox cell viability assay kit reagent (ItsBio, Seoul, Korea) in the cultured cell and cultured for 1 h, we used a microplate reader (Bio-Rad, Hercules, CA, USA) to estimate absorbance in the scale of 490 nm and repeated three times and deduced the average value and standard deviation of cell viability.

cDNA manufacturing

After using Trizol reagent (Invitrogen/Life Technologies) to dissolve cells which we obtained through cell culture, we added 0.2 mL of chloroform (Biopure, Tulln, Austria) and then centrifuged for 20 min at 12,000 rpm at 4 °C to divide into pellets which include protein and supernatant which include mRNA. For the supernatant, we added 0.5 mL of isopropanol (Biopure) and left at room temperature for 10 min, and then centrifuged it at 12,000 rpm at 4 °C to precipitate RNA. Next, we used 75% ethanol to wash, and then we removed ethanol and dried at room temperature. We dissolved dried RNA in diethylpyrocarbonate (DEPC; Biopure) water, and among the extracted RNA, we only used pure RNA whose purity is more than a ratio 1.8 of 260/280 nm using Nanodrop (Maestrogen, Las Vegas, NV, USA).

For cDNA synthesis, we manufactured total 10 μ L of 1 μ g RNA, 0.5 ng oligo dT18, and DEPC water in a PCR tube, and then we treated them for 10 min at the degree of 70 °C to induce RNA denaturation. Next, we used M-MLV reverse transcriptase (Enzynomics, Dajeon, Korea) to react for 1 h at 37 °C to synthesize cDNA.

Quantitative real-time PCR

To analyze gene expression pattern quantitatively within HDFs caused by β -eudesmol, we used qRT-PCR method. The qRT-PCR is to synthesize 0.2 μ M primers, 50 mM KCl, 20 mM Tris/HCl pH 8.4, 0.8 mM dNTP, 0.5 U Taq DNA polymerase, 3 mM $MgCl_2$, and 1 \times SYBR green (Invitrogen/Life Technologies) in a PCR tube to manufacture the reaction solution, and is to use Linegene K (BioER,

Hangzhou, China). To denature DNA, the mixture is heated for 3 min at 94 °C, and then 40 cycles of denaturation (94 °C, 30 s), annealing (58 °C, 30 s), and polymerization (72 °C, 30 s) were performed. We used SYBR green to identify changing of each gene expression and verified effectiveness of PCR through the melting curve. We standardized expression of β -actin for comparative analysis on each gene expression. The primer used in the experiment is shown in Table 1.

DPPH radical scavenging activity assay

DPPH assay is a method to inject a sample diluent of 100 μ L of each concentration respectively on a 96-well plate and add 50 μ L of DPPH of 0.2 mM, and then shut off the light at room temperature and neglect it for 30 min. We used a microplate reader (Bio-Rad) to estimate absorbance caused by DPPH reduction in the scale of 514 nm and repeated to perform estimation three times and deduced average value and standard deviation of absorbance.

Intracellular reactive oxygen species (ROS) quantitative analysis

To estimate the changing of concentration of ROS within cells, we inoculated HDFs of 2×10^5 cells/well in a 60-mm culture medium and cultured for 24 h and afterwards we treated cells properly and then cultured for 24 h. Then we added 10 μ M of dichlorofluorescein diacetate (DCF-DA; Sigma-Aldrich) which is needed to estimate ROS within cells and cultured for 30 min. Subsequently, we added phosphate buffered saline (PBS) to obtained cells and set them free; finally, we estimated amount of changing of ROS by using a flow cytometer (BD Biosciences, San Jose, CA, USA). To verify ROS scavenging effects of β -eudesmol, we also treated L-ascorbic acid which acts as an ROS scavenger, and then we estimated through the same process.

Glutathione (GSH) estimation

As an indicator for toxicity reaction inducing apoptotic and oxidative stress, GSH level change has been estimated (Esposito et al. 2000; Zhang et al. 2010; Kil et al.

Table 1 Lists of primers used in this study

Gene	Forward primer	Reverse primer
<i>β-actin</i>	GGATTCCTATGTGGGCGACGA	CGCTCGGTGAGGATCTTCATG
<i>SOD1</i>	GGGAGATGGCCAACTACTG	CCAGTTGACATCGAACCGTT
<i>CAT</i>	ATGGTCCATGCTCTCAAACC	CAGGTATCCAATAGGAAGG
<i>TNF-α</i>	CCCAGGGACCTCTCTAATC	GGTTTGCTACAACATGGGCTACA
<i>IL-1β</i>	GATCCATTCTCCAGCTGCA	CAACCAAGTATTCTCCATG
<i>COL1A1</i>	AGGGCCAAGACGAAGACATC	AGATCACGCATCGCACACA
<i>MMP1</i>	GGGCTTAGATCATTCTCAGTGCC	CAGGGTGACACCACTGACTGCAC
<i>TIMP1</i>	AACCCACCCACAGACA	ACCCATGAATTTAGCCCTTA

2012). We used ThiolTracker™ Violet Glutathione Detection Reagent (Invitrogen) to estimate the amount of reduced GSH. Before pretreatment of β -eudesmol for 24 h, HDF cells were seeded as 2×10^5 cells/well in a 60-mm culture dish and cultured for 24 h. Next cells were treated 500 μ M H_2O_2 and cultured for 3 h more. After we obtained cultured cells, we centrifuged them at 5000 rpm at 4 °C for 5 min to precipitate cells, and we removed supernatant and set cell pellet free through PBS of 300 μ L. Then we added ThiolTracker™ Violet dye of 300 μ L to cells. After blending softly, we cultured in the darkroom at room temperature for 30 min. And then, we washed cells through PBS, and in the condition of 5000 rpm at 4 °C, cells were centrifuged for 5 min and supernatant removed. Using excitation and emission at 405 and 525 nm each, a flow cytometer (BD Biosciences) was used to estimate fluorescent value.

Cellular senescence estimation

We estimated senescence by using senescence-associated beta-galactosidase (SA- β -gal) assay which is a method using a senescence detection kit (Biovision, Milpitas, CA, USA). After we inoculated HDFs of 2×10^5 cells/well in a 60-mm culture dish and cultured for 24 h, we treated cells properly and cultured for 24 h more. Then we eliminated a culture medium from the plates and washed with 1 mL PBS, and added 0.5 mL fixing solution for 15 min. Next, we added 0.5 mL mixed staining solution (staining solution 470 μ L, staining supplement 5 μ L, 20 mg/mL X-gal in dimethylformamide (DMF) 25 μ L) to each fixed HDFs and cultured for 24 h at 37 °C. After washing dyed cells with PBS, we estimated numbers of dyed cells through an optical microscope (Olympus, Tokyo, Japan) to analyze senescent cell portion. We calculated the numbers of total cells and dyed cells and figured out the ratio of senescent cells to identify.

NF- κ B luciferase assay

To identify influence of β -eudesmol on NF- κ B activity, we used NF- κ B promoter luciferase assay in this experiment. We used NF- κ B reporter NIH-3T3 stable cell line (Panomics, Fremont, CA, USA) including reporter gene

(luciferase gene) which contains NF- κ B promoter consensus sequence at the promoter region. As for the promoter activity of NF- κ B, the transcription factor is proportional to the amount of luciferase gene expression; therefore, through this, we identified activity of such a transcription factor as NF- κ B has an influence on skin inflammatory and skin aging.

After we seeded NF- κ B reporter NIH-3T3 stable cell of 2×10^5 cells/well in a 60-mm culture dish and cultured for 24 h, we treated cells in the proper condition like the above experiments, and cultured for 24 h more. Then we obtained cultured cells, added passive lysis buffer (Promega, Madison, WI, USA), on ice for 10 min to dissolve; next, we centrifuged for 30 min in the condition of 12,000 rpm at 4 °C and collected supernatant. After aliquoting 80 μ L of the supernatant which contains the same amount of protein in each black 96-well plate, we added and mixed luciferin (Promega) subsequently. Because luciferin is sensitive to the light, we used a Veritas luminometer (Turner Designs, Sunnyvale, CA, USA) to estimate luminance of luciferin right after adding it to the sample.

Statistical process

All experiments of this research were performed more than three times separately under the same condition to get experimental results, and we used Student's *t* test for every experiment and analyzed that it is statistically significant when *p* value of every experimental result is less than 0.05.

Results

Cell viability

After we treated each HDF through β -eudesmol in various concentrations of 5, 10, 20, 40, and 80 μ M and cultured for 24 h, we used WST-1 assay to estimate cell viability. As the result that we identified cell viability, it showed viability of 101, 107, 98, 91, and 83%, respectively. For concentration of 20 μ M, it hardly had an influence on viability, and it showed that cell viability decreased at the case of treatment through β -eudesmol in their concentrations of 40 and 80 μ M (Fig. 1a). After

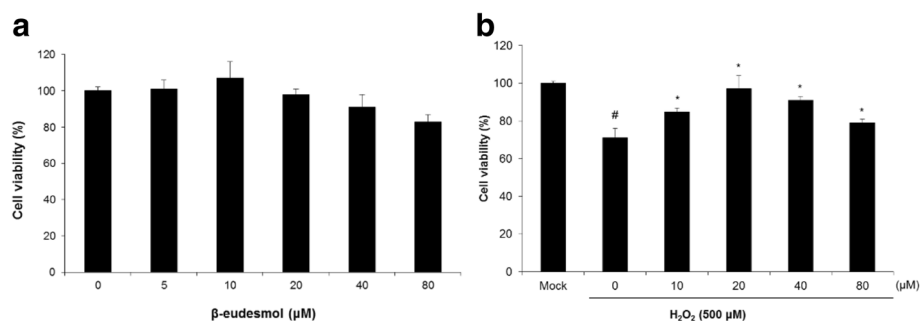


Fig. 1 Cell viability on β -eudesmol in HDFs. **a** Cytotoxicity on β -eudesmol in HDFs. **b** Cell viability on H_2O_2 in the HDFs pretreated with β -eudesmol. (* $p < 0.05$)

pretreating HDFs through β -eudesmol in concentrations of 5, 10, and 20 μ M respectively for 24 h and treating with 500 μ M H_2O_2 for 3 h, we made observations of cell viability. Assuming cell viability is 100%, in that case, we did not treat HDFs through both β -eudesmol and H_2O_2 , cell viability decreased to 71% when we did not treat HDFs through β -eudesmol but H_2O_2 of 500 μ M. However, cell viability was 85% at the case of pretreatment through β -eudesmol at 5 μ M, 85% at the case of pretreatment through β -eudesmol of 10 μ M, and 97% at the case of pretreatment through β -eudesmol of 20 μ M; therefore, through this result, we identified that cell viability of HDFs recovered depending on concentrations. On the contrary, cell viability at the case of pretreatment through β -eudesmol of 40 and 80 μ M was shown as 91 and 79% respectively; therefore, we identified that cell viability decreased in concentration more than 40 μ M. So we used concentration of β -eudesmol to 20 μ M at the most for HDFs in further experiments (Fig. 1b).

Antioxidative effects of β -eudesmol

Using DPPH radical scavenging activity assay, we identified radical scavenging effects of β -eudesmol. As positive control group, we performed comparative analysis on N-acetyl-L-cysteine (NAC) which is known as an ROS scavenger. At the case of treatment through β -eudesmol in concentrations of 5, 10, and 20 μ M, radical scavenging effects increased to 5, 31, and 49% respectively depending on concentration. At the case of treatment through NAC as positive control group in concentrations of 5, 10, and 20 μ M, we identified that radical scavenging effects are shown as 8, 25, and 46% respectively. Identified that β -eudesmol has similar radical scavenging effects with NAC, it can be admitted that β -eudesmol has positive radical scavenging effects (Fig. 2a). To identify effects of β -eudesmol on decrease of total amount of intracellular ROS which occurs in the course of cell metabolism and H_2O_2 addition, we used DCF-DA as a fluorescent probe to perform flow cytometry, and after pretreating HDFs through

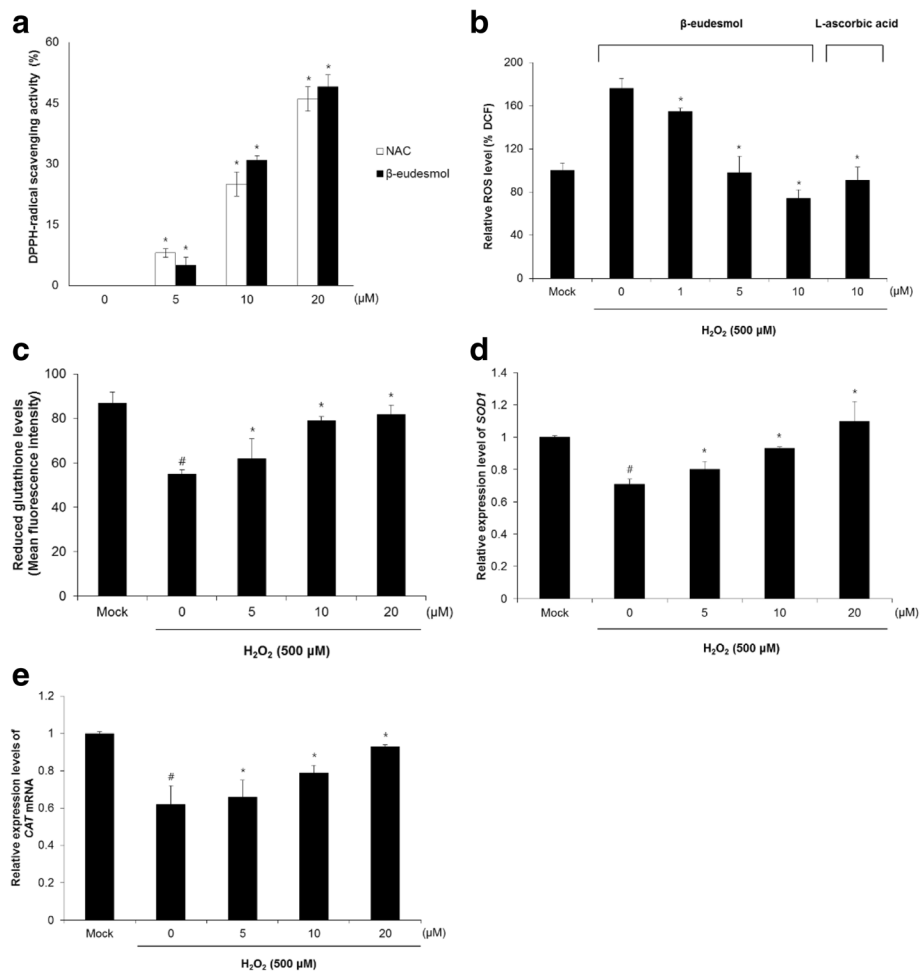


Fig. 2 Antioxidative effect of β -eudesmol against H_2O_2 . **a** The DPPH radical scavenging activity of β -eudesmol. **b** H_2O_2 -induced intracellular ROS scavenging activity of β -eudesmol in HDFs. **c** Reduced GSH level in HDFs with pretreated β -eudesmol before H_2O_2 treatment. **d** *SOD1* and **e** *CAT* mRNA gene expression analysis in HDFs with β -eudesmol over H_2O_2 treatment. (* $p < 0.05$)

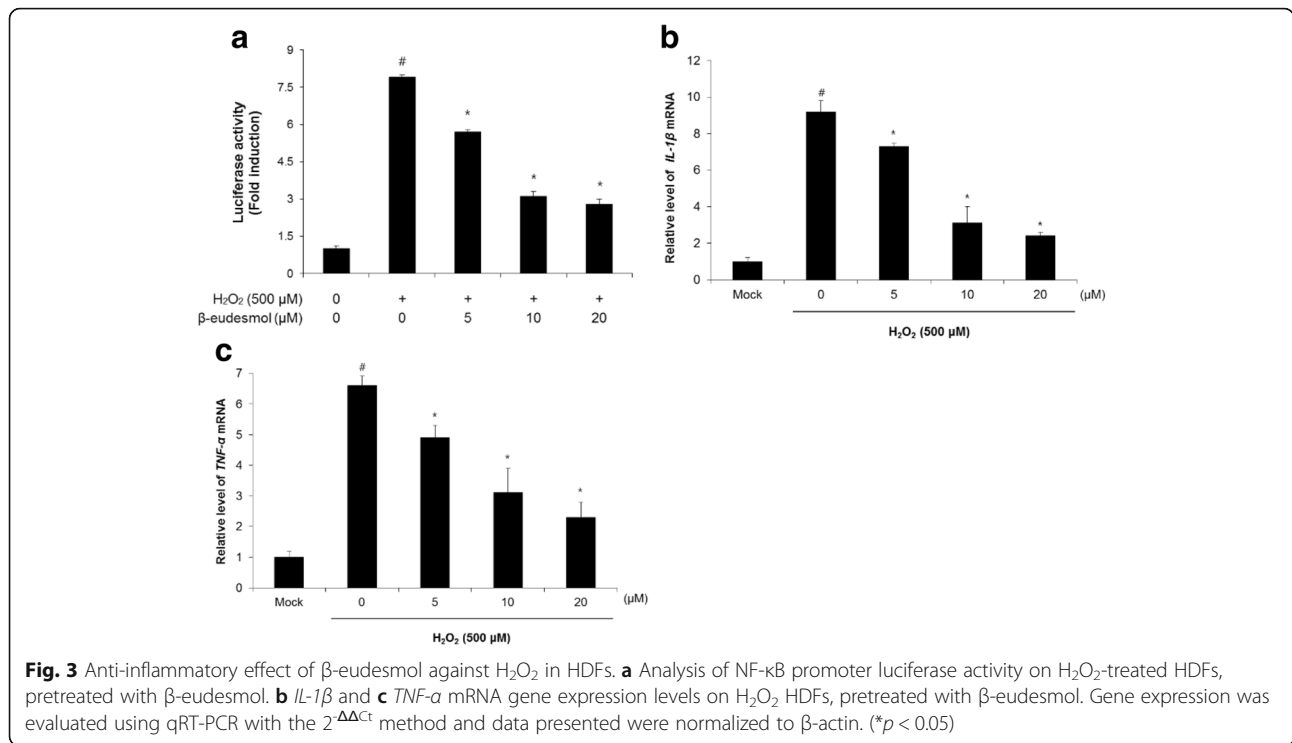
β -eudesmol for 24 h, we treated HDFs through H_2O_2 of 500 μ M. Its results which we analyzed after 3 h are as follows: Assuming it is 100%, in that case, we did not treat HDFs through both β -eudesmol and H_2O_2 , total amount of ROS increased to 176% at the case of treatment through H_2O_2 . However, at the case of treatment through β -eudesmol in each concentration of 1, 5, and 10 μ M, it was identified that the total amount of ROS decreases to 155, 98, and 74% respectively. At the case of L-ascorbic acid as the positive control group, total amount of ROS was shown as 91% when we treated in concentration of 10 μ M. This result is similar to the case of treatment through β -eudesmol in concentration of 5 μ M; therefore, we identified that β -eudesmol has an effect on decrease of total amount of intracellular ROS (Fig. 2b). Using the principle that ROS lowers intracellular GSH level through oxidative reaction or reaction to thiol group, we used Violet dye which reacts to thiol group to identify the glutathione level reduced form. As for the result that we pretreated HDFs through β -eudesmol for 24 h and then treated through H_2O_2 of 500 μ M and analyzed 3 h later, when we treated nothing, GSH level was shown as 87, but at the case of treatment through H_2O_2 , it decreases to 55. However, at the case of pretreatment through β -eudesmol in each concentration of 5, 10, and 20 μ M, GSH level increased to 62, 79, and 82 respectively; therefore, we identified that both results were similar to each other and GSH levels recovered at the same level (Fig. 2c). To identify antioxidation effects of β -eudesmol at a gene level, we identified changing of antioxidation enzyme *SOD1* gene expression through qRT-PCR. Assuming *SOD1* gene expression level is 1 at the case of no treatment for both β -eudesmol and H_2O_2 , *SOD1* gene expression decreased to 0.71 at the case of treatment through H_2O_2 . However, when we pretreated through β -eudesmol in each concentration of 5, 10, and 20 μ M for 24 h, treated through H_2O_2 , and then analyzed 3 h later, *SOD1* gene expression increased to 0.8, 0.93, and 1.1 respectively. Through this result, we were able to identify that β -eudesmol increased *SOD1* gene expression to have an influence on antioxidation effects in this experiment (Fig. 2d). Catalase (CAT) as the antioxidation enzyme has a role as catalyst to react to intracellular H_2O_2 and extinct radical to change into H_2O . To identify antioxidation effects of β -eudesmol, we checked changing of catalase gene, *CAT* expression. Assuming *CAT* gene expression level is 1 at the case of no treatment through both β -eudesmol and H_2O_2 , *CAT* gene expression decreased to 0.62 at the case of treatment through H_2O_2 . However, when we pretreated through β -eudesmol in each concentration of 5, 10, and 20 μ M and then treated through H_2O_2 , *CAT* gene expression increased to 0.66, 0.79, and 0.93 respectively as per a dose-dependent manner of β -eudesmol. Through this result, we identified that β -eudesmol increased *CAT* gene expression to have intracellular antioxidation effects (Fig. 2e).

Anti-inflammatory effects of β -eudesmol

Through NF- κ B luciferase assay, we identified promoter activity of NF- κ B, the transcription factor. We used NIH-3T3 stable cell line including luciferase gene which contains NF- κ B promoter consensus sequence in the field of promoter, added luciferin which reacts to luciferase to form fluorescer, and estimated the amount of luciferase gene expression through luciferin luminance estimation. When we treated through H_2O_2 , luminance of luciferin increased to 7.9 times compared with control group which were non-treated with HDFs, but when we treated through H_2O_2 after pretreating β -eudesmol in a concentration of 5 μ M, it decreased to 5.7 times. At the case of pretreatment through β -eudesmol in a concentration of 10 μ M and treatment through H_2O_2 , it decreased to 3.1 times; at the case of pretreatment through β -eudesmol in a concentration of 20 μ M and treatment through H_2O_2 , it decreased to 2.8 times; therefore, we identified that luciferin luminance decreased depending on the concentration of β -eudesmol. Therefore, we identified that β -eudesmol hindered transcriptional activity of NF- κ B (Fig. 3a). We checked changing of *IL-1 β* gene expression which is related to early inflammation reaction. *IL-1 β* gene expression increased to 9.2 times when we treated through H_2O_2 than when we did not treat through both β -eudesmol and H_2O_2 . However, when we pretreated through β -eudesmol in a concentration of 5 μ M and then treated through H_2O_2 , the amount of *IL-1 β* gene expression depending on concentrations of β -eudesmol decreased to 7.3 times. At the case of pretreatment through β -eudesmol in a concentration of 10 μ M and treatment through H_2O_2 , it decreased to 3.1 times; at the case of pretreatment through β -eudesmol in a concentration of 20 μ M and treatment through H_2O_2 , it decreased to 2.4 times (Fig. 3b). Subsequently, after we pretreated HDFs through β -eudesmol in each concentration for 24 h and treated through H_2O_2 for 3 h, we analyzed changing of *TNF- α* gene expression. When we treated HDFs through H_2O_2 , it increased to 6.6 times comparing with the control group which we did not treat through both β -eudesmol and H_2O_2 . However, when we pretreated through β -eudesmol in a concentration of 5 μ M for 24 h and then treated through H_2O_2 , the amount of *TNF- α* gene expression depending on concentration of β -eudesmol decreased to 4.9 times; at the case of pretreatment through β -eudesmol in concentration of 10 μ M for 24 h and treatment through H_2O_2 , it decreased to 3.1 times; and at the case of pretreatment through β -eudesmol in concentration of 20 μ M for 24 h and treatment through H_2O_2 , it decreased to 2.3 times (Fig. 3c).

Cellular senescence and extracellular matrix (ECM) modulation gene expression analysis of β -eudesmol

SA β -gal assay is particularly observed in senescent cells and is largely used as an indicator of senescence (Dimri et al. 1995). When we did not treat HDFs through both



β -eudesmol and H_2O_2 , β -galactosidase activity was shown as 6, and at the case of treatment through H_2O_2 , it increased to 69-folds. However, when we pretreated through β -eudesmol in a concentration of 5 μ M for 24 h and then treated through H_2O_2 , β -galactosidase activity depending on concentration of β -eudesmol decreased and was shown as 51; at the case of pretreatment through β -eudesmol in a concentration of 10 μ M for 24 h and treatment through H_2O_2 , it was shown as 27; and at the case of pretreatment through β -eudesmol in a concentration of 20 μ M for 24 h and treatment through H_2O_2 , it was shown as 18 (Fig. 4a). We used qRT-PCR method to identify changing of *COL1A1* gene expression which forms type I collagen constituting 80 to 85% of collagen. The relative *COL1A1* gene expression decreased to 0.39 at the case of treatment of H_2O_2 compared with non-treated cells. However, we identified that *COL1A1* gene expression increases depending on concentration of β -eudesmol, as the result that at the case of pretreatment through β -eudesmol in a concentration of 5 μ M and treatment through H_2O_2 , it was shown as 0.54; at the case of pretreatment through β -eudesmol in a concentration of 10 μ M and treatment through H_2O_2 , it was shown as 0.72; and at the case of pretreatment through β -eudesmol in a concentration of 20 μ M and treatment through H_2O_2 , it was shown as 0.88 (Fig. 4c). To identify changing of *MMP1* gene expression which decomposes type I collagen constituting most of dermis, we used qRT-PCR method to identify influence of β -

eudesmol on dermal tissue. When we did not treat HDFs through both β -eudesmol and H_2O_2 , *MMP1* gene expression was 0.51, and *MMP1* gene expression was doubled to be shown as 1 at the case of treatment through H_2O_2 . However, we identified that *MMP1* gene expression decreases depending on concentration, as the result that at the case of pretreatment through β -eudesmol in a concentration of 5 μ M and treatment through H_2O_2 , it was 0.84; at the case of pretreatment through β -eudesmol in a concentration of 10 μ M and treatment through H_2O_2 , it was 0.69; and at the case of pretreatment through β -eudesmol in a concentration of 20 μ M and treatment through H_2O_2 , it was 0.58. Particularly, *MMP1* gene expression at the case of treatment through β -eudesmol in a concentration of 20 μ M was similar to the result at the case of no treatment (Fig. 4c). We identified changing of *TIMP1* (Fisher Jr and Zheng 1996; Enjoji et al. 2000) gene expression which is a hindrance factor to MMP1 and MMP9 to identify its influence on collagen metabolism. Assuming *TIMP1* gene expression is 1 when we did not treat HDFs through both β -eudesmol and H_2O_2 , *TIMP1* gene expression decreased to 0.3 sharply at the case of treatment through H_2O_2 . However, when we pretreated through β -eudesmol in a concentration of 5 μ M and then treated through H_2O_2 , *TIMP1* gene expression depending on concentration increased to be shown as 0.35; at the case of pretreatment through β -eudesmol in a concentration of 10 μ M and treatment through H_2O_2 , it

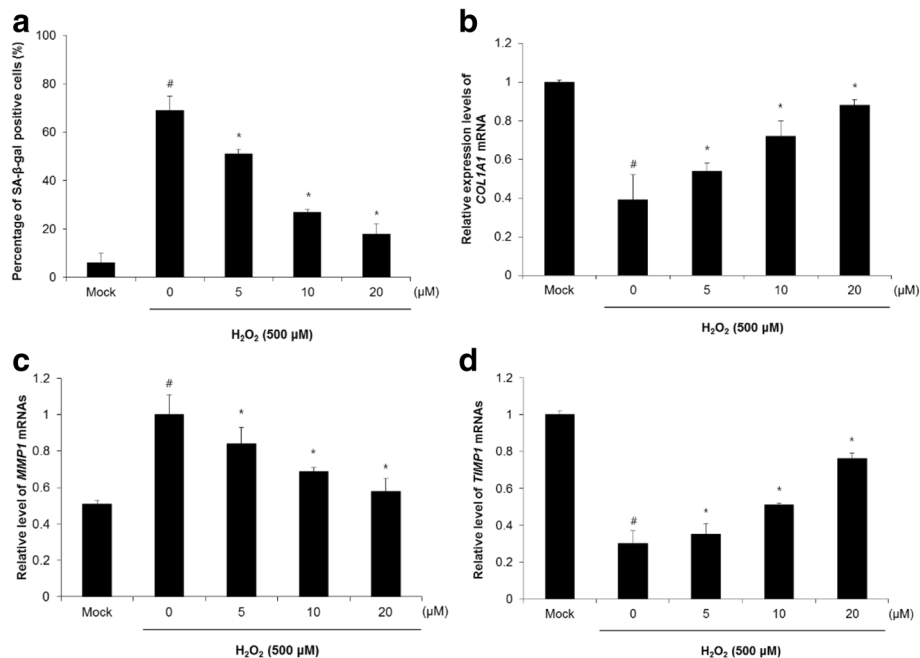


Fig. 4 Senescence attenuation effect of β -eudesmol against H_2O_2 in HDFs. **a** Protective effect of β -eudesmol on H_2O_2 -induced cellular senescence in HDFs using SA- β -gal assay. **b** *COL1A1*, **c** *MMP1*, and **d** *TIMP1* mRNA gene expression levels on H_2O_2 -treated HDFs, pretreated with β -eudesmol. Gene expression was evaluated using qRT-PCR with the $2^{-\Delta\Delta C_t}$ method and data presented were normalized to β -actin. (* $p < 0.05$)

was 0.51; and at the case of pretreatment through β -eudesmol in a concentration of 20 μ M and treatment through H_2O_2 , it was 0.76 (Fig. 4d).

Discussion

Antioxidation effects of β -eudesmol

When human skin is under oxidative stress caused by ROS, DNA, lipid, and protein damage can occur (Devasagayam and Kamat 2002). When intracellular ROS increases, through various intracellular signal transmission processes, it hinders collagen synthesis and promotes MMP expression which is an enzyme to decompose collagen to accelerate causing wrinkles and skin aging (Lavker and Kaidbey 1997). Also oxidative stress is deeply related to not only skin aging but also inflammation reaction so that it has an influence on controlling activity of various kinds of cells. If inflammation reaction is not controlled well, sometimes it will lead to a reaction to accelerate senescence (De Martinis et al. 2005).

To minimize damage caused by ROS, our body uses antioxidation enzymes such as SOD, CAT, and glutathione peroxidase (GPX) which are our defense mechanisms or methods to supply antioxidation substances such as vitamin (Vit) C, Vit E, and ubiquinone into our body, and research on development of antioxidation substances has been in progress continuously (Choi et al. 2007). In this research, we used β -eudesmol to apply to HDFs and identified its possibility as an antioxidant suitable for skin.

Through experimental methods such as DPPH radical activity assay, ROS quantitative analysis using DCF-DA, GSH estimation, *SOD1* gene expression, and *CAT* gene expression, we verified antioxidation effects of β -eudesmol. As the result of experiment through DPPH radical activity assay, we identified that radical extinction effects increase depending on concentration of β -eudesmol and that similar radical extinction effects appear comparing the case of N-acetyl-L-cysteine, the positive control group; therefore, we identified there were radical extinction effects. Although we identified antioxidation effects of β -eudesmol sample through DPPH radical activity assay, we also identified intracellular antioxidation effects through ROS quantitative analysis using DCF-DA and GSH estimation. The total amount of intracellular ROS in HDFs decreased depending on concentration of β -eudesmol, and particularly at the case of treatment through β -eudesmol in a concentration of 5 μ M, it showed a similar result to a positive control group which we treated through L-ascorbic acid used as antioxidation effect marker in a concentration of 10 μ M; therefore, it was considered that β -eudesmol had an effect to remove ROS. To identify effects of β -eudesmol to extinct intracellular H_2O_2 and hinder OH^- radical formation, we used a GSH measurement analysis method. As an indicator of toxicity reaction which induces apoptotic effect and oxidative stress, changing of GSH level has been estimated (Esposito et al. 2000; Zhang et al.

2010; Kil et al. 2012). It was identified that GSH level increased in accordance with an increase in concentration of β -eudesmol.

When human skin is under oxidative stress, signal transmission system is activated, in which cells react, increase radical formation, and decrease antioxidation enzyme expression (Masaki et al. 1995; Barber et al. 1998; Yasui and Sakurai 2000; Yamamoto and Gaynor 2001). To verify antioxidation effects of β -eudesmol at the gene level, we identified changing of antioxidation enzyme *SOD1* and *CAT* gene expression. It was identified that *SOD1* and *CAT* gene expression increased depending on concentration of β -eudesmol. Based on the above results, it was identified that β -eudesmol had not only antioxidation effects mediated with biological enzymes but also radical extinction effects with various methods. β -eudesmol sample itself had antioxidation effects and terminated intracellular ROS effectively. Also it was identified that it decreased antioxidation enzyme expression which is a ROS defense mechanism and controls ROS from the gene level to extinct ROS. Therefore, it was considered that β -eudesmol had hindrance effects to various damages and stimulations caused by oxidative stress on human skin (Fig. 5).

Anti-inflammation effects of β -eudesmol by controlling NF- κ B promoter activity

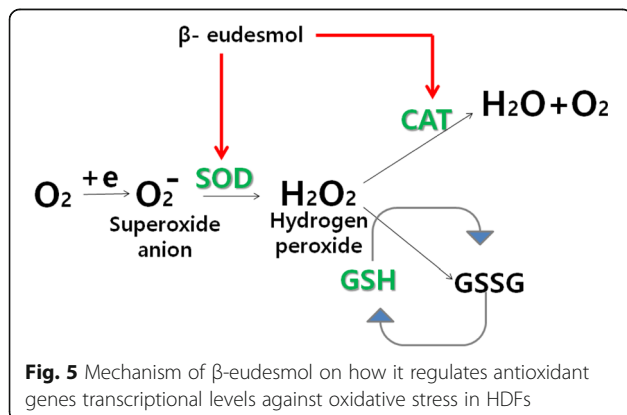
NF- κ B is the named transcription factor whose role is identified to be concerned with controlling immunoglobulin kappa chain gene expression in B cells (Sen and Baltimore 1986). Through classical pathway and alternative pathway, NF- κ B has a decisive role of inflammation reaction, immune reaction, cell proliferation, and apoptotic effects in various processes, and it has been identified as a survival factor in various cells against lipopolysaccharide (LPS), cytokine, ROS, and stimulation by ultraviolet (UV) rays (Bauerle and Henkel 1994; Siebenlist et al. 1994; Kopp and Ghosh 1995; Ghosh et al. 1998; Karin and Delhase 2000). NF- κ B is composed of homo or hetero-dimer and formed with five subunits

such as p65 (RelA), RelB, c-Rel, p50 (NF- κ B 1), and p52 (NF- κ B 2) (Gilmore 2006). NF- κ B is combined with inhibitor of NF- κ B (I κ B) to exist in the cytoplasm with its state of deactivation, but I κ B kinase (IKK) is activated through stimulation of UV rays or ROS on cells and phosphorylates I κ B to be separated. Phosphorylated I κ B is decomposed by ubiquitination and separated from proteasome, and NF- κ B flows into the nucleus and works as a transcription factor (Ghosh and Karin 2002; He and Karin 2011). TNF- α , IL, and chemokine, which are a product of inflammatory precursor, have a role to mediate an important course of cell activities such as cell proliferation, angiogenesis, and metastasis. Transcription of gene expression within which these molecules are coded is controlled by NF- κ B (Gloire et al. 2006). Through intracellular signal transmission, NF- κ B controls various cell metabolism, but in that case, it works as transcription factor of inflammation-mediated substances and it promotes transcription of COX2, E-selectin, inducible nitric oxide synthase (iNOS), intercellular adhesion molecule-1 (ICAM-1), IL-6, IL-12, IL-1 β , MMP, TNF- α , and vascular cell adhesion molecular 1 (VCAM-1) to accelerate inflammation reaction and cause tissue damage (Limuro et al. 1997; Bukman et al. 1998; Chen et al. 1999; Suschek et al. 2004; Yamamoto and Gaynor 2001; Farooqui et al. 2007).

To identify that influence of β -eudesmol on control of NF- κ B activity which has a significant role in inflammation reaction, we used NF- κ B luciferase assay method to identify NF- κ B activity, and it was found that NF- κ B activity decreased depending on concentration of β -eudesmol. In this experiment, we examined effects to hinder NF- κ B, and also, we checked changing of downstream gene expression which increases through NF- κ B activation. Through a qRT-PCR method, we checked changing of gene expression of cytokine IL-1 β , TNF- α (Fisher Jr and Zheng 1996; Dinarello 1991) which is a representative medium of early inflammation reaction. It was identified that *IL-1 β* , *TNF- α* inflammation induction gene expression decreased as concentration of β -eudesmol increases. Through this experiment, we identified that β -eudesmol shut off the NF- κ B pathway to hinder *IL-1 β* , *TNF- α* gene expression and inflammation reaction.

Regulation mechanism of ECM gene expression of β -eudesmol

Increase in ROS within dermal tissue has an influence on intracellular transforming growth factor-beta (TGF- β) cytokine, activator protein 1 (AP-1) transcription factor, NF- κ B transcription factor, and signal transmission pathway of Smad3/4; controls collagen formation gene expression; and increases gene expression of MMPs which is an enzyme-decomposing collagen so that it causes changing of ECM tissue and skin aging (Lee et al.



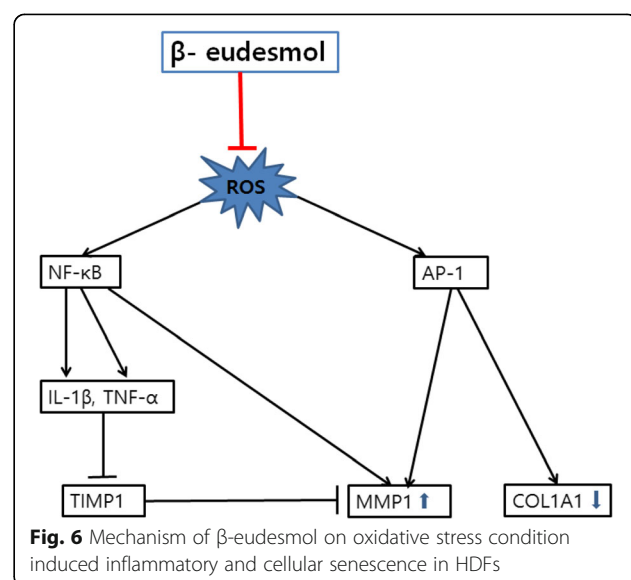
2012; Varani et al. 2002; Saito et al. 2004). AP-1, NF- κ B, and Smad from increase in ROS are activated by MAPK which works as an intracellular signal transmission factor. MAP-kinase p38, JNK, and extracellular signal-regulated kinase (ERK) exist in MAPK, and MAPK is concerned with various intracellular mechanisms such as cell growth, division, apoptosis, and gene expression (Kohl et al. 2011). When MAPK is activated by ROS, it activates NF- κ B to increase *MMP* gene expression and induce collagen decomposition (Lee et al. 2012; Bae et al. 2008). Moreover, it induces AP-1 activation, enters into cell nucleus, works as transcription factor to promote *MMP1*, *MMP3*, and *MMP9* gene expression, and hinders *COL1A1* gene expression (Quan et al. 2005; Lee et al. 2006). Increase in ROS causes hindering TGF- β which is a bifidus factor to control transcription activity of Smad (Leivonen and Kähäri 2007). Collagen gene expression which is one of their downstream genes is regulated and it induces collapse of ECM tissue within dermis to cause wrinkle, loss of elasticity, and skin aging (Quan et al. 2005). Within the dermis, TIMP which is an enzyme-hindering MMP activity, exists and with MMPs forms one to one complex to hinder MMP enzyme, and there are TIMP1, TIMP2, TIMP3, and TIMP4 as kinds of TIMP (Gomis-Rüth et al. 1997). MMP1 and MMP9 are enzymes which decompose type I collagen constituting 85% of collagen in the dermis, and it is TIMP1 enzyme which hinders MMP1 and MMP9 (Enjoi et al. 2000; Fisher et al. 1999). With its complicated mechanism, TIMPs control MMP activity, and when the balance between them is lost, it sometimes leads to tumor, and it has been reported that its gene expression increases through epidermal growth factor (EGF), TNF- α , IL-1, and TGF- β (Birkedal-Hansen 1993; Lee et al. 2016; Jang and Lee 2016).

Through changing of signal transmission process and gene expression of β -eudesmol in ECM, we identified its effects on human dermal fibroblast preservation and cell activation. Through experimental results, we identified that it controlled NF- κ B activity, and we checked changing of *COL1A1*, *MMP1* gene expression which are downstream genes of NF- κ B. *COL1A1* gene expression increased as concentration of β -eudesmol increased, and *MMP1* gene expression decreased depending on the concentration of β -eudesmol. We identified that *TIMP1* gene expression which hindered MMP1 increases depending on the concentration of β -eudesmol. Taking these experimental results together, it is identified that β -eudesmol shuts off the NF- κ B pathway to regulate *MMP1* gene expression and that *COL1A1* gene expression increases as the concentration of β -eudesmol increases. According to existing researches related to *COL1A1* gene expression, when AP-1 is activated by ROS, it promotes *MMP1*, *MMP3*, and *MMP9* gene

expression and hinders *COL1A1* gene expression (Quan et al. 2005; Lee et al. 2006). Also, it has been reported that ROS hinders a bifidus factor such as TGF- β to regulate transcription activity of Smad (Leivonen and Kähäri 2007) and control collagen gene expression which is a downstream gene (Quan et al. 2005). Therefore, it is considered that by β -eudesmol, AP-1 activity is hindered to regulate *MMP1*, *MMP3*, and *MMP9* gene expression, that β -eudesmol increases *COL1A1* gene expression and activates TGF- β , and that *COL1A1* gene expression increases through transcription activity of Smad. Through this experimental result, it is considered that β -eudesmol preserves tissue within the dermis and has effects to promote ECM formation to protect dermal cells. It has been identified that β -eudesmol has effects on cell preservation within the dermis at the gene level, but to find out whether this result proves effects to regulate cell senescence actually, through SA- β -gal assay experimental method, we identified human dermal fibroblast cell senescence. As for the experimental result of SA- β -gal assay, β -galactosidase activity decreased depending on concentration of β -eudesmol. Namely, it is found that β -eudesmol has effects to regulate cell senescence. Therefore, it is considered that β -eudesmol is highly useful as a cosmetic compound because it has antioxidation and anti-inflammation effects, delays dermal cell senescence, and has cell preservation effects (Fig. 6).

Conclusions

In conclusion, the present study showed that β -eudesmol has several effects on anti-inflammatory and ECM constructed gene expression in human dermal fibroblasts. Our results represented the antioxidant effect of β -eudesmol through improving intracellular antioxidant molecule and gene expressions, scavenging excessive generated ROS, and



radical scavenging capacity. Based on the above antioxidant results, we identified protective effect of β -eudesmol on NF- κ B-mediated inflammatory signaling activities. These preliminary results provided an interesting point in HDF: β -eudesmol may have impediment action to aging. Thus, SA- β -gal assay and ECM-related gene expression analysis were conducted, which suggested effective senescence-hindering capacity of β -eudesmol in dermal fibroblast cells. Additional further studies, *in vitro* and *in vivo*, will be necessary to verify molecular pathways and mechanisms of β -eudesmol in detail, but this study suggests β -eudesmol as a “cosmeceutical” compound.

Abbreviations

AP-1: Activator protein 1; CAT: Catalase; COL1A1: Collagen, type I, alpha 1; DCF-DA: 2',7'-Dichlorofluorescein diacetate; DEPC: Diethylpyrocarbonate; DMEM: Dulbecco's modified Eagle's medium; DMF: Dimethylformamide; DMSO: Dimethyl sulfoxide; DPPH: 2,2-Diphenyl-1-picrylhydrazyl; ECM: Extracellular matrix; EGF: Epidermal growth factor; ERK: Extracellular signal-regulated kinase; FBS: Fetal bovine serum; GPX: Glutathione peroxidase; GSH: Glutathione; HDF: Human dermal fibroblast; ICAM-1: Intercellular adhesion molecule-1; IKK: I κ B kinase; IL-12: Interleukin 12; IL-1 β : Interleukin 1 beta; IL-6: Interleukin 6; iNOS: Inducible nitric oxide synthase; JNK: c-Jun N-terminal kinases; LPS: Lipopolysaccharide; MAPK: Mitogen-activated protein kinase; MMP: Matrix metalloproteinase; NAC: N-Acetyl-L-cysteine; NF- κ B: Nuclear factor-kappa B; PC12: Pheochromocytoma; PBS: Phosphate buffered saline; qRT-PCR: Quantitative real-time polymerase chain reaction; ROS: Reactive oxygen species; SA- β -gal: Senescence-associated beta-galactosidase; SOD1: Superoxide dismutase 1; TIMP1: Tissue inhibitors of metalloproteinases-1; TNF- α : Tumor necrosis factor-alpha; TGF- β : Growth factor-beta; WST-1: Water-soluble tetrazolium salt; VCAM-1: Vascular cell adhesion molecular 1; Vit C: Vitamin C

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Author's contributions

KYK did all of research background such as experiments, data collecting, and statistical analysis as well as drafting the manuscript.

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Consent for publication

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Competing interests

The author declares no competing interests.

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