INTRODUCTION

Hair follicles are composed of cells that possess self-renewal capacity, which can undergo a repetitive regeneration process during hair growth (Yu et al., 2008). The ‘hair growth cycle’ has three phases: the anagen (growth), catagen (regression) and telogen (rest) phases (Stenn and Paus, 2001). During the anagen phase, the pigmented hair shaft is actively generated and the follicle reaches its maximal length and volume. At the end of the anagen phase, the pigment cells are actively generated and the follicle reaches its maximal length and volume. In the telogen phase, a relatively quiescent state, keratin production ceases and the club hair matures. After completion of the telogen phase, the hair begins to shed and the hair cycle restarts (Paus and Foitzik, 2004). It is known that the regulation of follicular morphogenesis and hair growth partly depends on the interaction between the epithelial and mesenchymal cells in hair follicles. The dermal papilla, a mesenchymal cell population located at the base of the hair follicle, plays an important role in regulating hair growth and cycling (Botchkarev and Kishimoto, 2003). Factors secreted by dermal papilla cells (DPCs) directly promote the surrounding matrix cells either to proliferate and differentiate or to stimulate hair stem cells to initiate a new anagen phase (Kang et al., 2010).

Recent studies in transgenic and knockout mouse models have revealed that the WNT/β-catenin-mediated signaling pathway plays a pivotal role in regulating hair follicle morphogenesis, hair shaft differentiation and follicular recycling (Kitagawa et al., 2009; Soma et al., 2012; Tsai et al., 2014). Reddy et al. (2001) demonstrated that certain WNT ligands, e.g., WNT-10a and WNT-10b, are overexpressed at the onset of the anagen phase and WNT-5a is selectively expressed in the dermal follicle.
cell at a later stage of follicular differentiation in a sonic hedgehog (SHH)-dependent manner. A WNT downstream signaling molecule, β-catenin has established a link between WNT signaling and SHH expression: the stabilization of epidermal β-catenin induced the formation of ectopic hair follicles and SHH expression (Gat et al., 1998), whereas the expression of Wnt/β-catenin signaling is mediated by signal transducer and activator of transcription (STAT) and its upstream regulator, Janus-activated kinase (JAK). A previous study demonstrated that mutation of mouse STAT3 prevents normal progression of telogen follicles into anagen (Sano et al., 2000). Pharmacological inhibition of the JAK-STAT pathway promoted rapid hair regrowth in alopecia areata (AA) in both mice and humans (Xing et al., 2014). Complying with this result, a clinically-approved JAK inhibitor, ruxolitinib, was also reported to reverse AA (Xing et al., 2014). In addition, another JAK inhibitor tofacitinib increases the growth rate of anagen hair shafts (skin grafts and organotypic culture assays) and enhances the inductivity of human dermal papilla spheres (neogenesis assays) (Zedler et al., 2016). Investigation of the molecular effects of tofacitinib treatment revealed that the treatment causes a molecular restoration of a subset of genes that are disrupted in culture but are present in fully inductive dermal papilla cells (HDPCs) (Harel et al., 2015).

3-Deoxysappanchalcone (3-DSC) is a naturally-occurring chalcone compound (Fig. 1A) isolated from Caesalpinia sappan L. (Leguminosae). C. sappan is commonly used as a herbal medicine to reduce inflammation and improve blood circulation (Shen et al., 2007; Liu et al., 2009; Yodsaoeue et al., 2009). Several studies have demonstrated that 3-DSC exerts several biological properties, including anti-allergic (Liu et al., 2009), anti-influenza virus (Yang et al., 2012), anti-inflammatory (Yodsaoeue et al., 2009), and antioxidant activities (Youn et al., 2011). In an effort to identify novel natural products that might promote hair growth, we observed that 3-DSC exerts stimulatory effects on hair growth in mice. Our study also demonstrates the potential molecular mechanisms of action of 3-DSC in the proliferation of HDPCs with a special focus on modulation of STAT and WNT/β-catenin signaling.

MATERIALS AND METHODS

Cell culture

The human hair follicle dermal papilla cells (HDPCs; Promo cell; ABM Inc., Richmond, BC, Canada) were cultured in PrigrowII (ABM Inc.) medium supplemented with 10% fetal bovine serum (FBS) and 100 U/ml penicillin and 100 μg/ml streptomycin. WNT reporter NIH3T3 cells lines were obtained from Enzo Life Sciences (Farmingdale, NY, USA) for TCF/LEF transcription factor to activate Wnt target gene expression. Stable STAT3 Luciferase-(LUCPorter™) reporter gene-expressing HEK293 cell lines were purchased from Novus (Littleton, CO, USA). Cells were incubated in accordance with the product manual and maintained in an incubator in a humidified atmosphere of 5% CO₂ at 37°C.

Chemicals and antibodies

3-DSC (purity >98%) was purchased from AK Scientific, Inc (Union City, CA, USA). CellTiter-Glo® Luminescent Cell Viability Assay kit was purchased from Promega Corporation (Madison, WI, USA). DMEM and fetal bovine serum (FBS) were procured from Invitrogen (Carlsbad, CA, USA). Interleukin (IL)-6 and IL-4 were purchased from R&D systems (Minneapolis, MN, USA). β-actin antibody was obtained from Sigma-Aldrich (St. Louis, MO, USA). Polyclonal antibodies against total β-catenin, phospho-specific β-catenin (Thr41/Ser45), total STAT3, STAT6, phospho-specific STAT3 (Tyr705) and STAT6 (Tyr641) were purchased from Cell Signaling Technology (Beverly, MA, USA). All other chemicals used in our experiments were molecular biology grade.

Real-time cell analyzer (RTCA) system

The xCELLigence System (ACEA Biosciences; San Diego, CA, USA) allows for label-free and real-time monitoring of cellular processes, such as cell proliferation, cytotoxicity, adhesion, viability, invasion, and migration, using the electronic cell sensor array technology (Ke et al., 2011). Electrode im-

Fig. 1. Effects of 3-DSC on hair cell growth. (A) Chemical structures of 3-DSC. Effects of 3-DSC on hair cell growth was examined by (B) real-time xCELLigence system and (C) CellTiter-Glo® luminescent cell growth assay as described in the Methods. All experiments were performed in triplicate. The asterisk indicates a significant statistical significance (*p<0.05).
pedance, which is displayed as cell index (CI) values, was used to provide quantitative information about the biological status of cells, including cell number, viability, and morphology. Changes in the cell status, such as cell morphology, cell adhesion, or cell viability led to a change in the CI, which is a quantitative measure of the number of cells present in a well. Subsequently, 150 µl of cell culture medium at room temperature was added to each well of E-plate 8 in the xCELLigence System. After that, the E-plate 8 was connected to the system and checked in the cell culture incubator for proper electrical contacts and the background impedance was measured during 24 hrs. Meanwhile, the HDPCs were resuspended in cell culture medium and adjusted to a cell number of 20,000 cells/well. Cell suspension (50 µl) was added to the 150 µl medium-containing wells in E-plate 8, in order to determine the optimum cell concentration. After 30-min incubation at room temperature, E-plate 8 was placed in the cell culture incubator. Then, adhesion, growth, and proliferation of the cells were monitored every 1 hr for a period of up to 24 h via the incorporated sensor electrode arrays in the E-Plate 8. After 24 hr, 0-3 M of 3-DSC was added to 200 µl cell culture medium and live cells were monitored every 15 min for a period of up to 96 hr. Electrical impedance was measured by the RTCA-integrated software of the xCELLigence system as a dimensionless parameter termed CI.

CellTiter-Glo® luminescent cell growth assay

Cell proliferation and cytotoxicity were assessed using a CellTiter-Glo® Luminescent Cell Viability Assay Kit (Promega Corporation), which is a homogeneous method to determine the number of viable cells in culture based on quantitation of the ATP present. Briefly, cells were seeded for 24 h in a 96-well plate (10,000 cells/well) and then attached cells were treated with 3-DSC (0-3 M) in serum free medium for 48 h. A volume of CellTiter-Glo® Reagent equal to the volume of cell culture medium present in each well was added and incubated at room temperature for 10 minutes to stabilize the luminescent signal. Amounts of ATP were determined by recording luminescence on a LuBi microplate luminometer (Micro Digital Ltd., Seoul, Republic of Korea).

RNA isolation and Quantitative real-time PCR (qPCR)

Cells were seeded for 24 h and then attached cells were treated with 3-DSC (0-3 M) in a serum free medium for 48 h. For mRNA quantification, total RNA was extracted using NucleoSpin® RNA Kit (Macherey-Nagel GmbH & Co., Düren, Germany). cDNA was synthesized using iScript™ cDNA Synthesis Kit (Bio-Rad, Hercules, CA, USA) according to the manufacturer’s instructions. Briefly, 2 µg of total RNA was used for cDNA preparation. The synthesized cDNA was amplified separately using primers for β-catenin, Lef/TCF, STAT3, STAT6, cyclin-dependent kinase (CDK)-4, fibroblast growth factor (FGF), vascular endothelial growth factor (VEGF) and GAPDH using GeneAmp PCR 9700 thermocycler (Thermo Fisher Scientific, Waltham, MA, USA). PCR products were analyzed by 1% agarose gel using 1X TAE buffer. Relative mRNA levels were quantified using myECL imager analysis software (Thermo Fisher Scientific). Quantitative real-time PCR was performed using the iQ™ SYBR® Green Supermix (Bio-Rad) specific for each gene. All reverse transcription reactions were run on a CFX96™ Real-Time System (Bio-Rad) using the following steps: 3 min at 95°C, 42 cycles of 10 s at 95°C, 15 s at 55°C, 30 s at 72°C, and then 10 s at 95°C. Relative expression levels were determined using the Bio-Rad CFX Manager 3.0 (Bio-Rad). The expression of target genes was normalized to that of GAPDH. The primer pairs for RT-PCR were as follows: β-catenin forward 5′-CCCCAAGACATTTCACACAGGG-3′, reverse 5′-AACAAGGCTTCTTACAGAGAG-3′; glycogen synthesis kinase (GSK)-3β forward 5′-AATCCATGGCCGAGCATAC-3′, reverse 5′-TGGTCTGGAGGGTTCTTCTT-3′; BAX forward 5′-CACCAGATATGATGATTCTT-3′, reverse 5′-GGACTTTATGAGTGTCTTGCTTTG-3′; STAT3 forward 5′-GAGATTTATGTTGAGCTTTTGTTGCTTACAG-3′, reverse 5′-ATGGCCAGGATATGATGCTT-3′; STAT4 forward 5′-AAGCTACCTTGTGATCAAACTCTA-3′, reverse 5′-TCTCTCTCTCCCTTTAAACA-3′; STAT5α forward 5′-CCCTGCTCTCTTAAGAGAGA-3′, reverse 5′-TGGATTGATCAACACAGAT-3′; STAT5β forward 5′-TATCTCTTCTTCAGATCTCTC-3′; reverse 5′-GGCGAGGGCGACTGG-3′; BMP4 forward 5′-CAGCTTGACTGAACTCTACAG-3′, reverse 5′-GGGACCCACATCCCTCTACT-3′; FGF forward 5′-GCTCTAGCAGACATTGGAG-3′, reverse 5′-GGTGTGTTGTCAACGTTTTCA-3′, VEGF forward 5′-GGGACCCACATCCCTCTACT-3′, reverse 5′-GGCGAGGGCGACTGG-3′; WNT-3A forward 5′-ACCTGAGATGATGATTCTT-3′, reverse 5′-AAGCGAGATTTGTTGAGCTTTG-3′, and GAPDH forward 5′-TGGC- AAATCTTATGGAAAC-3′, reverse 5′-CCATGTTGTTGAGCTTTG-3′.
medium were homogenized with a cell lysis buffer (100 mM Tris pH 7.5, 150 mM NaCl, 5 mM EDTA, 1% Triton X-100, 5 mM DTT, 0.1 mM PMSF, 10% Glycerol, protease inhibitor) and lysed with 2 h incubation on ice. The cell lysate was centrifuged at 13,000 rpm for 15 min at 4°C. Equal amounts of proteins (20 μg) were separated on a SDS/8%-polyacrylamide gel, and then transferred to a polyvinylidene difluoride (PVDF) membrane (Thermo Scientific). Blots were blocked for 1 h at room temperature with 5% (w/v) non-fat dried milk in Tris-Buffered Saline Tween-20 [TBS-T: 10 mM Tris (pH 8.0) and 150 mM NaCl solution containing 0.05% Tween-20]. After a short washing in TBST, the membranes were immunoblotted with specific antibodies. To detect target proteins, specific antibodies against stat3, stat6, phospho-stat3, phospho-stat6, β-catenin, phospho-β-catenin (1:1000, Cell Signaling Technology) and β-actin (1:5000, Sigma-Aldrich) were used. The blots were then incubated with the corresponding conjugated goat anti-rabbit or goat anti-mouse or donkey anti-goat IgG-horse-radish peroxidase (HRP) (1:5000; Santa Cruz Biotechnology Inc., Santa Cruz, CA, USA) secondary antibodies. Immunoreactive proteins were detected with an enhanced chemiluminescence western blotting detection system (myECL imager, Thermo Scientific).

**Hair growth activity in mice**

Seven-week-old female C57BL mice were purchased from Oriental Bio Co (Seoul, Republic of Korea). After a 7 day acclimation period for being automatically maintained at 21-25°C and a relative humidity of 45-65% with a controlled light-dark cycle, the animals were divided into 2 randomized groups (n=4) to investigate the hair growth promoting activity of 3-DSC. Two hundred microliters of 3-DSC (3 μM) were applied twice daily for 15 days. Reagents used for the hair growth test were dissolved in a vehicle containing 50% ethanol. All animals were cared for by using protocols approved by the Institutional Animal Care and Use Committee (Chungbuk, Republic of Korea).
RESULTS

The effects of 3-DSC on hair follicle dermal papilla cell growth

Since 3-DSC has been reported to elicit the activation of cell survival pathways (Kim et al., 2014), we examined the effects of 3-DSC on growth of HDPCs. Analysis of real-time cell proliferation using xCELLigence system revealed that the proliferation of HDPCs was promoted by 3-DSC treatment in a concentration-dependent manner (Fig. 1B). Likewise, 3-DSC resulted in a dose-dependent increase in the viability of HDPCs (Fig. 1C). Tofacitinib, which has been reported to promote growth of the hair shaft (Harel et al., 2015), was used as a positive control.

Effects of 3-DSC on the expression of genes involved in hair growth regulation

To investigate the effect of 3-DSC on hair growth regulation factors, we analyzed the transcriptional expression changes in various genes, using a conventional and quantitative reverse-transcriptase PCR (RT-PCR). As a result, 3-DSC caused transcriptional activation of β-catenin, tcf, fgf, vegf, cdk4 and stat3, whereas it decreased the level of stat4, stat5A/B and stat6.
mRNAs (Fig. 2A). We observed that the expression of Bax, Bcl-2, Sox-9 and BMP4 mRNAs was unaltered. This finding was further confirmed by conventional RT-PCR analysis and tofacitinib elicited a similar pattern of gene expression changes in HDPCs (Fig. 2B).

**Effects of 3-DSC on the transcriptional activity of TCF, STAT3 and STAT6 in HDPCs**

Next, we analyzed the effects of 3-DSC on TCF/LEF-mediated reporter activity, using NIH3T3-WNT-luciferase cells (Fig. 3A). As a result, we observed that 3-DSC resulted in TCF/LEF luciferase activation. In order to determine the effects of 3-DSC on STAT3-mediated transcriptional activity, we exposed the cells to 3-DSC and measured the STAT3-mediated luciferase activity in HEK293 cells that stably expressed a STAT3-regulated luciferase reporter plasmid (Fig. 3B). As a result, we observed that IL-6 stimulated STAT3 transcriptional activity in a concentration-dependent manner. In or-
der to determine whether the inhibitory effects of 3-DSC on STAT6 phosphorylation can be ascribed to the attenuation of the STAT6 transcriptional response, we analyzed the effects of 3-DSC on IL-4 receptor site-mediated reporter gene expression using a stably transfected HEK293 cell line that expresses the IL-4R site-TKluc/STAT6 regulated luciferase gene after treatment with IL-4 (Fig. 3C). 3-DSC decreased the IL-4R site-TKluc/STAT6 luciferase activity in a dose-dependent manner. It seems that 3-DSC increased the TCF/LEF activity by stabilizing β-catenin through the inhibition of β-catenin phosphorylation. 3-DSC also decreased IL-4-induced phosphorylation of STAT6 via interfering with the JAK1/3 pathway, and it activated phosphorylation of STAT3 via the IL-6-induced JAK2 pathway. These results indicate that 3-DSC-induced hair growth promotion is caused by regulating the molecular target of hair dermal follicular papilla cells.

**Effects of 3-DSC on phosphorylation of β-Catenin, STAT3 and STAT6 in HDPCs**

Based on the effects of 3-DSC on the mRNA expression of certain hair growth regulatory genes, we examined whether 3-DSC can modulate the phosphorylation of several key regulators of hair growth. As shown in Fig. 4A, 3-DSC decreased constitutive phosphorylation of β-catenin (Thr41 and Ser45). Treatment of HDPCs with WNT3a, which stabilizes β-catenin, also abolished constitutive β-catenin phosphorylation. Similar inhibition of β-catenin phosphorylation was noted when HDPCs were exposed to the standard drug tofacitinib. To examine the effects of 3-DSC on STAT3 and STAT6 phosphorylation, HDPCs were stimulated with IL6 and IL4, respectively, because STATs are not constitutively phosphorylated in HDPCs. As a result, 3-DSC exhibited inhibitory effects on IL6-induced STAT3 phosphorylation at the Tyr705 residue at lower concentrations (0.1 and 0.3 μM), but STAT3 phosphorylation was unchanged at a higher concentration of the compound (Fig. 4B). However, 3-DSC attenuated IL4-induced STAT6 phosphorylation at the Tyr641 residue in a concentration-dependent manner and it was comparable to tofacitinib (Fig. 4C).

**Effects of 3-DSC on hair growth in mice**

Finally, we attempted to examine the effects of 3-DSC on hair growth in mice. The back of C57BL/6 mice was shaved and topically treated with a vehicle (ethanol) or 3-DSC for 15 days. Compared to the vehicle control, 3-DSC promoted rapid and intense hair growth in mice (Fig. 5A, 5B). Histopathological analysis of mouse skin including the follicular and dermal layers at autopsy showed that the diameter and depth of the hair follicles were remarkably higher in mice that were administered with 3-DSC. A representative scheme of 3-DSC regulation of WNT/β-catenin and JAK-STAT pathway in human hair dermal follicle papilla cells (Fig. 5C).

**DISCUSSION**

*Caesalpinia sappan* L. has been reported to have various beneficial pharmacological activities such as immune function modulation, depression of the central nervous system, anti-inflammation, and vasorelaxation. The present study was designed to investigate the effect of 3-DSC on hair growth and it revealed that 3-DSC promoted in vivo hair growth and modulated intracellular signaling pathways, implicated in hair cycle regulation. The remodeling of hair follicles involves cyclical periods of growth (anagen), regression (catagen), rest (telogen) and shedding (exogen) (Paus and Cotsarelis, 1999). Many follicles undergo programmed cell death during catagen that leads to reduced hair size at the beginning of the telogen phase (Cotsarelis, 1997). Follicular regeneration at the onset of the subsequent anagen phase requires activation of rarely cycling epithelial stem cells located in the permanent, bulge region of the follicle (Cotsarelis *et al.*, 1990). This stem cell progeny forms a new follicular matrix during early anagen, and the hair shaft and inner root sheath are derived from these relatively undifferentiated matrix cells (Oshima *et al.*, 2001).
The size and length of the hair shaft correspond to the size of the hair follicle and to the duration of anagen, respectively. It is well established that STAT3 is one of the factors required for anagen onset (Sano et al., 2000). Activation of another intracellular signal pathway mediated via WNT-β-catenin is also required for the proliferation and differentiation of the hair shaft (Millar et al., 1999; Kishimoto et al., 2000; Cotsarelis and Millar, 2001). In fact, the volume of the dermal papilla reflects the number of matrix cells and it determines the size of the resulting hair shaft (Hardy, 1992). Because the size of the follicle is determined during the early stages of anagen, this could be a critical time for hair follicles undergoing miniaturization in androgenetic alopecia. Some factors (e.g., hormones, drugs, morphogens) might act by enhancing or preventing miniaturization only during this span of time at anagen onset, thereby requiring prolonged periods of time to alter a significant number of follicles. This might partially explain why the process of miniaturization takes years to both develop and treat.

During the progression of anagen stage, the maintenance of follicular epithelium requires transduction of signals from the dermal papilla to the follicular epithelium. Interestingly, dermal papilla cells cultured in the presence of β-catenin protein maintain their inductive abilities over many rounds of culture, suggesting that the epithelial signal is comprised of one or more WNT/β-catenin family members (Kishimoto et al., 2000; Cotsarelis and Millar, 2001). Thus, the activation of STAT3 and β-catenin appears to be the underlying mechanism of hair growth stimulation by 3-DSC. Besides STAT3, several other members of the STAT family and their upstream JAK kinase have been known to regulate hair growth. Several studies have reported that suppression of JAK signaling in mice activates a pro-growth/anti-quiescence signal during telogen (Pilikus et al., 2008; Festa et al., 2011; Jahoda and Christiano, 2011), thereby allowing entry into anagen. In particular, Harel et al. (2015) reported that inhibition of JAK-STAT signaling promotes hair growth by stimulating the activation and/or proliferation of hair follicular stem cells, highlighting the role of this pathway in maintenance of hair follicular quiescence. Increased proliferation or differentiation of stem/progenitor cells upon inhibition of JAK-STAT signals is not unique only for hair follicular cells, but it has also been observed in other types of progenitor cells to the hair follicles. For example, the loss of STAT5 in hematopoietic stem cells induces exit from a quiescent state, leading to increased bone marrow-repopulating capacity after irradiation (Wang et al., 2009). Likewise, inhibition of JAK-STAT signaling improves skeletal muscle regeneration in aged mice by promoting expansion of symmetric satellite cells under their engraftment in vivo (Price et al., 2014). These findings are consistent with the involvement of the JAK-STAT pathway in the maintenance of quiescence in the hair follicular cells (Lin et al., 2004) and the role of STAT3 in progression of the normal hair cycle in adult mice (Sano et al., 2000). Moreover, recent studies have shown that increased JAK-STAT signaling in aged mice inhibits hair follicular stem cell function in vitro (Doles et al., 2012). Goldstein et al. (2014) reported that quiescence of hair growth during pregnancy and lactation was partly mediated through prolactin-induced phosphorylation of Stat5. Therefore, JAK-STAT signaling appears to play a generalized role in promoting quiescence in adult stem cell populations.

As a first report, our study revealed that 3-DSC attenuated mRNA expression, and IL-4-induced phosphorylation of Stat5, which is a signaling molecule downstream of JAK3 in cultured HDPCs. Although our study showed that tofacitinib, a known JAK inhibitor, suppressed mRNA expression and IL-4-induced phosphorylation of STAT6, whether 3-DSC, which has a similar effect, can attenuate JAK activation is yet to be examined. However, the present study delineates the effects of 3-DSC on STATs, which are effectors signaling molecules downstream of JAK, thus indicating the key molecular aspect of hair follicle stimulation by 3-DSC via modulation of STATs. Histopathological analysis of mouse dermis also revealed that 3-DSC increased the diameter and depth of the hair follicle in the dermis. In conclusion, our study suggests that 3-DSC promotes proliferation of dermal papilae and stimulates hair growth partly via activation of Wnt/β-catenin signaling and inhibition of STAT6-mediated quiescence of hair follicular cells.

CONFLICT OF INTEREST

Authors declare no competing financial interests.

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