SHORT COMMUNICATION



Mushroom extract inhibits ultraviolet B-induced cellular senescence in human keratinocytes

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Abstract Mushrooms possess various bioactivities and are used as nutritional supplements and medicinal products. Twenty-nine bioactive components have been extracted recently from mushrooms grown in Nepal. In this study, we evaluated the ability of these mushroom extracts to augment SIRT1, a mammalian SIR2 homologue localized in cytosol and nuclei. We established a system for screening food ingredients that augment the SIRT1 promoter in HaCaT cells, and identified a SIRT1-augmenting mushroom extract (number 28, *Trametes versicolor*). UVB irradiation

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K. Shimizu · Y. Katakura (⊠) Faculty of Agriculture, Kyushu University, 6-10-1 Hakozaki, Higashi-ku, Fukuoka 812-8581, Japan e-mail: katakura@grt.kyushu-u.ac.jp induced cellular senescence in HaCaT cells, as evidenced by increased activity and expression of cellular senescence markers including senescenceassociated β -galactosidase, p21, p16, phosphorylated p38, and γ H2AX. Results clearly showed that the mushroom extract (No. 28) suppressed the ultraviolet B irradiation-induced cellular senescence in HaCaT cells possibly through augmenting SIRT1 expression.

Keywords SIRT1 · Mushroom · Cellular senescence · Keratinocytes · Skin · Ultraviolet B

Introduction

Accumulating evidence has demonstrated that ultraviolet radiation (UV) is the most common environmental factor that damages the human skin, leading to conditions such as sunburn, solar erythema, inflammation, skin carcinogenesis, and premature senescence (Pillai et al. 2005). In particular, UVB (a wavelength range of 280–320 nm) has been shown to permeate through the epidermis into the dermis and induce reactive oxygen species (ROS) production in the skin and cultured skin cells, thus contributing to gene mutations, abnormal cellular proliferation, and skin aging (Lavker et al. 1995; Hattori et al. 1996; Ahmed et al. 1999).

Sirtuins, a family of NAD⁺-dependent enzymes, are well-known modulators of lifespan in many species (Kaeberlein et al. 1999). Sirtuin1 (SIRT1), the closest homologue to yeast Sir2, is the most widely studied of the seven sirtuin family members (Frye 2000). Evidence has shown that SIRT1 is involved in cardiovascular disease, neurodegenerative diseases such as Alzheimer's disease and amyotrophic lateral sclerosis, and inflammatory diseases (Lavu et al. 2008). Furthermore, several lines of evidence have shown that SIRT1 prevents cellular senescence in endothelial cells and fibroblasts via various mechanisms (Huang et al. 2008; Zu et al. 2010; Yamashita et al. 2012). However, experimental evidence regarding the role of SIRT1 in stress-induced fibroblast senescence is sparse, particularly with respect to the protective effect of SIRT1 against UVB-induced premature cell senescence.

Mushrooms, as one of the potential sources of dietary antioxidants, have been valued not only for their nutritional properties (Barros et al. 2008), but also for various medicinal benefits (Lindequist et al. 2005; Ajith and Janardhanan 2007). Around 140,000 species of mushrooms are believed to exist (Wasser 2002); among these, wild mushrooms from Nepal are report to be diverse and play vital roles in many local communities (Adhikari et al. 2006). However, few studies of the pharmacological potential or bioactivity of mushrooms grown in Nepal have been published. Recently, Hai Bang and colleagues described the antioxidant activity of 29 mushrooms collected from the mountainous areas of Nepal (Hai Bang et al. 2014; Tamrakar et al. 2016). To our knowledge, the effects of such mushroom extracts on cellular senescence have not been investigated to date. In this study, we evaluated the protective effects of these twenty-nine mushroom extracts on UVB-induced cellular senescence in human keratinocytes.

Materials and methods

Cell culture and reagents

The HaCaT human keratinocyte cell line (Riken Bioresource Center, Tsukuba, Japan) was cultured in Dulbecco's modified Eagle's medium (DMEM; Nissui, Tokyo, Japan) supplemented with 10% fetal bovine serum (Life Technologies, Gaithersburg, MD, USA) at 37 °C in an atmosphere containing 5% CO₂.

Screening system for foods that activate the SIRT1 promoter

We amplified the human SIRT1 promoter (-1593 to - 1) by PCR using human genomic DNA as the template; the amplified fragment was cloned into the promoter-less pEGFP-C3 (Takara, Shiga, Japan) and designated pSIRT1p-EGFP (Harada et al. 2016). HaCaT cells transduced with pSIRT1p-EGFP (HaCaT (SIRT1p-EGFP)) were treated with mushroom extracts, and human SIRT1 promoter activity was then evaluated. Changes in the EGFP fluorescence derived from pSIRT1p-EGFP were monitored using an IN Cell Analyzer 1000 (GE Healthcare, Amersham Place, UK) (Udono et al. 2012).

Ultraviolet B (UVB) irradiation

The medium was removed and cells were exposed to UVB (Integrated Irradiance Level: 10 mJ/cm²; CL-1000 Ultraviolet Crosslinker, UVP, Upland, CA, USA). This exposure was repeated at an interval of 24 h. Soon after UVB irradiation, the medium in which cells were cultured was replaced with fresh medium.

Mushroom samples

Mushrooms were collected from several forests in various parts of Nepal. Scientific names, locations, and accession numbers of the mushrooms are provided in Table S1. Mushroom samples were identified on the basis of morphological features and/or genetic analysis, as described previously (Tamrakar et al. 2016). The samples were dried in an air-ventilated oven at 35 °C for 10 h, followed by at 45 °C for 1 h. The dried samples were ground to a fine powder and extracted with ethanol. The ethanol extracts were rotary-evaporated at 45 °C.

Fluorescence senescence-associated β -galactosidase (SA- β -Gal) assay

Fluorescence SA- β -Gal assay was carried out, as described previously (Udono et al. 2012). Briefly, HaCaT cells were fixed with 2% formaldehyde/0.2%

glutaraldehyde, then stained with 33 µM ImaGene Green C₁₂FDG (Life Technologies) and 1 µg/mL Hoechst 33342 solution (Dojin, Kumamoto, Japan). Image acquisition and analysis of imaging data was performed using an IN Cell Analyzer 1000 (GE Healthcare), as described previously. The imaging data were reported as SA-\beta-Gal intensity (mean fluorescence intensity per cell) and the SA-β-Galpositive/negative cells by setting the threshold intensity of the SA- β -Gal staining. The threshold of SA- β -Gal intensity was set to the point at which about 75% of total cells were negative in the control cells. The data collected using Developer were analyzed by Spotfire DecisionSite Client 8.2 (PerkinElmer, Waltham, MA, USA) software to visualize the results. This experiment was repeated at least 3 times, and the representative data are shown.

Immunofluorescence

Immunofluorescence was performed as described previously (Udono et al. 2012). Briefly, HaCaT cells were seeded onto a µClear Fluorescence Black plate (Greiner Bio-One, Tokyo, Japan). After fixing the cells with 4% paraformaldehyde, cells were blocked with 5% goat serum and 0.3% Triton-X100. After washing the cells, specific antibodies (anti-p16INK4a (ab81278, Abcam, Cambridge, UK), anti-p21 (2947, Cell Signaling, Danvers, MA, USA), anti-phospho p38 (4511, Cell Signaling), and anti-phospho histone H2A.X (9718, Cell Signaling) were added. After washing the cells, secondary antibody (Alexa Fluor 555 F(ab') fragments of goat anti-rabbit IgG (Life Technologies) were added. After washing the cells, Hoechst 33342 solution was added, and an image of each well was acquired using an IN Cell Analyzer 1000. Imaging data were analysed using the Multi Target Analysis tool and visualized using Spotfire DecisionSite Client 8.2 software. This experiment was repeated at least 3 times, and the representative data were shown.

Quantitative RT-PCR (qRT-PCR)

RNA was isolated using the High Pure RNA Isolation Kit (Roche, Mannheim, Germany). cDNA was generated from the isolated RNA using the ReverTra Ace (Toyobo, Osaka, Japan). qRT-PCR was performed using the KAPA SYBR Fast qPCR Kit (KAPA Biosystems, Woburn, MA, USA) and the Thermal Cycler Dice Real Time System TP-800 instrument, as described previously (Udono et al. 2012). The samples were analyzed in triplicate, and the SIRT1 level was normalized to the corresponding β -actin level. The PCR primer sequences used were as follows: SIRT1 forward primer 5'-GCCTCACATGCAAGCTCTAG TGAC-3' and reverse primer 5'-TTCGAGGATCT GTGCCAATCATAA-3'; β -actin forward primer 5'-TGGCACCCAGCACAATGAA-3' and reverse primer 5'-CTAAGTCATAGTCCGCCTAGA AGCA-3'.

Retrovirus production and transduction

Viral supernatants were produced after transfecting 293 T cells with pGag-pol, pVSV-G, and individual expression vectors (pBABE-puro, pBABE-puro-SIRT1) using the HilyMax reagent (Dojindo), as previously described (Udono et al. 2012). The cells were cultured at 37 °C in DMEM supplemented with 10% FBS for 24 h. Medium was then replaced with fresh DMEM supplemented with 2% FBS and incubated for an additional 24 h. Viral supernatant was collected and supplemented with 10 mg/mL polybrene (Merck Millipore, Billerica, MA, USA). The target cells were infected with this viral supernatant for 24 h at 37 °C. After infection, the cells were selected using 3 μ g/mL puromycin (Enzo Life Sciences, Farmingdale, NY, USA) for 3 days.

Statistical analysis

All experiments were performed at least 3 times. The results are presented as mean \pm SD. Statistical significance was determined using a two-sided Student's *t* test. Multiple comparisons between groups were made by one-way ANOVA with Turkey's post hoc test. Statistical significance was defined as p < 0.05.

Result and discussion

SIRT1 overexpression protects HaCaT cells from UVB-induced senescence

Although UVB irradiation is known to induce apoptosis in HaCaT cells (Li et al. 2017), keratinocytes possess the ability to completely repair DNA damage induced by low-energy UVB (Integrated Irradiance Level: 5 mJ/cm²) exposure and maintain normal cellular function (Kim et al. 2016). In the present study, we repeatedly treated HaCaT cells with lowenergy UVB, and evaluated the induction of cellular senescence by measuring SA- β -Gal activity, a biomarker of cellular senescence, using an IN Cell Analyzer 1000 (Udono et al. 2012). Cellular senescence induction was evaluated by analyzing cell size (x-axis) as well as the percentage of SA- β -Galpositive/negative cells (y-axis) (Fig. 1a). In comparison with that observed in non-treated HaCaT cells, the percentage of SA- β -Gal-positive cells markedly

increased among the UVB-irradiated population (Fig. 1a; pie chart). In addition, as shown in Fig. 1a, an increase in the size of UVB-exposed HaCaT cells was observed (Fig. 1a; pie chart). These data indicate that repeated irradiation with low-energy UVB radiation induced cellular senescence in HaCaT cells.

SIRT1, popularly known as the longevity gene, has been expressed in cultured skin keratinocytes; the activity of this gene was shown to be downregulated by both UV and H_2O_2 treatment in a time- and dosedependent manner (Cao et al. 2009). To investigate the effects of SIRT1 on UVB-induced senescence in



Fig. 1 Effects of SIRT1 on UVB-induced cellular senescence in HaCaT cells. **a** Effects of UVB on SA-β-Gal activity in HaCaT cells. Cells (9 × 10⁵ cells) were seeded on to culture dish (φ 35 mm), and cultured for 24 h. Medium was removed and cells were irradiated with UVB (Integrated Irradiance Level: 10 mJ/cm²) and cultured in DMEM supplemented with 10% serum. Next day this UVB irradiation procedure was repeated, and cells were cultured for 1 day, followed by detection of SA-β-Gal activity using an IN Cell Analyzer 1000. Cellular SA-β-Gal (y-axis) and cell size (x-axis) were plotted on a scatter plot. The relative number of cells with high SA-β-Gal activity are shown in the pie chart. Statistical significance was determined using a two-sided Student's *t* test. Significant difference is denoted by ^{\$\$\$\$} *p* < 0.001. **b** Effects of

SIRT1 on UVB-induced cellular senescence in HaCaT cells. Recombinant HaCaT cells with ectopic SIRT1 expression or mock expression were irradiated with UVB (10 mJ/cm²). Senescence-related biomarkers, including SA- β -Gal activity, γ H2AX, p16, p21, and phospho-p38 MAP kinase, were detected using an IN Cell Analyzer 1000. The relative number of cells with high activity or expression of senescence biomarkers is shown in black in the pie chart. The size of the pie chart demonstrates the average cellular size. Multiple comparisons against non-treated mock-transfected HaCaT (#) or against UVB-treated mock-transfected HaCaT (*) were calculated by one-way ANOVA with Turkey's post hoc test. Significant differences are denoted by ###, *** p < 0.05 HaCaT cells, we first established recombinant HaCaT cells with ectopic expression of SIRT1. Senescencerelated biomarkers, such as SA- β -Gal activity, γ H2AX, p16, p21, and phospho-p38 (p-p38), were examined in HaCaT cells irradiated with UVB using an IN Cell Analyzer 1000. Figure 1b clearly shows that UVB induced cellular senescence in HaCaT cells, and, further, that SIRT1 attenuated UVB-induced cellular senescence. Taken together, these data suggest that SIRT1 protects HaCaT cells from UVBinduced cellular senescence.

Identification of mushrooms that augment SIRT1 transcription in HaCaT cells

Previously, we established a system for screening foods/food ingredients that augment SIRT1 and SIRT3 expression in Caco-2 cells, and identified Lactobacillus brevis T2102 and pomegraniin A as SIRT1- and SIRT3-activating food components (Zhao et al. 2016; Harada et al. 2016). Here, we established a similar system for screening foods/food ingredients that activate SIRT1 expression using HaCaT cells that express EGFP under the control of the SIRT1 promoter. We tested the SIRT1-augmenting activity of twenty-nine extracts of mushrooms grown in Nepal using this cell line (Table S1). As shown in Fig. 2a, extract numbers 9, 27, and 28 significantly augmented SIRT1 promoter activity in HaCaT cells. In this assay, EGFP fluorescence intensity depends on promoter activity of SIRT1 and cell condition. Thus, we tested whether these mushroom extracts augment endogenous SIRT1 expression in HaCaT cells by qRT-PCR (Fig. 2b). Results showed that extract number 28 significantly augmented endogenous SIRT1 expression in HaCaT cells.

Mushroom extract inhibits UVB-induced senescence in HaCaT cells

Next, we tested whether mushroom extract number 28 inhibited UVB-induced senescence in HaCaT cells. UVB induced cellular senescence in HaCaT cells, as evidenced by the activation of SA- β -Gal activity and increase in cellular senescence markers (Fig. 2c). Extract number 28 (*Trametes versicolor*) markedly attenuated UVB-induced cellular senescence in HaCaT cells; however, resveratrol, which is a known

SIRT1-activating polyphenol, had no effects on UVBinduced cellular senescence.

Trametes versicolor, also known as Yunzhi, is known to possess a wide range of beneficial medical properties such as protective effects against oxidative damage, cancer, and bacterial/viral infection as well as immune-potentiating activity; further, this mushroom has been shown to elicit improvement in bone properties in diabetic rats (Lindequist et al. 2005; Chen et al. 2015; Jhan et al. 2016). The best-known commercial polysaccharopeptide preparations derived from T. versicolor are polysaccharopeptide Krestin (PSK) and polysaccharopeptide (PSP). PSPs, which are among the main bioactive constituents of T. versicolor, are recognized as biological response modifiers that are useful adjuncts to conventional therapy. PSPs have been reported to increase the activity and gene expression of antioxidant enzymes and reduce lipid peroxidation in senescence-accelerated mice (Li et al. 2007), as well as exert immunomodulatory effects on blood lymphocytes and the breast cancer cell line MCF-7 (Kowalczewska et al. 2016). Until now, several responsible bioactive compounds in mushrooms including polysaccharides, triterpenoids, proteins, phenolic compounds have been reported (Hai Bang et al. 2014). We need to identify active compounds in the extract in the future study. In the present study, we identified novel activity of extracts derived from T. versicolor, which is expected to be of value in applications related to skin maintenance and repair of skin damage induced by UVB, as well as in the production of novel cosmetics targeting skin aging.

UVB-exposed edible white button mushrooms have been proven to be a safe and effective source of vitamin D2, which supports bone growth and regulates the immune response (Kalaras et al. 2012; Calvo et al. 2013). In the present study, mushrooms were collected from high-altitude regions in Nepal, and therefore considered to possess strong pharmacological potential as a result of UV exposure.

ROS production represents the main source of damage in UVB-irradiated cells. Numerous bioactive components protect HaCaT cells from UVB-induced photo-damage through ROS clearance (Zheng et al. 2016; Oh et al. 2016; Yuan et al. 2017; Zhu et al. 2017; Li et al. 2017). Furthermore, it has been reported that SIRT1 expression is downregulated by UV irradiation through ROS-mediated JNK activation (Cao et al.



Fig. 2 Effects of mushroom extract number 28 on UVBinduced senescence in HaCaT cells. a Identification of mushroom extracts that augment SIRT1 transcription in HaCaT cells. HaCaT cells transduced with vector expressing EGFP under the control of the SIRT1 promoter $(2.0 \times 10^4 \text{ cells})$ were seeded on to 96-well µClear Fluorescence Black plate and cultured for 24 h. Mushroom extracts (10 µg/mL) were added and changes in EGFP fluorescence were monitored using an IN Cell Analyzer 1000. b Effects of mushroom extracts on the expression of endogenous SIRT1 in HaCaT cells. HaCaT cells were treated with 10 µg/mL of mushroom extracts. Next day, RNA was prepared and quantitative RT-PCR was conducted. The results are expressed as mean \pm SD. Statistical significance was defined as *p < 0.05, ***p < 0.001. c Effects of mushroom

extract number 28 on UVB-induced senescence in HaCaT cells. Cells were firstly treated with resveratrol and mushroom extract number 28 (10 μg/mL) every 3 days, irradiated with 10 mJ/cm² UVB in the presence of extract, and cultured for 3 days. Senescence-related biomarkers, including SA-β-Gal activity, γH2AX, p16, p21, and phospho-p38 MAP kinase, were detected using an IN Cell Analyzer 1000. Resveratrol (5 μM) was used as a positive control. The relative number of cells with high activity or expression of senescence biomarkers is shown in black in the pie chart. Multiple comparisons against non-treated HaCaT ([#]) or against UVB-treated HaCaT ([§]) were calculated by one-way ANOVA with Turkey's post hoc test. Significant differences are denoted by ^{###, SSS} p < 0.001

2009), and that SIRT1 overexpression prevents UVBinduced fibroblast senescence by suppressing oxidative stress via the deacetylation of FOXO3 α and p53 (Chung et al. 2015). However, mushroom extract number 28 (*T. versicolor*) with strong SIRT1-augmenting ability has been previously reported to possess limited antioxidant activity (Hai Bang et al. 2014), suggesting that this mushroom extract inhibited UVB-induced cellular senescence via a ROS- independent pathway. Further studies aimed at elucidating the underlying mechanisms are therefore warranted.

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Compliance with ethical standards

Conflict of interest The authors declare that they have no conflict of interest.

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